DISSERTATION

MULTI-CRITERIA ANALYSIS IN MODERN INFORMATION MANAGEMENT

Submitted by

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WE HEREBY RECOMMEND THAT THE DISSERTATION PREPARED UNDER OUR SUPERVISION BY RINKU DEWRI ENTITLED MULTI-CRITERIA ANALYSIS IN MODERN INFORMATION MANAGEMENT BE ACCEPTED AS FULFILLING IN PART REQUIREMENTS FOR THE DEGREE OF DOCTOR OF PHILOSOPHY.

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ABSTRACT OF DISSERTATION

MULTI-CRITERIA ANALYSIS IN MODERN INFORMATION MANAGEMENT

The past few years have witnessed an overwhelming amount of research in the field of information security and privacy. An encouraging outcome of this research is the vast accumulation of theoretical models that help to capture the various threats that persistently hinder the best possible usage of today's powerful communication infrastructure. While theoretical models are essential to understanding the impact of any breakdown in the infrastructure, they are of limited application if the underlying business centric view is ignored. Information management in this context is the strategic management of the infrastructure, incorporating the knowledge about causes and consequences to arrive at the right balance between risk and profit.

Modern information management systems are home to a vast repository of sensitive personal information. While these systems depend on quality data to boost the Quality of Service (QoS), they also run the risk of violating privacy regulations. The presence of network vulnerabilities also weaken these systems since security policies cannot always be enforced to prevent all forms of exploitation. This problem is more strongly grounded in the insufficient availability of resources, rather than the inability to predict zero-day attacks. System resources also impact the availability of access to information, which in itself is becoming more and more ubiquitous day by day. Information access times in such ubiquitous environments must be maintained within a specified QoS level. In short, modern information management must consider the mutual interactions between risks, resources and services to achieve wide scale acceptance.

This dissertation explores these problems in the context of three important domains, namely disclosure control, security risk management and wireless data broadcasting. Re-

search in these domains has been put together under the umbrella of multi-criteria decision making to signify that "business survival" is an equally important factor to consider while analyzing risks and providing solutions for their resolution. We emphasize that businesses are always bound by constraints in their effort to mitigate risks and therefore benefit the most from a framework that allows the exploration of solutions that abide by the constraints. Towards this end, we revisit the optimization problems being solved in these domains and argue that they oversee the underlying cost-benefit relationship.

Our approach in this work is motivated by the inherent multi-objective nature of the problems. We propose formulations that help expose the cost-benefit relationship across the different objectives that must be met in these problems. Such an analysis provides a decision maker with the necessary information to make an informed decision on the impact of choosing a control measure over the business goals of an organization. The theories and tools necessary to perform this analysis are introduced to the community.

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CHAPTER 1

Multi-Criteria Information Management

Information management is the process of collecting information from one or more sources, followed by its dissemination to one or more consumers. This process also implicitly assumes the transient storage of the collected data prior to its distribution. The term "information" is not bound to a specific form or media, but instead generalizes to any data asset in which the managing organization has a stake. This perception is used to differentiate information management from data management where the latter typically involves management of the data needs within an organization. The former, however, also includes preserving the enterprise motives during and after the data transits from a source to a consumer.

Any modern information management system entails three broad dimensions – *risk*, *resources*, and *services*. Risk manifests itself when the data that is collected and then distributed is potentially sensitive, or deemed sensitive by the primary source of collection. In this case, the management system must have some safety guarantees associated with them. Risk also encapsulates the case where data assets may be improperly accessed either during storage or after distribution. The second dimension, services, involves the guarantees of access to data by a legitimate consumer. This factor is commonly known as "availability" of a system. However, a system must also try to assure that the information is received in a state that is usable to the consumer. The quality of service offered by the

Figure 1.1: Application domains in multi-criteria information management.

system is therefore an evaluation of its efficiency in implementing these requirements. The third dimension, resources, is related to the infrastructure that goes into realizing this process of collection, storage and dissemination of information. The focus is on how well this infrastructure is put to use, and whether all available resources are utilized to their full potential.

As in many real-world systems, balancing the multiple aspects of an information system is crucial to its proper functioning. The best approach to do so is not straightforward owing to underlying contentions. An organization may effectively increase the resource requirement or degrade information quality and availability while attempting to maintain low levels of risk. Similarly, service quality suffers when resources are limited or stringent risk related safeguards are installed. The available infrastructure capabilities also dictate the quality of service and what potential threats can be eliminated. Multi-criteria information management is therefore defined as the strategic management of an organization's data assets, including but not limited to its protection, accessibility and utility, while satisfying existing business goals and constraints.

We shall exemplify the conceived notion of multi-criteria information management using three domains in computer science (Fig. 1.1) — *disclosure control*, *security risk management* and *wireless data broadcasting*. The remainder of this dissertation is broadly divided into three parts as described next, followed by a summary of contributions and potential future work in Chapter 15.

Part I: Disclosure Control

Privacy violations emanating from the public sharing of personally identifying information have raised serious concerns over the past few years. The nature of these violations indicate that information privacy is difficult to guarantee even after the removal of unique identification data. Multiple real world instances have exemplified this aspect in recent years.

- 35% of victims were re-identified in Chicago's de-identified homicide database by comparing the records with those in the Social Security Death Index [137].

- The health records of the Governor of Massachusetts were re-identified from Group Insurance Commission's anonymized data by using a voter registration list [165].

- Unique identification is possible for approximately 70% of the population from familial database records using genealogies extracted from newspaper obituaries and death records [122].

- Users were uniquely identified based on their search queries released as part of an anonymized Web queries data set (contained over twenty million search keywords) by AOL for research purposes [14].

- A person's date of birth, gender and ZIP code forms a unique identifier for 63% of the US population reported in the 2000 census [76].

- Researchers uniquely identified Netflix$^{\circledR}$ users from an anonymized movie ratings data set (contained nearly half million records) released by the rental firm to facilitate research to improve its movie-recommendation engine [134].

- Public availability of the Social Security Administration's Death Master File and the widespread accessibility of personal information from sources such as data brokers or profiles on social networking sites enable researchers to fully predict the social security number [4].

The seriousness of these violations is well-understood in the research community and has generated significant interest in the field of disclosure control. This field of research

has looked into the formulation of data modification techniques and privacy models that can prevent possible unique linkage between a person and the corresponding information contained in a database record. However, the challenge lies in the enforcement of these techniques, predominantly due to the fact that the quality of the shared data often determines its usefulness for the legitimate purpose for which it is intended. The privacy primitives can reduce the risk of re-identification, but at the same time degrades the service quality that an organization can guarantee based on the data quality. We shall look into the limitations of the current decision making framework used in this domain, and present our formulations that implicitly assume the existence of reciprocal interactions between risks and services in disclosure control.

Part I of this dissertation is organized as follows. Chapter 2 introduces the privacy versus utility problem in disclosure control and provides an extensive survey of currently known solution techniques for solving this problem. This chapter also discusses the multi-objective nature of the problem and presents our decision making model based on multi-criteria analysis [55, 57]. Chapter 3 continues the discussion in the context of a data modification representation widely used by current techniques [59]. Chapter 4 extends the multi-criteria analysis by enabling the inclusion of data publisher preferences in the optimization process [53, 65]. Inclusion of these preferences allows a data publisher to focus on solutions that meet certain pre-specified requirements in terms of risk mitigation and quality achievement. Chapter 5 discusses the methodology that should be adopted for a comparative study in this domain, given that an algorithm cannot cater to both the privacy and utility objectives simultaneously [58]. Chapter 6 collects our observations and presents a unified framework to perform multi-criteria decision making in data anonymization [61]. Chapter 7 looks at some of the privacy issues in the management of information originating from the ubiquitous usage of mobile services [60, 62].

Part II: Security Risk Management

Networked systems constantly run under the risk of compromise. The security in these systems is only as good as the availability of known exploits and corresponding mitigation controls. Modern systems also have an extensive degree of inter-connectivity

that complicates the identification of contributions made by an existing vulnerability towards system compromise. Significant research has therefore gone into the formulation of security models for networked systems using paradigms like attack graphs and attack trees. These models help identify the cause-consequence relationship between system states, and enumerate the different attack paths that can lead to a compromised state. They also allow a system administrator to identify the minimum set of control measures necessary to prevent known exploits.

While risk assessment is vital to the security of any networked system, risk mitigation is often constrained by the availability of sufficient resources. Efficiency in risk mitigation is therefore also dependent on what control measures are chosen within given cost constraints. Security risk management encompasses this decision making paradigm where resource availability impacts the extent of risk to which a networked system will always be exposed. However, existing techniques deviate from this core principle and assume that the system administrator has the resources required to effectuate a "completely secure" system. We argue that this is an impractical assumption and the decision to achieve a certain level of security can only be made after obtaining a comprehensive understanding of the risks–resources trade-offs.

Part II of this dissertation is organized as follows. Chapter 8 looks at how security hardening is typically approached in existing works, and highlights the requirement for a multi-criteria analysis [51]. This analysis is crucial to any organization that operates under tight budget constraints, but at the same time, must make a best effort to protect its network assets. Chapter 9 discusses a similar problem in the domain of pervasive systems, where resource constraints are imposed by the heterogeneous nature of the environment [52]. Chapter 10 discusses how the hardening process can be made more robust by including attack probabilities in a cause-consequence model. These probabilities indicate an attack's difficulty level and are used to identify system states that have a high likelihood of compromise. Chapter 11 introduces the evolving attacker model and positions the efficacy of the static approach to security risk management under constantly changing attacker-defender dynamics.

Part III: Wireless Data Broadcasting

Fast access to information is becoming increasingly vital to support today's mobile infrastructures. As mobile devices gain more and more compute power, the variety of applications that can be supported by these devices are also becoming diverse. One of the implications of this trend for modern information management systems is to have the ability to provide clients fast access to the vast repository of data over a wireless medium. Challenges in doing so arise from the fact that wireless bandwidth is a limited resource. Wireless data broadcasting is becoming a popular mechanism in this scenario due to its scalability properties. Broadcasting reduces the amount of data to be transferred over a wireless channel when multiple clients are interested in the same information. Extensive research have therefore been carried out to determine optimal broadcast schedules satisfying a variety of access time constraints.

Since the broadcast schedules are built in real time, the QoS criteria must be well represented in order to minimize access delays and maximize resource utilization. The QoS criteria must encapsulate the possibility that access time constraints cannot always be met, in which case, the data should be made available based on its possible utility. We shall look at a practical perception of information availability in terms of the utility derived from it by the consumers. A number of utility driven optimization problems are explored here.

Part III of this dissertation is organized as follows. Chapter 12 discusses the design of broadcast schedulers that attempt to maximize the utility of broadcast data given the bandwidth limitations of a system [56]. Chapter 13 considers the additional constraint of ordering in the data items [63, 64]. Chapter 14 introduces a stochastic variant of the broadcasting problem in the context of a non-local data storage model [54].

Part I

Disclosure Control

CHAPTER 2

Managing Data Privacy and Utility

Various scientific studies, business processes and legal procedures depend on quality data. Large companies have evolved whose sole business is gathering data from various sources, building large data repositories, and then selling the data or their statistical summary for profit. Examples of such large data publishers are credit reporting agencies, financial companies, demographic data providers and so on. These data repositories often contain sensitive personal information, including medical records and financial profiles, which if disclosed and/or misused, can have alarming ramifications. Thus, not only the storage of this data done with strong security controls, but the dissemination is also frequently governed by various privacy requirements and subjected to disclosure control. For privacy protection, the data need to be sanitized of personally identifying attributes before it can be shared. Anonymizing the data, however, is quite challenging. Re-identifying the values in sanitized attributes is not impossible when other publicly available information or an adversary's background knowledge can be linked with the shared data. A classic example of such *linking attacks* was demonstrated by Sweeney [163], in which the author used a readily purchased voter list to re-identify medical records. In fact, a recent study on the year 2000 census data of the U.S. population reveals that 53% of the individuals can be uniquely identified by their gender, city and date of birth; 63% if the ZIP code is known in addition [76].

Database researchers have worked hard over the past several years to address such privacy concerns. Earlier techniques such as scrambling and adding noise to the data values [5] address the inference problem in statistical databases without reducing the value of the data. More recently, Samarati and Sweeney proposed the concept of *k–anonymity* to address the privacy problem. *k*–anonymity reduces the chances of a linking attack being successful [151, 152, 165]. The anonymization process involves transforming the original data set into a form unrecognizable in terms of the exact data values by using *generalization* and *suppression* schemes. A generalization performs a one-way mapping of the values of personally identifiable attributes, called *quasi-identifiers*, to a form non differentiable from the original values or to a form that induces uncertainty in recognizing them. An example of this is replacing a specific age by an age range. More often than not, it may be impossible to enforce a chosen level of privacy due to the presence of outliers in the data set. Outliers are not pre-defined in a given data set. Rather they depend on the generalization scheme that one is applying on the data. Given a particular generalization, outliers may emerge, making it difficult to achieve a desired level of privacy. In such a situation, a suppression scheme gets rid of the outliers. Suppression works by removing entire tuples making them no longer existent in the data set. A transformed data set of this nature is said to be *k*–anonymous if each record in it is same as at least $k - 1$ other records with respect to the quasi-identifiers. The higher the value of k, the stronger the privacy that the model offers.

An unavoidable consequence of performing such anonymization is a loss in the quality of the information content of the data set. Statistical inferences suffer as more and more diverse data are recoded to the same value, or records are deleted by a suppression scheme. A summary statistic relying on accurate individual information automatically deteriorates when stronger privacy is implemented. Researchers have therefore looked at different methods to obtain an optimal anonymization that results in a minimal loss of information [16, 89, 92, 150, 164, 178]. Since deciding on an anonymization for *k*–anonymity is NP-hard [127], most studies so far have focused on algorithms to minimize the information loss for a fixed value of k.

As research in this front progressed, other types of attacks have also been identified

— *homogeneity attack, background knowledge attack, skewness attack* and *similarity attack* [112, 120]. Models beyond *k*–anonymity have been proposed to counter these new forms of attacks on anonymized data sets and the hidden sensitive attributes. Two of the more well known models in this class are the *ℓ–diversity* model [120] and the *t–closeness* model [112]. While these models enable one to better guarantee the preservation of privacy in the disseminated data, they still come at a cost of reduced quality of the information.

The (possibly) unavoidable loss in data quality due to anonymizing techniques presents a dilemma to the data publisher. Since the information they provide forms the basis of their revenue, its whole purpose would be lost if the privacy controls prohibit any kind of fruitful inferences being made from the distributed data. In other words, although the organization needs to use some anonymization technique when disseminating the data, it also needs to maintain a pre-determined level of utility in the published data. Proper anonymization thus involves weighing the risk of publicly disseminated information against the statistical utility of the content. In such a situation, it is imperative that the data publisher understands the implications of setting a parameter in a privacy model (for example, k in k–anonymity or ℓ in ℓ–diversity) to a particular value. There is clearly a trade-off involved. Setting the parameter to a "very low" value impacts the privacy of individuals in the database. Picking a "very high" value disrupts the inference of any significant statistical information from the anonymized data set. Furthermore, a data publisher may at times be confronted with a choice of several values of a parameter. This will arise in situations where individuals are allowed an opportunity to specify their desired privacy levels. For example, some users may be content with $k = 2$ (in the k–anonymity model) while others may want $k = 4$. In such cases, the publisher needs to determine if some higher parameter value than initially selected is (or is not) possible with the same level of information loss. If a higher value is possible it will do a bonafide service to the individuals whose personal data are in the repository.

We believe that in order to understand the impact of setting the relevant parameters, a data publisher needs to answer questions similar to the following.

1. What level of privacy can one assure given that one may not suppress any record in the data set and can only tolerate an information loss of 25% (say)?

2. What is a good value for k (assuming the k–anonymity model) when one may suppress 10% (say) of the records and be able to tolerate an information loss of (maybe) 25%?

3. Under the "linking attacks" threat model and assuming that the users of the published data sets are likely to have background knowledge about some of the individuals represented in the data set, is it possible to combine the k–anonymity and the ℓ–diversity models to obtain a generalization that protects against the privacy problems one is worried about?

4. Is there a generalization that gives a high k and a high ℓ value if one is ready to suppress (maybe) 10% of the records and tolerate (say) 20% of information loss?

Unfortunately, answering these questions using existing techniques will require us to try out different k (or ℓ) values to determine what is suitable. Additionally, since the k–anonymity and ℓ–diversity models have been developed to address different types of attacks on privacy, one may want to combine the two models. This will require more possibilities to be tried out. Further, such a methodology does not guarantee that better privacy results cannot be obtained without incurring any or an acceptable increase in the information loss. Although recent studies have looked into the development of fast algorithms to minimize the information loss for a particular anonymization technique with a given value for the corresponding parameter, very few of them explore the data publisher's dilemma – given an acceptable level of information loss, determine the best k and/or ℓ value that satisfy the privacy requirements of the data set.

This chapter introduces the multi-objective nature of data anonymization and proposes the requisite formulations to address the data publisher's dilemma. First, we discuss the formulation of a series of multi-objective optimization problems, the solutions to which provide an in-depth understanding of the trade-off present between the level of privacy and the quality of the anonymized data set. We note that one important feature often overlooked in the specification of a privacy model is the distribution of the privacy parameter across the anonymized data set. The privacy parameter reported on an anonymized data set, for example k in k-anonymity, is a quantifier of the least property

satisfied by all tuples in the data set. Very often, this quantity is an inexact characterization of the privacy level, for it may be the case that a majority of the tuples in the data set actually satisfy a higher privacy property – a higher k value for example. Failure to capture the distribution of the parameter values thereby makes differentiating between two equivalent (in the sense of a privacy model parameter value and utility) anonymizations a difficult task. Techniques are therefore required to capture (or specify) this distribution and make it a part of the optimization process. As our second contribution, we build on the concept of *weighted-k anonymity* and include it in one of the multi-objective problem formulations. Third, we provide an analytical discussion on the formulated problems to show how information on this trade-off behavior can be utilized to adequately answer the data publisher's questions. Fourth, we exemplify our approach by using a popular evolutionary algorithm to solve the multi-objective optimization problems relevant to this study. Our last contribution is the design of a multi-objective formulation that can be used to search for generalizations that result in acceptable adherence to more than one privacy property within acceptable utility levels. Towards this end, we show how decision making is affected when trying to use the k–anonymity and ℓ-diversity models simultaneously.

The remainder of the chapter is organized as follows: Section 2.1 reviews some of the existing research in disclosure control. The required background on multi-objective optimization is presented in Section 2.2. We introduce the terminology used in the chapter in Section 2.3. Section 2.4 provides a description of the four multi-objective problems we formulate and the underlying motivation behind them. The specifics of the solution methodology with respect to solving the problems using an evolutionary algorithm is given in Section 2.5, and a discussion of the results so obtained is presented in Section 2.6. Finally, Section 2.7 summarizes and concludes the chapter.

2.1   Related Work

Several algorithms have been proposed to find effective k–anonymization. The μ-argus algorithm is based on the greedy generalization of infrequently occurring combinations of quasi-identifiers and suppresses outliers to meet the k–anonymity requirement [89].

μ-argus suffers from the shortcoming that larger combinations of quasi-identifiers are not checked for *k*–anonymity and hence the property is not always guaranteed [164].

Sweeney's *Datafly* approach uses a heuristic method to generalize the attribute containing the most distinct sequence of values for a provided subset of quasi-identifiers [164]. Sequences occurring less than *k* times are suppressed. Samarati's algorithm [150] can identify all *k*–minimal generalizations, out of which an optimal generalization can be chosen based on certain preference information provided by the data recipient. A similar *full-domain generalization* is also proposed in *Incognito* [109]. The basic Incognito algorithm starts with the generalization lattice of a single attribute and performs a modified bottom-up breadth-first search to determine the possible generalized domains of the attribute that satisfy *k*–anonymity. Thereafter, the generalization lattice is updated to include more and more number of attributes.

Iyengar proposes a flexible generalization scheme and uses a genetic algorithm to perform *k*–anonymization on the larger search space that resulted from it [92]. Although the method can maintain a good solution quality, it has been criticized for being a slow iterative process. In this context, Lunacek et al. introduce a new crossover operator that can be used with a genetic algorithm for constrained attribute generalization, and effectively show that Iyengar's approach can be made faster [118]. As another stochastic approach, Winkler proposes using simulated annealing to do the optimization [178].

On the more theoretical side, Sweeney proposes the *MinGen* algorithm [164] that exhaustively examines all potential generalizations to identify the generalization that minimally satisfies the anonymity requirement, acknowledging that the approach is impractical even on modest sized data sets. Meyerson and Williams have recently proposed an approximation algorithm that achieves an anonymization with $O(k \log k)$ of optimal [127]. However, the method is not suitable when larger values of *k* is desired.

Most of the previous approaches start from the original data set and systematically or greedily generalize it into one that is *k*–anonymous. Bayardo and Agrawal propose a complete search method that iteratively constructs less generalized solutions starting from a completely generalized data set [16]. The algorithm starts with a fully generalized data set and systematically specializes it into one that is minimally *k*–anonymous. It uses a tree

search strategy exploiting both cost-based pruning and dynamic search rearrangement [17]. The idea of a "solution cut" is presented by Fung et al. in their approach to top down specialization [72]. A generalization is visualized as a "cut" through the taxonomy tree of each attribute. A cut of a tree is a subset of values in the tree that contains exactly one value on each root-to-leaf path. A solution cut is a cut that satisfies the anonymity requirement.

A more general view of k-anonymization is clustering with a constraint on the minimum number of objects in every cluster [6, 32, 67, 111]. Clustering techniques use metrics that quantify the distance between tuples and distance between equivalence classes. The basic idea for the algorithm is to find an arbitrary equivalence class of size smaller than k and merge it with the closest equivalence classes to form a larger equivalence class with the smallest distortion. The process is repeated recursively until each equivalence class contains at least k tuples. Minimum distortion is enforced by choosing the closest common generalizations of attributes.

LeFevre et al. extend the notion of generalizations on attributes to generalization on tuples in the data set [110]. The authors argue that such multidimensional partitioning of the generalization domain show better performance in capturing the underlying multivariate distribution of the attributes, often advantageous in answering queries with predicates on more than just one attribute.

The drawbacks of using k–anonymity are first described by Machanavajjhala et al. [120]. They identify that k–anonymized data sets are susceptible to privacy violations when there is little diversity in the sensitive attributes of a k–anonymous equivalence class. In order to alleviate such privacy breaches, they propose the model of ℓ–diversity which obtains anonymizations with an emphasis on the diversity of sensitive attribute values on a k–anonymous equivalence class. Further work presented by Li et al. show that the ℓ–diversity model is also susceptible to certain types of attacks [112]. To this effect, they emphasize having the t–closeness property that maintains the same distribution of sensitive attribute values in an equivalence class as is present in the entire data set, with a tolerance level of t. They realize that t–closeness does not deal with identity disclosure scenarios and propose that it should be used in conjunction with k–anonymity.

The k–anonymity model assumes an adversary with full knowledge about the public information (quasi-identifiers) of all individuals in the microdata. This is an unrealistic assumption and is often not required. Variations have thus been proposed with relaxations on the knowledge of an adversary. If an adversary knows the public information of a single individual, then we may generalize the table so that the original public data of every individual maps to the generalized public data of at least k tuples. This is a relaxation of the k–anonymity requirement since the k tuples are not required to form an equivalence class. Such an anonymization results in $(1,k)$–anonymity. Another notion is that of $(k,1)$–anonymity where any tuple in the anonymized data set maps to at least k tuples in the original data. When both forms of anonymity are satisfied, it is called (k,k)–anonymity [74]. Every k–anonymous table is (k,k)–anonymous, but the reverse is not necessarily true. (k,k)–anonymizations are secure if an adversary has knowledge on a limited number of individuals in the data set, but can be insecure if the adversary has full knowledge on all individuals. Quantified notions of adversarial knowledge is also used in *skyline privacy* [36]. A vector (ℓ,k,m) is used to say that the adversary knows ℓ sensitive values that a target individual does not have, the sensitive values of k individuals other than the target, and m individuals such that if any one has a specific sensitive value then the target also has it. Given such a representation of adversarial knowledge, a released table must guarantee multidimensional privacy such that the adversary's confidence that an individual has a certain sensitive value should not exceed a given threshold.

Combination of different privacy characteristics have also been attempted in certain models. The (α,k)–anonymity model extends k–anonymity such that the confidence of associating quasi-identifiers to sensitive values is limited to within α [181]. Under this model, any k–anonymization must also satisfy the *α-deassociation* requirement, i.e. the relative frequency of a given value of a sensitive attribute in an equivalence class must be less than or equal to a user-defined threshold α, where $0 < \alpha < 1$. However, the model has limitations similar to ℓ–diversity. When the frequency of sensitive values in the whole data set is not well-proportioned, the presence of some highly sensitive values with lower frequency will force α to be very close to one. Therefore a generic model, called the *complete (α,k)–anonymity,* uses different α values for different sensitive values [95].

Another model proposed to protect against attribute disclosure is *p–sensitive k–anonymity* [173]. A data set satisfies *p*–sensitive *k*–anonymity if it satisfies *k*–anonymity and the number of distinct values for each sensitive attribute is at least *p* in each equivalence class. *Extended p–sensitive k–anonymity* further enforces the requirement that no two values of a sensitive attribute in an equivalence class are descendants of a common protected node in a hierarchy tree specified over the sensitive attribute domain [34]. The hierarchy tree captures semantic relationships between possible values. A similar approach to combine *k*–anonymity and ℓ–diversity is adopted in the (k, ℓ)–*anonymity* model [113].

Most anonymity models focus on an universal approach to privacy where the same amount of preservation is sought for all individuals. As a consequence, such models may be offering insufficient protection to a subset of individuals while applying excessive privacy control to another subset. The notion of *personalized anonymity* eliminates such problems by performing generalizations that satisfy personal requirements [185]. Personal preferences on sensitive attributes are captured as "guarding nodes". For example, a personal preference may allow "flu" to be disclosed as the illness of a patient but no disclosure of a "lung cancer" patient or the inference that the illness is "cancer" may be allowed. In such cases, the released microdata must limit the probability of inference of information beyond what is allowed within a threshold.

A different approach to data generalization is adopted in *Anatomy* [184]. It is a data dissemination method designed on top of ℓ–diversity with the difference that quasi-identifiers are not generalized and released in their original form. The quasi-identifiers and the sensitive attribute are released in two different tables, called the *QIT* and *ST* respectively. The tuples in a microdata are first partitioned (without generalizing the quasi-identifiers) into groups such that the ℓ–diversity property holds in every group. QIT comprises of only the quasi-identifier values of the tuples along with the group number to which a tuple belongs. ST lists, for every group, the distinct values of the sensitive attribute in the group along with a count of the number of tuples in the group that has a specific value. Since no generalization is performed on the quasi-identifiers, anatomy preserves any distributions/correlations in the attributes. Enforcing the ℓ–diversity property while grouping the tuples protects ST against homogeneity and background knowledge

attacks. Besides such fragmentation, perturbation (addition of noise) is another technique that is used to anonymize data. Although data perturbation is slowly being discarded in microdata anonymization, these methods are still used in statistical disclosure control where answers to summary queries on a statistical database should not reveal individual values. However, under the light of recent results by Dwork and Yekhanin [68], the efficacy of the technique even for statistical disclosure control has become questionable.

Quantification of data utility has been approached from different perspectives by researchers. Early notion of information loss is based on the number of generalization steps one has to perform to achieve a given privacy requirement [150]. Such a method assumes that attribute domains can be progressively generalized and a partial order can be imposed on the domain of all generalizations for an attribute. For instance, ZIP codes can be generalized by dropping a digit from right to left at each generalization step. Postal addresses can be generalized to the street, then to the city, to the county, to the state, and so on. Given that such orderings can be imposed, a distance can be computed for each attribute between the microdata and a generalized version of it. The result is a distance vector with an entry for each attribute. Dominance relationships are used to determine if one distance vector is better than other – a better distance vector will have lower distance values for each attribute.

Information loss is also measured in terms of the amount of distortion in a generalized table. In a cell of a generalized table, the ratio of the domain of the value found in the cell to the height of the attribute's hierarchy reports the amount of generalization and thereby measures the cell's distortion [164]. *Precision* of a generalized table is then computed as one minus the sum of all cell distortions (normalized by the total number of cells). Generalizations based on attributes with longer generalization hierarchies typically maintain precision better than generalizations based on attributes with shorter hierarchies. Further, hierarchies with different heights can provide different precision measures for the same table. A similar distortion measure is also used in [111].

Similar estimation of information loss is used in the *general loss metric* [92]. The general loss metric computes a normalized information loss for each data value in the generalized data set. The requirement here is that information in every column is potentially impor-

tant and hence a flexible scheme to compute the loss for both numeric and categorical data is required. Consider an attribute containing categorical information that is generalized based on a hierarchy tree (e.g. Fig. 2.2). The generalized value of this attribute for a certain tuple corresponds to node P in the tree. If the total number of leaf nodes in the tree is M and the number of leaf nodes in the subtree rooted at node P is M_P, then loss for this data value is computed as $(M_P - 1)/(M - 1)$. Similarly, if U_i and L_i are the upper and lower bounds of the interval to which a numerical attribute value gets mapped to, the loss is $(U_i - L_i)/(U - L)$, where U and L are the upper and lower bounds of the attribute. The general loss is the sum of loss over all data values. In some attempts, a summation of losses computed by these methods (interval based and tree based) is instead used [32].

A widely used loss metric, called the *discernibility metric*, assigns a penalty to each tuple based on the number of tuples in the anonymized data set that are indistinguishable to each other [16]. Thus, a tuple belonging to an equivalence class of size j is assigned a penalty of j. A suppressed tuple is assigned a penalty equal to the number of tuples in the data set. The idea behind using the size of the equivalence class as a measure of information loss is to penalize generalizations that result in equivalence classes bigger than what is required to enforce a given privacy requirement. A variant of this is to use the *normalized average cluster size* [110].

Data utility is often measured in conjunction with privacy in an attempt to combine both objectives into a single metric. A metric of such nature favors generalizations that result in maximum gain in the information entropy for each unit of anonymity loss resulting from the generalization. Methods employing such a metric progressively increase the amount of generalization, called a *bottom up generalization* approach [176], or decrease it, called a *top down specialization* approach [72], with the objective of maximizing the metric without violating the anonymity requirement. Utility assessment in these methods are motivated by the ability to perform correct classification tasks as typical in data mining applications. A classification metric is also proposed by Iyengar where a tuple is penalized if it gets suppressed or of its class label does not match the majority class label [92].

Another metric, called *usefulness*, measures utility as the average diversity in the tuples

belonging to an equivalence class [117]. This measurement is similar to the general loss metric, with differences being in the treatment of interval based attribute domains. For such domains, the loss is assigned as the normalized distance between the maximum and minimum values of the attribute in the equivalence class. A complementary metric, called *protection*, uses the inverse of the tuple diversities as a measure of the privacy factor. The two metrics inherently exhibit a reciprocal relationship, useful when a data publisher wants to modulate the anonymization process towards one objective or the other.

Preliminary metrics to evaluate the effectiveness of anonymized data in answering aggregate queries have also been proposed. Quality here is derived from a normalized difference between the result of evaluating a query on the anonymous data and the result on the original data [110]. Given a totally ordered set of sensitive attribute values, these metrics measure the range of values that is encompassed in a query result [193]. The smaller the range, the higher is the query answering accuracy.

Loss metrics should not only capture the information loss caused by the generalization but also account for the importance of the different attributes. For example, given a disease analysis data set, an "age" attribute may be considered more critical than a "ZIP code" attribute. In such a case, generalizations that are able to maintain the "age" attribute more accurately should be favored. The *weighted normalized certainty penalty* metric uses a weighted sum of the loss measurements in different attributes of the data set [187]. The loss measurement is similar as in the general loss metric and the usefulness metric. The introduction of such preference characteristics indicates that the measurement of utility can be a very subjective matter after all.

The first known attempt of exploring the privacy and utility trade-offs is undertaken by Dewri et al. [55]. The work focuses on a multi-objective optimization formulation based on a model called weighted-k anonymity. A similar trade-off analysis is presented by Huang and Du in the problem of optimizing randomized response schemes for privacy protection [88].

The potential problem in using the above approaches is that they are targeted towards obtaining an optimal generalization for a fixed value of k, ℓ, or t, sometimes in conjunction. Besides running the algorithms multiple times with different values of the

parameter(s), no attempt is known to have been made to understand how the generalizations and the related cost metrics change with changes in the parameter values. Our work seeks to fill this gap.

2.2   Multi-objective Optimization

In real world scenarios, often a problem is formulated to cater to several criteria or design objectives, and a decision choice to optimize these objectives is sought for. An optimum design problem must then be solved with multiple objectives in consideration. This type of decision making falls under the broad category of multi-criteria, multi-objective, or vector optimization problems.

Multi-objective optimization differs from single-objective ones in the cardinality of the optimal set of solutions. Single-objective optimization techniques are aimed towards finding the global optima. In case of multi-objective optimization, there is no such concept of a single optimum solution. This is due to the fact that a solution that optimizes one of the objectives may not have the desired effect on the others. As a result, it is not always possible to determine an optimum that corresponds in the same way to all the objectives under consideration. Decision making under such situations thus require some domain expertise to choose from multiple trade-off solutions depending on the feasibility of implementation.

Formally we can state a multi-objective optimization problem (MOOP) in microdata disclosure control (MDC) as follows.

Definition 2.1 MDC MOOP. Let $f_1, \ldots, f_M$ denote M objective functions to maximize while performing a modification of a given table T. Find a generalized table T' of T which optimizes the M-dimensional vector function

$$f(\mathsf{T}^*) = [f_1(\mathsf{T}^*), f_2(\mathsf{T}^*), \ldots, f_M(\mathsf{T}^*)], \tag{2.1}$$

where T^* is a generalized version of T.

The objective functions in this case are either related to the privacy or utility level maintained in an anonymized table. Note that the privacy level can be inferred with

respect to different privacy models. Hence the number of objectives can be more than two. In order to find an optimal solution to the MDC MOOP, we must be able to compare anonymizations with respect to all the objectives in hand. However, due to the conflicting nature of the objective functions, a simple objective value comparison between two anonymizations cannot be performed. Most multi-objective algorithms thus use the concept of dominance to compare feasible solutions.

Definition 2.2 DOMINANCE AND PARETO-OPTIMAL SET. Given a table T and M objectives to maximize, a generalized table T_1 of T is said to dominate another generalized table T_2 of T if

1. $\forall i \in \{1,2,\ldots,M\}$ $f_i(\mathsf{T}_1) \geq f_i(\mathsf{T}_2)$, and

2. $\exists j \in \{1,2,\ldots,M\}$ $f_j(\mathsf{T}_1) > f_j(\mathsf{T}_2)$.

T_2 is then said to be dominated by T_1, denoted by $\mathsf{T}_2 \preceq \mathsf{T}_1$. If the two conditions do not hold, T_1 and T_2 are said to be non-dominated w.r.t. each other, denoted by the $\npreceq$ symbol. Further, all generalized tables of T which are not dominated by any possible generalized version of T constitutes the Pareto-optimal set.

In other words, a Pareto-optimal solution is as good as other solutions in the Pareto-optimal set, and not worse than other feasible solutions outside the set. The surface generated by these solutions in the objective space is called the *Pareto-front* or *Pareto-surface*. Fig. 2.1 shows the Pareto-front for a hypothetical two-objective problem, with the dominance relationships between three feasible solutions.

In the context of the *k*–anonymity problem, the Pareto-front for the two objectives – maximize *k* and minimize *loss* – provides the decision maker an understanding of the changes in the information loss when *k* is varied. Consider two anonymized versions T_1 and T_2 of a data set, with corresponding *k* and *loss* as $(k_1, loss_1)$ and $(k_2, loss_2)$ respectively. Let us assume that $k_1 < k_2$ and $loss_1 = loss_2$. A decision maker using T_1, and unaware of T_2, misses on the fact that a higher *k* value is possible without incurring any increase in the loss. A multi-objective algorithm using the dominance concept can expose this relationship between T_1 and T_2, namely $\mathsf{T}_1 \preceq \mathsf{T}_2$. As another example, consider the

Figure 2.1: Pareto-front for a hypothetical two-objective problem.

case with $loss_2 - loss_1 = \epsilon > 0$. T_1 and T_2 are then non-dominated solutions, meaning that one objective cannot be improved without degrading the other. However, if ϵ is a relatively small quantity acceptable to the decision maker, T_2 might be preferable over T_1. Such trade-off characteristics are not visible to the decision maker until a multi-objective analysis is carried out. Thus, the objective of the analysis is to find the Pareto-optimal set from the set of all possible anonymized versions of a given data set.

The classical way to solve a multi-objective optimization problem is to follow the weighted-sum approach [172]. Many methods following this approach employ a scalarization of the multiple objectives at hand [129, 130]. A relative weight vector for the objectives can help reduce the problem to a single-objective instance, or impose orderings over the preference given to different objectives. However, such methods fail to provide a global picture of the choices available to the decision maker. In fact, the decision of preference has to be made before starting the optimization process. Relatively newer methods have been proposed to make the decision process more interactive.

Evolutionary algorithms for multi-objective optimization (EMO) have been extensively studied and applied to a wide spectrum of real-world problems. One of the major advantages of using evolutionary algorithms is their ability to scan through the global search space simultaneously, instead of restricting to localized regions of gradient shifts. An EMO works with a population of trial solutions trying to converge on to the Pareto-

optimal set by filtering out the infeasible or dominated ones. Having multiple solutions from a single run of an EMO is not only an efficient approach but also helps a decision maker obtain an intuitive understanding of the different trade-off options available at hand. The effectiveness of an EMO is thus characterized by its ability to converge to the true Pareto-front and maintain a good distribution of solutions on the front [81, 105].

A number of algorithms have been proposed in this context [40, 46] – NPGA [85], DPGA [139], PAES [102], SPEA2 [195] and NSGA-II [48], to name a few widely referred ones. We employ the Non-dominated Sorting Genetic Algorithm-II (NSGA-II) for the multi-objective optimization in this study. NSGA-II has gained wide popularity in the multi-objective optimization community, partly because of its efficiency in terms of the convergence and diversity of solutions obtained, and partly due to its extensive application to solve real-world problems[1]. However, we would like to highlight that the availability of an algorithm is not sufficient to apply it directly in this problem domain. As in many real world applications, our contribution comes in the form of appropriate formulations of the problem so that the algorithm can be applied.

2.3   Preliminaries

A data set D can be visualized as a tabular representation of a multi-set of tuples $r_1, r_2, \ldots, r_{n_{row}}$ where n_{row} is the number of rows in the table. Each tuple (row) r_i comprises of n_{col} values $\langle c_1, c_2, \ldots, c_{n_{col}} \rangle$ where n_{col} is the number of columns in the table. The values in column j correspond to an attribute a_j, the domain of which is represented by the ordered set $\Sigma_j = \{\sigma_1, \sigma_2, \ldots, \sigma_{n_j}\}$. The ordering of elements in the set can be implicit by nature of the data. For example, if the attribute is "age", the ordering can be done in increasing order of the values. For categorical data, obtaining an ordering requires the user to explicitly specify a hierarchy on the values. A hierarchy can be imposed based on how the values for the attribute can be grouped together. Fig. 2.2 shows an example hierarchy tree for the attribute "marital status". The leaf nodes in this example constitute

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[1]*ISI Essential Science Indicators* fast breaking paper in engineering for February 2004: http://www.esi-topics.com/fbp/fbp-february2004.html

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Figure 2.2: Hierarchy tree for the *marital status* attribute. Numbering on the leaf nodes indicate their ordering in $\Sigma_{marital\ status}$.

the actual values that the attribute can take. The ordering for these values can be assigned based on the order in which the leaf nodes are reached in a pre-order traversal of the hierarchy tree [92].

A generalization G_j for an attribute a_j is a partitioning of the set Σ_j into ordered subsets $\langle \Sigma_{j_1}, \Sigma_{j_2}, \ldots, \Sigma_{j_p} \rangle$ which preserves the ordering in Σ_j, i.e. if σ_a appears before σ_b in Σ_j then, for $\sigma_a \in \Sigma_{j_l}$ and $\sigma_b \in \Sigma_{j_m}$, $l \leq m$. Further, every element in Σ_j must appear in exactly one subset and the elements in the subsets maintain the same ordering as in Σ_j. For the "age" attribute having values in the range of $[10, 90]$, a possible generalization can be $\langle \{[10, 30]\}, \{(30, 50]\}, \{(50, 70]\}, \{(70, 90]\} \rangle$. A possible generalization for the "marital status" attribute can be $\langle$*{Not Married}, {spouse-absent}, {civ-spouse}, {AF-spouse}*$\rangle$. It is important to note that generalizations for categorical data is dependent on how the hierarchy is specified for it. Further, generalizations are restricted to only those that respect the hierarchy. The generalization is said to be *constrained* in such a case. For example, the generalization $\langle$*{Never Married, Divorced}, {Widowed, Separated}, {Married}*$\rangle$ is not valid for "marital status" since the hierarchy tree specifies that the values *{Divorced, Widowed, Separated}* can only be generalized as *Once-Married*, if at all.

Note that the number of partitions (or groups) of an attribute domain, i.e. P, signifies the extent of generalization that will be performed for the attribute. If $P = 1$ then all values of the attribute will be represented by the same subset, in which case all information in

that attribute is lost. On the other extreme, if $P = |\Sigma_j|$ for attribute a_j then every value will map to its own unique subset (no generalization) and all information in the attribute will be maintained in the original form.

Given the generalizations $G_1, G_2, \ldots, G_{n_{col}}$, the data set D can be transformed to the anonymized data set D' by replacing each value v at row i and column j in D by $G_j(v)$, where $G_j(v)$ gives the index of the subset to which v belongs to in the generalization G_j. Note that if a particular generalization G_j is equal to the domain of values Σ_j, all values of the corresponding attribute will be transformed to the same subset index 1, in which case all information in that attribute is lost and the cell is *suppressed.*

2.3.1 k–Anonymity

Tuples in D whose subset indices are equal in every column of D' can be grouped together into *QI-groups* or *equivalence classes*. In other words, a QI-group collects all tuples in D that has the same generalized form for the quasi-identifiers. The k–anonymity problem is then defined as follows.

Definition 2.3 k–ANONYMITY PROBLEM. Given a data set D, find a set of generalizations for the attributes in D such that the QI-groups induced by anonymizing D using the generalizations are all of size at least k.

The problem can also be explained as obtaining the generalizations under which every tuple in D' is same as at least $k - 1$ other tuples. Thus, a higher value of k evaluates to a lower chance of privacy breach.

2.3.2 ℓ–Diversity

The set of attributes can be divided into *sensitive* and *non-sensitive* classes. A sensitive attribute is one whose value must not be revealed (or get revealed) for any tuple in the data set. All other attributes are considered non-sensitive. Then, the ℓ–diversity principle says that every QI-group should contain at least ℓ "well-represented" values for a sensitive attribute [120]. The principle can be instantiated in many different forms depending on the meaning of "well-represented".

Let a_s be a sensitive attribute in a data set with the domain of values $\Sigma_s = \{\sigma_1, \sigma_2, \ldots, \sigma_{n_s}\}$. Further, let $Q_1, \ldots, Q_p$ be the QI-groups induced by a generalization. If $c(\sigma)_j$, where $\sigma \in \Sigma_s$, denotes the count of the number of tuples with the sensitive attribute value σ in Q_j, then one possible instantiation of the ℓ–diversity problem can be stated as follows.

Definition 2.4 ℓ–DIVERSITY PROBLEM. Given a data set D, find a set of generalizations for the attributes in D such that for each QI-group induced by anonymizing D using the generalizations, the relation

$$\frac{c(\sigma)_j}{|Q_j|} \leq \frac{1}{\ell} \tag{2.2}$$

holds for all $\sigma \in \Sigma_s$ and $j = 1, \ldots, p$.

In other words, the ℓ–diversity property guarantees that a sensitive attribute value cannot be associated with a particular tuple with a probability more than $1/\ell$. The higher the value of ℓ, the better is the privacy. Although this instantiation underlines the essence of the ℓ–diversity principle, two other formulations have been suggested in the original work. One of them is to assure that the entropy of each QI-group, defined as

$$Entropy(Q_j) = -\sum_{\sigma \in \Sigma_s} \frac{c(\sigma)_j}{|Q_j|} \log\left(\frac{c(\sigma)_j}{|Q_j|}\right), \tag{2.3}$$

is at least $\log \ell$. This requires that the entropy of the whole table is at least $\log \ell$ which may be difficult to ensure when a few values appear more frequently than others. Hence, another instantiation makes sure that the most frequent sensitive values do not appear too frequently in a QI-group and the less frequent values do not occur rarely. This is called *recursive (c, ℓ)–diversity*. If m is the number of sensitive attribute values in a QI-group and r_i is the count of the ith most frequent value in the QI-group, then the QI-group is said to be recursive (c, ℓ)–diverse if $r_1 < c(r_\ell, r_{\ell+1}, \ldots, r_m)$. Every QI-group must be recursive (c, ℓ)–diverse for a table to have recursive (c, ℓ)–diversity. The nature of the instantiation is not the focus of this work but the parameters involved in it. We shall use the instantiation given in Def. 2.4 to demonstrate our methodology.

2.3.3 (k, ℓ)–Safe

At this stage, we introduce the concept of a (k, ℓ)–*safe* anonymization. Along the lines of models such as general (α, k)–anonymity [181] and p–sensitive k–anonymity [173], a (k, ℓ)–safe data set incorporates the benefits of k–anonymity and ℓ–diversity in an anonymization. From a data publisher's perspective, it is meaningful since it allows one to ascertain the presence of the privacy guards obtainable from both k–anonymity and ℓ–diversity. A high k value in this case prohibits the likelihood of linking attacks, while a high ℓ value prohibits the likelihood of homogeneity and background knowledge attacks. When multiple types of attacks are possible on a data set, a data publisher would most certainly want to safeguard the published data against as many of them as possible.

Definition 2.5 (k, ℓ)–SAFE. An anonymized data set is (k, ℓ)–safe if it is k–anonymous and ℓ–diverse.

The distinction between (α, k)–anonymity and (k, ℓ)–safety lies in the representation of attribute disclosure risks. (α, k)–Anonymity requires that a possible value of a sensitive attribute should appear with a relative frequency not greater than α in a QI-group, also called the α-deassociation property. The motivation behind this representation is to prohibit the frequent occurrence of certain highly sensitive values in a QI-group, such as "HIV" in a disease data set, and hence control the inference probability on these values. (k, ℓ)–Safe, on the other hand, uses an instantiation of ℓ–diversity that specifies the minimum number of distinct values of a sensitive attribute that must appear in a QI-group. This is very much similar to p–sensitive k–anonymity. Nonetheless, ℓ–diversity can very well be instantiated in conformation to α-deassociation or any of the other forms mentioned in the original work. Hence, we use the generic name (k, ℓ)–safe.

2.3.4 Optimal generalization

The trivial generalization $G_j = \langle \Sigma_j \rangle$, where $j = 1, \ldots, n_{col}$, can provide the highest k and the highest ℓ value. However, such a generalization results in an anonymized data set with little or no statistically significant information. It is often desired that the anonymized data set be useful for some level of statistical analysis. In such a case, the

decision on the value of k (or ℓ) is subjected to a loss measurement of the information content in the anonymized data set, usually done using some metric. An optimization problem defined for a given value of k (or ℓ) tries to find the generalizations that result in a minimal loss given by the metric.

Depending on the distribution of data values in a data set, obtaining a generalization with an acceptable loss for a given value of k (or ℓ) may or may not be possible. This happens when the data set has outliers that cannot be anonymized without overly generalizing the remaining data points. Therefore, it becomes a requirement that such outliers be suppressed completely in order to avoid an over-generalization. A suppressed tuple is usually considered nonexistent in the data set. The loss metric can account for this suppression in its loss measurement.

When suppression is allowed, an anonymized data set can be made k–anonymous by suppressing all tuples that belong to QI-groups of size less than k. Similar suppression methods can be used to enforce the ℓ–diversity property. The case without suppression can be modeled into the earlier scenario (with suppression) by assigning an infinite loss when suppression is performed [16]. However, it should be noted that the presence of outliers will always force the requirement for suppression, in which case the loss measurement will always become infinite. Furthermore, even though suppression is not allowed, such an approach enforces the k–anonymity (or ℓ–diversity) property by suppressing outliers. If all the data points in the data set must stay in the anonymized data set as well, the desired privacy properties cannot be ascertained even after adopting such modeling.

We now proceed to formulate a set of four problems, the solutions to which provide an in-depth understanding of the publisher's dilemma. This is accompanied by an example that illustrates the scope of the problem and the corresponding analysis we envisage to help alleviate the problem.

2.4   Problem Formulation

As stated in the previous section, an optimization algorithm requires a numeric representation of the information loss associated with a particular generalization. A quantified loss value enables the optimization algorithm to compare two generalizations for their

relative effectiveness. Loss (cost) metrics assign some notion of penalty to each tuple whose data values get generalized or suppressed, thereby reflecting the total information lost in the anonymization process. In this work, we use the general loss metric proposed by Iyengar [92]. The general loss metric computes a normalized information loss for each of the data values in an anonymized data set. Note that our use of the general loss metric is only a matter of choice. The optimization process we use is a black box method and is not affected by how information loss is measured.

2.4.1 Generalization loss

Consider the data value $v_{i,j}$ at row i and column j in the data set D. The general loss metric assigns a penalty to this data value based on the extent to which it gets generalized during anonymization. Let $g_{i,j} = G_j(v_{i,j})$ be the index of the subset to which $v_{i,j}$ belongs in the generalization G_j, i.e. $v_{i,j} \in \Sigma_{j_{g_{i,j}}}$. The penalty for information loss associated with $v_{i,j}$ is then given as follows.

$$loss(v_{i,j}) = \frac{|\Sigma_{j_{g_{i,j}}}| - 1}{|\Sigma_j| - 1} \tag{2.4}$$

For categorical data, the loss for a cell is proportional to the number of leaf nodes rooted at an internal node (the generalized node) of the hierarchy tree. The loss attains a maximum value of one when the cell is suppressed ($G_j = \langle \Sigma_j \rangle$), or in other words, when the root of the tree is the generalized node. Subtracting one ensures that a non-generalized value incurs zero loss since the cardinality of the subset to which it belongs would be one. The generalization loss is then obtained as the total loss over all the data values in the data set.

$$GL = \sum_{i=1}^{n_{row}} \sum_{j=1}^{n_{col}} loss(v_{i,j}) \tag{2.5}$$

2.4.2 Suppression loss

Although the loss due to suppression can be incorporated into the generalization loss, we decide to separate it out for the purpose of our study. When a row is suppressed, all cells in the row are suppressed irrespective of the generalization. Each cell thereby incurs

a loss of one (consequence of Eq. (2.4)). Let n_{sup} be the number of rows to be suppressed in the data set. The suppression loss for the data set is then given as

$$SL = n_{col} \times n_{sup}. \tag{2.6}$$

2.4.3 The multi-objective problems

The multi-objective problems we formulate in this chapter are intended to analyze and understand the trade-off nature of the generalization and suppression losses when k (or ℓ) is varied. A single objective optimization problem to minimize the generalization loss with a fixed k (or ℓ) will require multiple runs of the algorithm to understand this trade-off. By adopting a multi-objective approach, we can generate a fairly good approximation of the Pareto-front in a single run of the algorithm, which in turn provides us with the requisite information to make a better decision on the choice of k (or ℓ) for anonymization. In this context, we formulate a series of multi-objective problems for our analysis. Although, Problems A, B, and C are described for the k–anonymity problem, similar analysis can be carried out for ℓ–diversity as well. Problem D caters to (k, ℓ)–safety.

Also note that the problems under study are not intended to provide the data publisher a "best" value for the parameter(s) involved in the anonymization technique. Rather, we put forward a methodology to understand the implications of choosing a particular value for the parameter(s) in terms of the resulting privacy and the data utility. Hence, we shall often find that one or more solutions returned by the optimization process are trivially not acceptable either in terms of privacy or utility, or in some cases, both. It is not our objective to consider such solutions as degenerate and prohibit them from appearing in the solution set. After all, they are also a manifestation of the privacy-utility trade-off, which would likely be never selected as a choice by the data publisher, but still possible. For example, an extreme solution will correspond to a situation where every tuple in the data set belongs to its own QI-group, thereby resulting in no privacy and maximum utility. Another extremity is the case where all tuples are grouped together in a single QI-group resulting in maximum privacy but no utility. One cannot deny the fact that in the case of privacy versus utility, both of these are possible solutions. The multi-objective

optimization formulations do not incorporate the required domain knowledge to identify these extremities (or other such solutions) as being impractical. Only the data publisher has the requisite knowledge to make such identification and disregard such solutions. This is often a post-optimization process. Hence, the focus in the solution set should be concentrated on the practical solutions reported by the method. Quite often there will be more than one, and the methodology provides the data publisher a distinctive picture of the differences arising between privacy and utility when it makes a decision to choose one solution over another.

2.4.3.1   Problem A: Zero suppression

The presence of outliers in a data set makes it difficult to find a suitable value of k when suppression of data is not allowed. In the first problem formulation, we strictly adhere to the requirement that no tuple in the data set can be deleted. Intuitively, such a strict requirement makes the k–anonymity problem insensible to solve for a given k as the optimization algorithm will be forced to overly generalize the data set in its effort to ensure k–anonymity. The outliers usually belong to very small QI-groups and the only way to merge them into a bigger one is by having more generalization.

Although solving the k–anonymity problem is not possible in terms of its strict definition, it is worth noting that a generalization can still affect the distribution of the size of the QI-groups even when suppression is not allowed.

Let us define an equi-privacy class E_i as the set of all tuples in QI-groups of size i. Hence an arbitrary equi-privacy class E_i shall contain all tuples that are i–anonymous. For brevity, we shall use the term "small equi-privacy class" or "equi-privacy class with lower i" to mean equi-privacy classes E_is with relative smaller i than that used in the term "large equi-privacy class" or "equi-privacy class with high i". An ideal generalization would then maintain an acceptable level of loss and also keep the number of rows in equi-privacy classes with lower i relatively fewer than in equi-privacy classes with higher i. Although this does not guarantee complete k–anonymity, the issue of privacy breach can be solved to a limited extent by reducing the probability that a randomly chosen row would belong to a small QI-group.

With this motivation, we define the weighted-k–anonymity multi-objective problem to find generalizations that produce a high weighted-k value and low generalization loss. Each equi-privacy class E_i defines a privacy level for its member tuples – every tuple in the equi-privacy class is the same as exactly $i - 1$ other tuples in the same class and has a probability of breach equal to $1/i$. For an intuitive comparison, the k in k–anonymity is the smallest value of i such that E_i is non-empty.

The weighted-k for a particular generalization inducing the equi-privacy classes $E_1, E_2,$ $\ldots, E_{n_{row}}$ on the anonymized data set is then obtained as follows.

$$k_{weighted} = \frac{\sum_{i=1}^{n_{row}} (i \cdot |E_i|)}{\sum_{i=1}^{n_{row}} |E_i|} \tag{2.7}$$

Refer to the concept of local recoding [168] in this context. A local recoding scheme produces a k–anonymization by using an individual generalization function (instead of a global one) for each tuple in the data set. This is a more powerful scheme compared to having a single generalization function since outliers can be easily suppressed without the drawbacks of an over generalization. Hence, data utility can be maintained. The weighted-k–anonymity based generalization is orthogonal to this concept in certain ways. Local recoding explores the domain of generalization functions and uses multiple points in this domain to recode different subsets of the data set differently. This puts outliers in their own subset(s), thereby making it easy to enforce a given minimum QI-group size. Weighted-k–anonymity, on the other hand, works with a single generalization function and, instead of trying to enforce a fixed minimum QI-group size, it flexibly creates QI-groups of different sizes with no minimum size constraint. The outliers then must lie in smaller QI-groups in order to maximize data utility. The similarity between the two methods is that the outliers get treated differently than the rest of the data set.

The weighted-k is an estimation of the QI-group size distribution, and hence, although the chances are very rare, a high value need not always indicate that there exists no tuple in the anonymized data set appearing in its original form, i.e. in QI-groups of size one. Since the method results in an average case analysis, rather than worst case, such generalizations can appear. We present three justifications to the use of average privacy in multi-objective analysis, rather than worst case privacy.

First, the use of minimum privacy induces a very dense search space for the purpose of multi-objective optimization. Let us assume that an encoding for a candidate generalization is represented by n bits, resulting in a search space of size 2^n. Given that a data set of size N can be generalized to have possible k values in the range of 1 (no generalization) to N (all suppressed), on the average, there are $2^n/N$ points on the search space that will map to the same k value, i.e. these points will not be distinguishable from each other on measures of privacy. Even with the modest assumption of $n = 50$ and $N = 10^6$, this average number is in the magnitude of 10^9. In addition, this average estimate is only wishful thinking. The mapping will be much denser for smaller k values. On one side, most points in the search space will induce very similar (and low) privacy levels, while on the other, finding one with significantly higher values of k will be extremely difficult. We can say that there is very little diversity in the search points when using minimum privacy as an objective to maximize. This diversity is a desired property to analyze trade-offs. Using average privacy introduces diversity since the distribution of QI-group sizes have significant avenues to be different for different possible generalizations.

Second, in the context of multi-objective optimization, we cannot exclude solutions with QI-group sizes of one since they just represent an aspect of the privacy-utility trade-off. Ideally, if another generalization with the same (or better) level of utility but without the isolated tuples exists, i.e. the tuples are embedded in QI-groups of size more than one, then it would result in a higher value of weighted-k. Moreover, such a generalization will dominate the previous one and replace it from the solution set.

Third, conformation to worst-case privacy requirements can be achieved once the trade-off front is approximated. Enforcements of worst-case requirements such as minimum QI-group size is essential during optimization in a single objective framework. However, when performing a multi-objective analysis, these enforcements are part of the post-optimization strategy. The requirement essentially helps filter out solutions from the Pareto-front approximated in the analysis. In fact, a minimum utility level can also be one such requirement.

Note that in most cases not all QI-groups with all possible sizes will be generated. Hence, certain equi-privacy classes will be empty. The weighted-k value provides a suffi-

ciently good estimate of the distribution of the QI-group sizes. A high weighted-*k* value implies that QI-groups with bigger sizes are relatively more abundant than those with very few tuples. The multi-objective problem is then formulated as finding the generalization that maximizes the weighted-k and minimizes the generalization loss for a given data set.

2.4.3.2 Problem B: Maximum allowed suppression

In this problem, we enable suppression and allow the user to specify an acceptable fraction η of the maximum suppression loss possible ($n_{row} \cdot n_{col}$). Such an approach imposes a hard limit on the number of suppressions allowed [16]. However, unlike earlier approaches, by allowing the user to specify a suppression loss limit independent of *k*, the optimization procedure can be made to explore the trade-off properties of *k* and generalization loss within the constraint of the imposed suppression loss limitation.

When suppression is allowed within a user specified limit, all tuples belonging to the equi-privacy classes $E_1, \ldots, E_d$ can be suppressed, where *d* satisfies the relation

$$\sum_{i=1}^{d}(|E_i| \cdot n_{col}) \leq \eta \cdot n_{row} \cdot n_{col} < \sum_{i=1}^{d+1}(|E_i| \cdot n_{col}). \tag{2.8}$$

Thereafter, the resulting data set becomes $(d+1)$–anonymous and also satisfies the suppression loss constraint. We can now define our optimization problem as finding the generalization that maximizes d and minimizes the generalization loss. The problem can also be viewed as the maximization of *k* and minimization of *GL* satisfying the constraint $SL \leq \eta \cdot n_{row} \cdot n_{col}$. Note that the problem formulation allows the optimization procedure to find generalizations that create equi-privacy classes with lower *i*'s of smaller size and thereby increase *d*.

2.4.3.3 Problem C: Any suppression

The third problem is formulated as an extension of the second one where the user does not provide a maximum limit on the suppression loss. The challenge here is the computation of *k*, *GL* and *SL* for a generalization without having a baseline from which to start. Since the three quantities are dependent on each other for their computation, it is important that we have some base *k* value to proceed. We adopt the weighted-*k*

value at this point. Although not very precise, the weighted-k value provides a good estimate of the distribution of the QI-groups. If a very high weighted-k value is obtained for a generalization, then the number of tuples in small equi-privacy classes is sufficiently low, in which case we can suppress them. If the weighted-k value is low, then most of the tuples belong to equi-privacy classes with low i. In this case, a higher amount of suppression is required to achieve an acceptable k for the anonymized data set. Also, high weighted-k generally implies a high generalization loss. Such trade-off characteristics are the point of analysis in this problem.

To start with, a particular generalization's weighted-k value is first computed. Thereafter, all tuples belonging to an equi-privacy class E_i with $i < k_{weighted}$ are suppressed, enabling the computation of SL. This makes the k for the anonymized data set equal to at least $k_{weighted}$. The generalization loss GL is then computed from the remaining data set. The multi-objective problem is defined as finding the generalization that maximizes $k_{weighted}$, and, minimizes GL and SL.

2.4.3.4 Problem D: (k, ℓ)–safety

This problem is motivated by the requirement that a data publisher may impose on obtaining an anonymized data set that is (k, ℓ)–safe. To formulate the problem, we define the equi-privacy class $E_{i,j}$. A tuple belongs to this equi-privacy class if it belongs to an i–anonymous and j–diverse QI-group. Next, an order of importance is imposed on the k and ℓ properties. Such an order specifies which property is more desired by the data publisher and enables us to define a total ordering on the equi-privacy classes $E_{i,j}$. The ordering is obtained by first arranging the equi-privacy classes w.r.t. an increasing value in the least desired property, and then for a given value in this property, the equi-privacy classes are rearranged w.r.t. an increasing value in the most desired property. For example, if the ℓ property is more desired (denoted by $k \ll \ell$), then an example total ordering could be $E_{1,1} < E_{1,2} < \ldots < E_{2,1} < E_{2,2} < \ldots$. Otherwise, the ordering would be $E_{1,1} < E_{2,1} < \ldots < E_{1,2} < E_{2,2} < \ldots$. The objective behind such an ordering is to find the first equi-privacy class E_{d_1, d_2} in that order such that, for a given acceptable fraction of

Table 2.1: Original data set T.

Tuple ID	Age	Marital Status
1	15	Never Married
2	17	Never Married
3	20	Civ-Spouse
4	26	AF-Spouse
5	28	AF-Spouse
6	30	Civ-Spouse
7	30	AF-Spouse

suppression loss η,

$$k \ll \ell : n_{col} \cdot \left(\sum_{i=1}^{d_1-1} \sum_{j=1}^{\mathcal{L}} |E_{i,j}| + \sum_{j=1}^{d_2} |E_{d_1,j}| \right) > \eta \cdot n_{row} \cdot n_{col} \tag{2.9}$$

$$\ell \ll k : n_{col} \cdot \left(\sum_{i=1}^{\mathcal{K}} \sum_{j=1}^{d_2-1} |E_{i,j}| + \sum_{i=1}^{d_1} |E_{i,d_2}| \right) > \eta \cdot n_{row} \cdot n_{col}, \tag{2.10}$$

where $\mathcal{L}$ and $\mathcal{K}$ are the maximum obtainable values for ℓ and k respectively for the given data set. In other words, all tuples belonging to equi-privacy classes prior to E_{d_1,d_2} in the order can be suppressed without violating the suppression loss constraint. The data set then becomes (d_1, d_2)–safe and satisfies the suppression loss constraint. The multi-objective optimization problem is then defined as finding the generalization that maximizes d_1, maximizes d_2, and minimizes GL.

2.4.4 Illustrative example

Before discussing the solution methodology, we present a small example that illustrates the scope of these problems and type of analysis that we envisage to evolve from our solution. Consider the data tuples shown in Table 2.1. Obtaining a 3-anonymous generalization would require tuples 1 and 2 to be in an equi-privacy class with one or more of the other tuples. Given the hierarchy tree of the "marital status" attribute (see Fig. 2.2), such a generalization would result in the suppression of the attribute, since the only ancestor node common to the values is the root node of the hierarchy tree. Tuples 1 and 2 act as outliers in this case which prohibit obtaining 3-anonymity without overly generalizing the "marital status" attribute.

Table 2.2: 2-anonymous generalization of original data set T.

	ID	Age	Marital Status		ID	Age	Marital Status
	1	10-19	Not Married		1	10-19	Not Married
	2	10-19	Not Married		2	10-19	Not Married
T$_1$:	3	20-39	Married	T$_2$:	3	20-29	Married
	4	20-39	Married		4	20-29	Married
	5	20-39	Married		5	20-29	Married
	6	20-39	Married		6	30-39	Married
	7	20-39	Married		7	30-39	Married

Table 2.2 shows two different 2-anonymous generalizations possible on the data set. The only difference between the two anonymizations is the extra information present in T$_2$ regarding the age range of the married individuals. Let us assume that such information is not pertinent in the utility measure, and hence both anonymizations have the same utility value. As a result, an algorithm searching for an anonymization based on 2-anonymity can return either of T$_1$ or T$_2$. However, note that anonymization T$_1$ offers better privacy preservation when tuples 3 through 7 in the original data set are concerned. Although both anonymizations are 2-anonymous, which is actually the least privacy level in all QI-groups, there does exist groups with higher k values. In T$_2$, tuples 3, 4 and 5 are associated with probability $\frac{1}{3}$ of re-identification, while tuples 6 and 7 are associated with a probability $\frac{1}{2}$. This probability is $\frac{1}{5}$ for all tuples 3 through 7 in T$_1$. Such distinctions are not visible when generalizations are sought only to satisfy a fixed minimum size for all QI-groups, ignoring the actual distribution of the sizes induced. We use the weighted-k as a measure of this distribution. The weighted-k in T$_1$ evaluates to $\frac{29}{7}$, while that in T$_2$ evaluates to $\frac{17}{7}$, clearly marking that T$_1$ has better privacy preserving potential than T$_2$. If utility in both anonymizations is same then we shall have T$_1 \preceq$ T$_2$. Otherwise, there is trade-off present between the privacy and utility factors in the two anonymizations.

2.5   Solution Methodology

Classical approaches developed to handle multiple objectives concentrate on transforming the multi-objective problem into a special form of a single objective problem formulated using certain user-based preferences. However, because of the trade-off na-

Figure 2.3: Example generalization encoding for the *workclass* constrained attribute.

ture of multi-objective solutions, the quality of a solution obtained from a transformed single objective problem is contingent on the user-defined parameters. Evolutionary algorithms for multi-objective optimization are *multi-point methods* usually working with a population of solutions and concentrate on obtaining multiple optimal solutions in a single run. We thus employ the NSGA-II [48] algorithm to solve the multi-objective problems defined in the previous section.

2.5.1 Solution encoding

Before NSGA-II can be applied, a viable representation of the generalization has to be designed for the algorithm. Here we adopt the encoding suggested by Iyengar [92]. Consider the numeric attribute "age" with values in the domain $[10,90]$. Since this domain can have infinite values, the first task is to granularize the domain into a finite number of intervals. For example, a granularity level of 5 shall discretize the domain to $\{[10,15],(15,20],\dots,(85,90]\}$. Note that this is not the generalization used to anonymize the data set. The discretized domain can then be numbered as $1:[10,15],2:(15,20],\dots,16:(85,90]$. The discretized domain still maintains the same ordering as in the continuous domain. A binary string of 15 bits can now be used to represent all possible generalizations for the attribute. The ith bit in this string is 0 if the ith and $(i+1)^{th}$ intervals are supposed to be combined, otherwise 1. For attributes with a small domain and a defined ordering of the values, the granularization step can be skipped. For categorical data, a

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Figure 2.4: One generation of NSGA-II.

similar encoding can be obtained once an ordering on the domain values is imposed as discussed in Section 2.3. Fig. 2.3 shows an example generalization encoding for a "workclass" attribute. The individual encoding for each attribute is concatenated to create the overall encoding for a generalization involving all attributes.

2.5.2 NSGA-II

Similar to a simple genetic algorithm, NSGA-II starts with a population P_0 of N_{pop} random generalizations. A generation index $t = 0, 1, \ldots, Gen_{MAX}$ keeps track of the number of iterations of the algorithm. Each trial generalization is used to create the anonymized data set and the corresponding values of the quantities to be optimized are calculated. Each generation of NSGA-II then proceeds as follows. An offspring population Q_t is first created from the parent population P_t by applying the usual genetic operations of selection, crossover and mutation [75]. For constrained attributes, a special crossover operator is used as discussed in the next subsection. The offspring population also gets evaluated. The parent and offspring populations are then combined to form a population $R_t = P_t \cup Q_t$ of size $2N_{pop}$. A non-dominated sorting is applied to R_t to rank each solution based on the number of solutions that dominate it. Rank 1 solutions are all nondominated solutions in the population. A rank r solution is only dominated by solutions of lower ranks.

Figure 2.5: Usual single point crossover (left) and special crossover for constrained attributes (right).

The population P_{t+1} is generated by selecting N_{pop} solutions from R_t. The preference of a solution is decided based on its rank; lower the rank, higher the preference. By combining the parent and offspring population, and selecting from them using a non-dominance ranking, NSGA-II implements an elite-preservation strategy where the best solutions obtained are always passed on to the next generation. However, since not all solutions from R_t can be accommodated in P_{t+1}, a choice is likely to be made when the number of solutions of the currently considered rank is more than the remaining positions in P_{t+1}. Instead of making an arbitrary choice, NSGA-II uses an explicit diversity-preservation mechanism. The mechanism, based on a *crowding distance metric* [48], gives more preference to a solution with a lesser density of solutions surrounding it, thereby enforcing diversity in the population. The NSGA-II crowding distance metric for a solution is the sum of the average side-lengths of the cuboid generated by its neighboring solutions. Fig. 2.4 depicts a single generation of the algorithm.

2.5.3 Crossover for constrained attributes

The usual single point crossover operator in a genetic algorithm randomly chooses a crossover point and creates two offspring by combining parts of the bit string before and after the crossover point from two different parents. As shown in Fig. 2.5 (left), such an operation can result in an invalid generalization for constrained attributes. Iyengar proposes modifying such invalid generalizations to the nearest valid generalization [92]. However, finding the nearest valid generalization can be time consuming, besides destroying the properties on which the crossover operator is based on. In this regard, Lu-

nacek et al. propose a special crossover operator that always creates a valid offspring for constrained attributes [118]. Instead of randomly choosing a crossover point, their operator forces the crossover point to be chosen at a location where the bit value is one for both parents. By doing so, both parts (before and after the crossover point) of both parents can be guaranteed to be valid generalizations individually, which can then be combined without destroying the hierarchy requirement. Fig. 2.5 (right) shows an instance of this operator.

2.5.4   Population initialization

In order to be able to use Lunacek et al.'s crossover operator, the validity of the parent solutions must be guaranteed. This implies that the initial population that NSGA-II starts with must contain all valid generalizations for the constrained attributes. For a given hierarchy tree, we use the following algorithm to generate valid generalizations for the constrained attributes in the initial population.

Starting from the root node, a node randomly decides if it would allow its subtrees to be distinguishable. If it decides not to then all nodes in its subtrees are assigned the same identifier. Otherwise the root of each subtree receives a unique identifier. The decision is then translated to the root nodes of its subtrees and the process is repeated recursively. Once all leaf nodes are assigned an identifier, two adjacent leaf nodes in the imposed ordering are combined only if they have the same identifier. Since a parent node always make the decision if child nodes will be combined or not, all generalizations so produced will be valid.

2.5.5   Experimental setup

We apply our methodology to the "adult.data" benchmark data set available from the UCI machine learning repository [103]. The data was extracted from a census bureau database and has been extensively used in studies related to k–anonymization. We prepared the data set as described in [16, 92]. All rows with missing values are removed from the data set to finally have a total of 30162 rows. The attributes "age", "education", "race", "gender" and "salary class" are kept unconstrained, while the attributes "work-

class", "marital status", "occupation" and "native country" are constrained by defining a hierarchy tree on them. The remaining attributes in the data set are ignored. For Problem D, the "occupation" attribute is considered sensitive.

For NSGA-II, we set the population size as 200 for Problem A and B, and 500 for Problem C and D. The maximum number of iterations is set as 250. A single point crossover is used for unconstrained attributes while Lunacek et al.'s crossover operator is used for constrained attributes. Also, mutation is only performed on the unconstrained attributes. The remaining parameters of the algorithm are set as follow: crossover rate $= 0.9$, mutation rate $= 0.1$ with binary tournament selection. We ran the algorithm with different initial populations but did not notice any significant difference in the solutions obtained. The results reported here are from one such run.

2.6   Results and Discussion

Before presenting our results and the analysis, we would like to emphasize that the rationale behind doing the multi-objective analysis is not to come up with a way of determining the best possible value of a model parameter. Our intention is focused at providing a global perspective of what values of the parameter are possible at different levels of data utility. The final choice of a solution depends on other feasibility criteria as well, for example, if the parameter value found at a particular utility level is acceptable to the human subjects involved or not. An inherent human factor (the data publisher or the human subjects) is thus involved in the selection of a final solution. Further, the use of NSGA-II may raise questions on whether the obtained solutions are optimal. It is possible that another algorithm, or a different metric, provides better solutions. However, our problem formulation neither has any dependency on the methodology chosen to solve them nor is particular to the loss metric used. Further, we want to emphasize that the solutions generated by the NSGA-II implementation are *valid* at each iteration of the algorithm owing to the approach we undertake in formulating the problems. For example, a solution always gives a generalization resulting in $(d+1)$–anonymity in Problem B, and, d_1–anonymous and d_2–diverse in Problem D. Of course the objective is to maximize these quantities along with the minimization of the information loss.

Figure 2.6: Solutions to Problem A found by NSGA-II. Inset figures show cumulative distribution of $|E_i|$ as i increases.

The parameters associated with NSGA-II did not have any significant affect on the quality of the solutions obtained. We believe that the special crossover operator provides a much faster rate of convergence as compared to the genetic algorithm implementation by Iyengar [92]. The following results are obtained from the standard settings as mentioned in the previous section.

The term *loss* in the following discussion signify the total information loss as a result of generalization and suppression, i.e. $loss = GL + SL$. The differentiation between GL and SL is made wherever appropriate.

2.6.1 Problem A: Zero suppression

Fig. 2.6 shows the different trade-off solutions obtained by NSGA-II. A point in the plot corresponds to a solution that induces a particular distribution of QI-group sizes on the anonymized data set. As expected, the generalization loss increases as the distribution gets more inclined towards bigger sizes. In the absence of suppression, a single group size is often hard to enforce for all tuples in the data set. Thus, a solution here results in tuples being distributed in QI-groups of varying sizes. A higher $k_{weighted}$ value signifies that most tuples belong to QI-groups of relatively much bigger sizes, in which case, the generalization loss is higher. A solution with low $k_{weighted}$ value results in tuples being distributed mostly in small QI-groups.

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The inset figures in the plot depict the cumulative distribution of the number of tuples belonging to an equi-privacy class E_i (y-axis) with respect to different i values (x-axis). The distributions of two extreme solutions corroborate the speculation that a higher generalization loss must be incurred to assure a greater level of privacy for a larger section of the data set. Low generalization losses are only possible when most tuples belong to equi-privacy classes of lower i value.

However, it should be noted that the distributions for the two example solutions are not complementary in nature. For the solution with lower generalization loss, the distribution has a continuously increasing trend, implying that equi-privacy classes of different i values exist for the solution. The other solution shows an abrupt increase signifying that the tuples either belong to equi-privacy classes with very small i or ones with very large i. The sought balance in the distribution may therefore exist with an acceptable level of generalization loss.

2.6.2   Problem B: Maximum allowed suppression

Fig. 2.7 shows the trade-off between k and *loss* in Problem B when a maximum of 10% suppression loss is allowed. Recall that the k value in this problem is $d + 1$. The top-leftmost plot shows all the solutions obtained for the problem. Each subsequent plot (follow arrows) is a magnification of the steepest part in the previous plot. Each plot shows the presence of locally flat regions where a substantial increase in the k value does not have a comparatively high increase in the *loss*. These regions can be of interest to a data publisher since it allows one to provide higher levels of data privacy without compromising much on the information content. Also, since the solutions corresponding to these flat regions evaluate to distantly separated k values, an analysis based on a single objective formulation with a fixed k shall require a much higher number of runs of the algorithm to identify such trade-off characteristics.

Interestingly, the trend of the solutions is similar in each plot. The existence of such repeated characteristics on the non-dominated front suggests that a data publisher's choice of a specific k, no matter how big or small, can have avenues for improvement, specially when the choice falls in the locally flat regions. A choice of k made on the rising parts

Figure 2.7: Solutions to Problem B ($\eta = 10\%$) found by NSGA-II. Top-leftmost plot shows all obtained solutions. Each subsequent plot (follow arrows) is a magnification of a region of the previous plot.

of the front is seemingly not a good choice since the user would then be paying a high cost in degraded data quality without getting much improvement on the privacy factor. The rational decision choice in such a case would be to lower the *k* value to a flat region of the front. We observed similar trends in the solutions when the suppression loss was reduced to a low 1%.

2.6.3 Problem C: Any suppression

The trade-off characteristics in Problem C are depicted in Fig. 2.8. Preliminary observations from the plot indicate that an increase in generalization loss results in a decrease in the suppression loss. A similar trend is observed when the *k* value increases. Since the *k* values in this case are computed directly from the weighted-*k*, an explanation for these observations is possible. A high generalization loss signifies that most tuples in the data set belong to large equi-privacy classes, thereby inducing a high weighted-*k*. This implies a low accumulation in suppression loss resulting from the deletion of tuples in equi-privacy classes with $i < k_{weighted}$. Also, as $k_{weighted}$ increases, the equi-privacy class size distribution incline more towards the ones with high *i* values resulting in lesser number

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Figure 2.8: Solutions to Problem C found by NSGA-II. Read x-axis label for a plot from the text-box along the same column and y-axis label from the text-box along the same row. Trade-off characteristics are visible across different pairs of the objective functions.

of tuples available for suppression.

The benefit of solving this problem comes in the form of an approximate solution set available for first-level analysis. For example, Fig. 2.9a shows the solutions from the set when the suppression loss is set at a maximum allowable limit of 20%. Although *GL* and *SL* are conflicting objectives here, the analysis is intended to see if an acceptable level of balance can be obtained between the two with a reasonably good value of *k*. The encircled region in the plot show that three solutions around the point ($k = 5000, GL = 35\%, SL = 17\%$) are available in this case, and hence a more specific analysis can be performed. A similar solution is found when the analysis is performed by setting the generalization loss limit to 30% (Fig. 2.9b).

2.6.4 Problem D: (k, ℓ)–safety

Fig. 2.10 depicts a subset of the solutions obtained for Problem D. The solutions correspond to a suppression loss limit set as $\eta = 0.1$. Further, the plots only show solutions

Figure 2.9: Problem C solutions for (a) $\%SL < 0.2$ and (b) $\%GL < 0.3$. Encircled solutions can be of interest to a user.

Figure 2.10: Problem D solutions for $\eta = 0.1$ and $\%loss < 0.3$. (a) k–anonymity is more desired and (b) ℓ–diversity is more desired.

for which the *loss* is less than 30%. The existence of multiple solutions for a fixed value of ℓ (or k) signifies that there is a trade-off involved in the amount of information loss and the value of k (or ℓ). An observation to make here is the number of solutions obtained for varying values of k (or ℓ). When the preference is inclined towards the k–anonymity property, the solutions obtained give more choices for the parameter k than ℓ (Fig. 2.10a). Similarly, when preference ordering is changed to $k \ll \ell$ (Fig. 2.10b), more choices are available for the ℓ parameter. More importantly, any solution in either of the the two plots satisfy the 30% constraint on *loss* and hence is a viable solution to a data publisher's request with the same constraints. To choose a single solution, we can follow

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Figure 2.11: Problem D solutions for $\eta = 0.1$ and $k \ll \ell$. (a) Maximum *loss* allowed is 20%. (b) Maximum *loss* is increased to 21%. The quality of solutions as well as the number of choices available increase considerably for a slight increase in the *loss* limit.

the same preference ordering as was used while defining the optimization problem. For example, if the ℓ–diversity property is more desirable, we can choose the solution with the highest value of k from the set of solutions with the highest value of ℓ (a (15,7)–safe solution in the plot).

We can extend our analysis to see if improvements are obtainable without much increase in the information loss. Often, the data publisher's constraint on the information loss is specified without an understanding of the trade-offs possible. A multi-objective analysis reveals the nature of these trade-offs among different objectives and can provide suggestions on how one might be improved. For example, Fig. 2.11a shows solutions when the data publisher has given a 20% constraint on the information loss. For a good balance between the two desired privacy properties, say the data publisher chooses the (20,4)–safe solution. Solutions with $\ell = 3, 5$, or 6 are avoided at this point because of the low value of k associated with them. Fig. 2.11b shows the solutions to the same problem but with a slightly higher *loss* limit of 21%. The encircled solutions depict the new choices that are now revealed to the data publisher. For a small amount of increase in the *loss* limit, the data publisher now has a choice – (20,5)–safe – which offers the same value of k as in the old choice, but with a higher value of ℓ. Further, the earlier reluctance to

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choose a solution with $\ell = 3$ is reduced after the revelation of the $(22,3)$–safe solution. In fact, there is even a candidate solution with $\ell = 6$ and a much better value in k. With this information, the data publisher now has some idea about the trade-offs possible between privacy and information loss. In fact, the trade-off analysis in this case may motivate the data publisher to relax the loss constraint, which is not a big relaxation in itself, and reap the benefits of better privacy.

2.6.5 Resolving the dilemma

It is possible to analyze the trade-off surface generated for a particular data set and provide the data publisher with an analytical form for it. Given that an approximation of the Pareto-front is known, an analytical form can be derived through a polynomial curve fitting approach. However, it must be noted that analyzing the privacy-utility trade-off in this manner is rather problem specific. It cannot be ascertained that the Pareto-front always has a definitive structure for all data sets. Any theoretical analysis motivated from Pareto behavior in a data set is limited to that particular data set, and is not directly extensible to another. We cannot say for sure if the Pareto front will be similar for different data sets with similar distributions. Understanding the trade-off is rather empirical in this work, represented directly in terms of the parameter values and the resulting utility (represented by the value of the utility metric of choice). Nonetheless, it is not always true that empirical results are just tuples of ⟨privacy,utility⟩ values without providing much insight into the trade-off involved. Observing the Pareto-front on a graphical plot can reveal underlying traits. A classic case of this is seen in Fig. 2.7. Our observations indicate here that a major part of the Pareto-front is flat, or steep, in this data set. This signify that the data publisher has more flexibility in choosing a k value for a given level of utility. The steep jumps in utility signify that there exist points for which utility can often be improved significantly with a slight deterioration in privacy.

To summarize the above discussion, we go back to the questions asked by the data publisher in the beginning of this chapter, and try to provide answers to them w.r.t. the benchmark data set.

1. We can generalize the data set in such a way that more number of tuples have a

low probability of being identified by a linking attack. There is a generalization that results in 22% loss and attains a weighted average k value of 2528 (from Fig. 2.6). The inset figure shows that a substantial fraction of the tuples belong to sufficiently big QI-groups.

2. For the constraints given, a generalization with $k = 14$ is known (from Fig. 2.7). However, if the information loss constraint can be relaxed to 26%, a solution with $k = 36$ is known. Note that an analysis of the nature performed in Problem C can be used to provide further suggestions on the trade-offs available for suppression loss.

3. A (k, ℓ)–safe solution can provide the benefits of both k–anonymity and ℓ–diversity. A generalization with a high value of k and ℓ can be an answer. However, it is required that the more desired property be specified for better analysis.

4. For the given constraints, and assuming that ℓ–diversity is more desired, a solution with $k = 20$ and $\ell = 4$ offers a good balance between the two privacy measures (from Fig. 2.11). There are other solutions with trade-offs in the k and ℓ values. However, if the information loss constraint is relaxed to 21%, the ℓ value can be increased to 5. Besides, this will also allow two additional solutions: $(22, 3)$–safe and $(18, 6)$–safe.

2.7   Conclusions

In this chapter, we investigate the problem of data privacy preservation from the perspective of a data publisher who must guarantee a specific level of utility on the disseminated data in addition to preserving the privacy of individuals represented in the data set. The conflicting nature of the two objectives motivate us to perform a multi-objective analysis on the problem, the solutions to which present a data publisher with the knowledge of different trade-off properties existent between the privacy and utility factors.

We present empirical results to demonstrate that the choice of the parameter value in the k–anonymity problem can be made in an informed manner rather than arbitrarily. The multi-objective problems are formulated to cater to differing requirements of a decision

maker, primarily focused on the maximization of the k value and minimization of the losses.

For generalizations without suppression, a unique k may not be available. However, the analysis indicates that generalizations are possible that provide a higher level of privacy for a higher fraction of the data set without compromising much on its information content. When suppression is allowed up to a hard limit, the user's choice of k should be based on an analysis similar to that performed in Problem B. Typically, the nature of the non-dominated solution set provides invaluable information on whether an anonymization exists to improve a particular value of the model parameter without much degradation in quality of the data. First-level explorations in this context can begin with gaining an overall understanding of the trade-off characteristics in the search space.

Our results also indicate that different privacy models can be combined and optimized to result in minimal information loss. However, the trade-off picture is better portrayed in cases when the model parameters are kept separated and formulated as multiple objectives. We use (k, ℓ)–safety as the combination of k–anonymity and ℓ–diversity models with the objective of demonstrating how a multi-objective formulation can be devised to search for generalizations that result in acceptable adherence to more than one privacy property and within acceptable utility levels. Trade-off analysis lends credence to the fact that often a data publisher's choice of a particular solution can be augmented with information to suggest possible improvements on one or more objectives.

The formulations presented in the chapter also address the data publisher's dilemma. They provide a methodology to analyze the problem of data anonymization in manners that appeal to the actual entity that disseminates the data. We believe that such an analysis not only reinstates the data publisher's confidence in its choice of a particular privacy model parameter, but also identifies ways of examining if the level of privacy requested by a human subject is achievable within the acceptable limits of perturbing data quality.

Future work in this direction can start with examination of the framework with other models of privacy preservation. Real valued parametrization of t makes the t–closeness model an interesting subsequent candidate. Hybrid models catering to different forms of attacks are also required. Work on this can begin with an exploration of what trade-

offs are generated when looking for the existence of two, or more, privacy properties simultaneously. We believe that transitioning these different models into the real world would require us to synchronize our perspective of the problem with those that actually deal with it.

CHAPTER 3

Pareto Optimization on a Generalization Lattice

Representation of the generalization space plays a crucial role in the design of algorithms attempting to maximize the data utility. Although Iyengar's bit vector representation is one of the most flexible schemes to represent the search space, it does not readily provide a known structure that algorithms can exploit during the search. Therefore, a majority of the anonymization algorithms [16, 72, 89, 92, 109, 110, 117, 150, 176] are known to work with a generalization space arranged in the form of a graph, called the *domain generalization lattice*, where every node is a vector signifying the amount of generalization for each quasi-identifier. Efficient traversal of the lattice has been particularly explored so that the number of nodes that undergo evaluation (determining equivalence classes and computing loss) can be reduced.

In this chapter, we extend our analysis based on the Pareto concept to identify optimal generalizations on a domain generalization lattice. Our principle contribution, the *Pareto-Optimal k-Anonymization (POkA)* algorithm, utilizes a combination of depth first traversals of the lattice to efficiently move from one Pareto-optimal node to another. Two key properties, namely *height boundary property* and *ground node property*, have been proposed to guide the search in a manner that requires minimal node evaluations while assuring that all Pareto-optimal nodes are identified. Performance analysis on a benchmark data set shows that POkA can prune a large number of sub-optimal nodes from being evaluated

Figure 3.1: (a) Example domain generalization hierarchies of attributes *ZIP, SEX* and *SALARY*. (b) Domain generalization lattice using example DGHs. (c) Generalization and specialization graphs of node $(1,0,1)$ in the lattice.

and can still obtain all Pareto-optimal nodes.

The rest of the chapter is organized as follows. Section 3.1 introduces the background concepts. Theoretical groundwork for the algorithm is presented in Section 3.2. Section 3.3 describes our algorithm. Performance analysis on a standard benchmark data set is presented in Section 3.4. Finally, Section 3.5 concludes the chapter.

3.1 Preliminaries

A data set is conceptually arranged as a table of rows (or tuples) and columns (or attributes). Each attribute denotes a semantic category of information that is a set of possible values. Attributes are unique within a table. Each row is a tuple of s values $\langle v_1, \ldots, v_s \rangle$, s being the number of attributes in the data set, such that the value v_j is in the domain of the jth attribute A_j, for $j = 1, \ldots, s$. The domain of attribute A_j is denoted by the singleton sets $A_j = \{a_{j1}\}, \ldots, \{a_{jS_j}\}$, where S_j is the size of the domain.

A generalization of attribute A_j is a union of its domain into supersets. Hence the generalized domain of A_j can be written as $H_j^1 = A_{j1}, \ldots, A_{jm}$ where $\bigcup_i A_{ji} = \cup A_j$ and $A_{jp} \cap A_{jq} = \phi$ for $p \neq q$. We then say H_j^1 is a generalized domain of A_j, denoted as $H_j^1 <_G A_j$. The domain H_j^1 can be further generalized in a similar manner to the domain H_j^2. Generalization of an attribute's domain in this manner gives rise to a *domain generalization hierarchy* (DGH) $H_j^{N_j} <_G \ldots <_G H_j^1 < H_j^0$, where $H_j^0 = A_j$. N_j is called the *length* of the attribute's DGH. Refer to Fig. 3.1a for an example DGH. The DGH is a specification of

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how an attribute's values can be combined progressively to bigger sets. H_j^0 is called a *full specialization* of attribute A_j, meaning that no two values belong to a single set. The other extreme of this is a *full generalization* $H_j^{N_j}$ where all values of the attribute belong to a single set. The *generalization level* of the attribute is signified by an integer between 0 and N_j. A generalization level of 0 signifies that all values are distinguishable from each other, while a level of N_j signifies that no two values can be distinguished from each other.

Given a DGH for each quasi-identifier in the data set, a tuple is said to be in an anonymized form when a generalization is applied on the attribute values. The anonymized form is represented as follows. Let us assume a tuple $\langle v_1, \ldots, v_s \rangle$ in the data set. Let $(n_1, \ldots, n_s); 0 \le n_i \le N_i$ be the vector representing the generalization level for each attribute; n_i is the level to use in the DGH for attribute A_i. To map the value v_1 to its generalized form we replace it by the index of the set to which it belongs in the generalized domain at level n_1. For example, if $H_1^{n_1} = A_{11}, \ldots, A_{1m}$ and $v_1 \in A_{1p_1}$, then v_1 is replaced by p_1. After performing similar operations for the other attribute values, the tuple is anonymized to the form $\langle p_1, \ldots, p_s \rangle$, p_i being the set index for value v_i in $H_i^{n_i}$. Transforming all tuples in the data set in this manner results in an anonymized data set.

The anonymized tuples of a data set can be grouped together into equivalence classes. Two anonymized tuples $\langle p_1, \ldots, p_s \rangle$ and $\langle q_1, \ldots, q_s \rangle$ belong to the same equivalence class if $p_i = q_i; 1 \le i \le s$. The *k*-anonymity property requires that every such equivalence class should be of size at least *k*. Table 3.1 shows an example of this property. With reference to Fig. 3.1a, *ZIP* is generalized at level 1, *SEX* at level 1, and *SALARY* at level 0. Note that as higher *k* values are desired, higher generalization levels need to be used for the attributes. In principle, the anonymized tuples have categorical labels instead of the set index. Thus, two anonymized tuples in the same equivalence class have the same labels making them indistinguishable from each other.

3.1.1 Domain generalization lattice

A domain generalization lattice DGL is a graph with $\prod_i (N_i + 1)$ nodes. Every node $(n_1, \ldots, n_s); 0 \le n_i \le N_i$ is a vector of *s* dimensions where the ith element n_i specifies the generalization level for attribute A_i. The 3-anonymous table in Table 3.1 corresponds

Table 3.1: 3-anonymous version (right) of a table.

ZIP	SEX	SALARY	DISEASE		ZIP	SEX	SALARY	DISEASE
12345	M	<50K	Cancer		1234*	*	<50K	Cancer
12346	M	<50K	Gastritis		1234*	*	<50K	Gastritis
12345	F	<50K	HIV		1234*	*	<50K	HIV
12355	F	≥50K	Cancer		1235*	*	≥50K	Cancer
12355	M	≥50K	HIV		1235*	*	≥50K	HIV
12356	M	≥50K	Cancer		1235*	*	≥50K	Cancer

to the node $(1,1,0)$. An edge exists between two nodes $(n_1,\ldots,n_s)$ and $(m_1,\ldots,m_s)$ if and only if the vectors differ in exactly one element and the difference is one. To put it formally, $\sum_i |n_i - m_i| = 1$. The node $(0,\ldots,0)$ (s times) is the *fully specialized* node of the lattice and corresponds to the un-anonymized data set. The node $(N_1,\ldots,N_s)$ is the *fully generalized* node and corresponds to no disclosure of the data. Fig. 3.1b illustrates these terms.

In a typical *k*-anonymization algorithm, a node is sought in this lattice such that it satisfies *k*-anonymity and results in minimum information loss for the specified value of *k*. An exhaustive search is often not desired since evaluation of equivalence class sizes on a moderately sized data set can be computationally intensive. Most algorithms therefore perform some form of pruning of the lattice. Note that the algorithms we refer to here are meant to find optimal generalization levels for a given value of *k*. Pruning of the lattice when no *k* value is specified is an unresolved problem until now.

Given a domain generalization lattice and a node $(n_1,\ldots,n_s)$, we can also define a *specialization graph* which contains the nodes $(p_1,\ldots,p_s)$ such that $p_i \leq n_i; 1 \leq i \leq s$. Edges between nodes in this graph are drawn similar to as in a DGL. The node $(n_1,\ldots,n_s)$ is called the *root node* of the specialization graph. Along similar lines, we can also define a *generalization graph* if nodes instead satisfy $p_i \geq n_i; 1 \leq i \leq s$. Fig. 3.1c highlights the specialization and generalization graph of the node $(1,0,1)$. Intuitively, the specialization graph of a node contains all other nodes which are more specialized in one or more attributes, and in effect induce comparatively smaller *k* and lower loss. Contrary to that, a generalization graph of a node contains all other nodes which are more generalized in one or more attributes, and in effect induce comparatively larger *k* and higher loss.

Figure 3.2: Depiction of Pareto-optimal nodes.

Hence, we shall often use the term "move down" and "move up" while traversing a specialization and generalization graph respectively. These two graphs will be used later while performing the search for optimal nodes.

3.1.2 Pareto-optimal generalization

Let k_N and $\mathcal{L}_N$ signify the k and information loss associated with a node N in the domain generalization lattice. The node N is a Pareto-optimal generalization if there is no other node M in the lattice such that one of the following two conditions hold.

- $k_M \geq k_N$ and $\mathcal{L}_M < \mathcal{L}_N$, or

- $k_M > k_N$ and $\mathcal{L}_M \leq \mathcal{L}_N$.

If one of the conditions is true, then M is said to dominate N. Therefore, a Pareto-optimal node is one whose k value cannot be improved upon by another node without increasing the information loss, and the information loss at the induced k value is minimal. Note that Pareto-optimal nodes need not always exist at every possible value of k. For example, in Fig. 3.2, no Pareto-optimal node appears with $k = 4$ since the node with $k = 5$ offers a higher value of k but with lower information loss. As is evident from the figure, trade-off behavior becomes clear when all Pareto-optimal solutions are known. The advantage of finding Pareto-optimal nodes is two fold. First, the minimal information loss at relevant k values is computed. Second, the choice of a particular solution can be based on the change of information loss rather than on arbitrary selection of k. In

the figure, the choice of $k = 5$ can perhaps be made since there is not much difference in information loss from the $k = 3$ node. Therefore, our objective is to search the DGL in an efficient manner and identify the Pareto-optimal nodes. Note that this process does not require the specification of a k value by the data publisher. Instead, optimal k values are reported as part of the set of Pareto-optimal generalizations.

3.2 Pareto Search

The basic search strategy we adopt is to start from an already known Pareto-optimal node and prune nodes that cannot be Pareto-optimal. We shall start with the Pareto-optimal node with the highest possible k value and then use it as a starting step to find the *next Pareto-optimal node*. The next Pareto-optimal node is the one with a k value closest to that of the previous one but with lower loss. Hence, given a Pareto-optimal node $N = (n_1,\ldots,n_s)$, the next Pareto-optimal node M is the one with minimum $(k_N - k_M); k_M < k_N$ and $\mathcal{L}_M < \mathcal{L}_N$. We shall call the node N a *base node.* In Fig. 3.2, the next Pareto-optimal node for the node with $k = 5$ is the node with $k = 3$. The node M is found by combining a depth-first search (DFS) traversal of the specialization graph of N with a DFS traversal of the generalization graph of another node.

The first step in this process is to have a known Pareto-optimal node to begin with – the first base node. This is not difficult since the node $(N_1,\ldots,N_s)$ is bound to be Pareto-optimal. This node induces a k value equal to the size of the data set since all tuples get transformed to the same anonymized form. No other node can produce a k value higher than this. Once the next Pareto-optimal node is found, the process is repeated using this newly found node as the base node.

The next step is to assure that the node M can be reached from the base node N. Note that $k_M < k_N$. Hence, M will not be present in the generalization graph of N. Recall that all nodes in the generalization graph of N will have more generalization in one or more attributes and will result in a higher k value. This observation results in the first level of pruning of the DGL. Given the Pareto-optimal node N, the number of nodes pruned by not searching the generalization graph of N is $\prod_i (N_i - n_i + 1)$. At first glance it may seem that the node M should be somewhere in the specialization graph of N and

Figure 3.3: Use of depth search and height search to reach node *M* from node *N* through a ground node.

is hence directly reachable from *N*. In fact, if the node *M* is in the specialization graph of *N*, then it would be the one with the highest *k* value in the set $\{(n_1, \ldots, n_{j-1}, n_j - 1, n_{j+1}, \ldots, n_s) | 1 \leq j \leq s; n_j \neq 0\}$ – immediate neighbors of *N* in the graph. Since other nodes in the specialization graph are indeed specializations of a node in this set, they would have *k* values lower than the highest possible *k* in these immediate neighbors.

However, there is still a set of nodes that are not present in the specialization graph of *N* but can potentially include *M*. These nodes are the ones that can be generated from *N* by performing generalization in some attributes, specialization in some and no change in others. A positive observation in this context is that the node *M* can still be reached from *N* through a node common in the specialization graph of *M* and *N*, called a *ground node*.

3.2.1 Ground nodes

A ground node is a node common in the specialization graphs of two nodes in the domain generalization lattice. A trivial ground node for any two nodes is the fully specialized node $(0, \ldots, 0)$. Other non-trivial nodes also do exist. Given the nodes *N* and *M*, any node in the set $\mathcal{G} = \{(g_1, \ldots, g_s) | 0 \leq g_i \leq \min(n_i, m_i)\}$ is a ground node for *N* and *M*. *M* can then be reached from *N* by first moving down the specialization graph of *N* to a ground node and then moving up in the generalization graph of the ground node. The first phase of this process, i.e. moving down to the ground node, is called a *depth search* rooted at *N*. The phase of moving up from the ground node is called a *height search* rooted at the ground node. Fig. 3.3 illustrates this process.

Note that although a depth search is essential to find a ground node, nodes traversed in the process need not be evaluated if they are not immediate neighbors of the base node. This follows from the earlier observation that if M resides in the specialization graph of N, then it will be one of the immediate neighbors. This observation leads to the second level of pruning in node evaluation. However, nodes in the height search are to be evaluated since k values will increase progressively. The only exception are nodes that are also part of the depth search but not immediate neighbors of the base node. A brute force method to perform the searches would mean traversing the specialization graph of N all the way down to the fully specialized node and then traversing the generalization graph of the trivial ground node. Clearly, this is an exhaustive search. The following two sections discuss the theoretical properties that bound the extent of search to be performed in both phases. Efficiently determining the minimum extent to search will further reduce the number of nodes to be evaluated.

3.2.2 Height search

Height search from a ground node is a DFS traversal of the generalization graph with the ground node as the root. To clarify any ambiguity in language, we say that height search is a height first traversal of the graph. The search is said to be at height h when the current node H is h steps away from the ground node G, i.e $\sum_i |h_i - g_i| = h$. The height h is a dynamically chosen parameter in our approach. The assumption we make is that both k and information loss are non-decreasing quantities when performing more generalization in an attribute. Based on this assumption, the following property states how "high" should the height search proceed before coming back to its parent node.

Theorem 3.1 Height Boundary Property. Let $\mathcal{A} = \{A_1, \ldots, A_s\}$ be a set of attributes. Given a Pareto-optimal node $N = (n_1, \ldots, n_s)$, the next Pareto-optimal node M can be found by a height search of a node in the specialization graph of N, further search up a node H being terminated whenever $k_H \geq k_N$ or $\mathcal{L}_H \geq \mathcal{L}_N$.

Proof The node to start the height search from is a ground node G present in the specialization graph of N. Since M is the next Pareto-optimal node, we have $k_M < k_N$ and

$\mathcal{L}_M < \mathcal{L}_N$. Based on the aforementioned assumption, if M is in the generalization graph of G, then it must be reached in a height search before a node H with $k_H \geq k_N$ or $\mathcal{L}_H \geq \mathcal{L}_N$ is reached. Hence, whenever such a node is encountered, we have reached the maximum height along that path. $\square$

The height to search is therefore bounded by the k and loss values of the base node. We shall start with the ground node and choose a child for subsequent search only if it has k and loss lower than that of N. Child nodes that are common to the specialization graph of N (and not immediate neighbors of N) are not evaluated and are always selected. All selected child nodes are subjected to the same evaluation for further search. Exact specification of how the next Pareto-optimal node is identified from height searches is presented in Section 3.3. The further down the ground node is from the base node, the greater will be the number of nodes that will require evaluation in the height search. We therefore need a good estimate of the ground node closest to the next Pareto-optimal node. This is achieved by the depth search.

3.2.3 Depth search

Depth search from a base node N is a DFS traversal of the specialization graph with N as the root. The *depth d* of the search is the maximum total difference in generalization levels from the base node to an internal node. Therefore, a depth search of depth d implies the traversal of nodes D such that $\sum_i |n_i - d_i| \leq d$. Under this specification, an internal node is searched further only if its maximum total difference in generalization levels from the base node is less than d. Each node at depth d in the specialization graph of N is considered a candidate ground node and is subjected to a height search. Note that the DGL is a graph (not a tree) and hence the general intuition that the number of nodes subjected to height search will increase exponentially with larger d is not true. In the following, we deduce the depth at which a ground node (not necessarily the closest one) for the next Pareto-optimal node is bound to exist and derive an estimate of the depth to actually search.

Lemma 3.1 The minimum equivalence class size of a data set with s attributes $\mathcal{A} =$

$\{A_1,\ldots,A_s\}$ is the same as the minimum equivalence class size of a data set with two attributes $\mathcal{A}_1$ and $\mathcal{A}_2$, where $\mathcal{A}_1 = A_j$ and $\mathcal{A}_2 = A_1 \times \ldots \times A_{j-1} \times A_{j+1} \times \ldots \times A_s$.

Proof The proof follows from the fact that $\mathcal{A}_2$ is simply a concatenated version of the values of the attributes $A_1,\ldots,A_{j-1},A_{j+1},\ldots,A_s$. Determining the equivalence class sizes in a data set with s attributes require finding the frequency of occurrence of every possible combination of values for $s-1$ attributes. In the case with two attributes, this step is performed while finding the equivalence class sizes for the attribute $\mathcal{A}_2$. Hence in both cases, a tuple with a particular sequence of values for $A_1,\ldots,A_s$ will belong to an equivalence class of the same size. This means that the minimum equivalence class size will also be same. $\square$

Lemma 3.2 Let A_1 and A_2 be two attributes with DGH lengths of N_1 and N_2 respectively. If $N = (n_1,n_2)$, such that $0 \le n_1 \le N_1$ and $0 \le n_2 \le N_2$, is a Pareto-optimal node, then one of $G_1 = (n_1,0)$ and $G_2 = (0,n_2)$ is a ground node for the next Pareto-optimal node.

Proof The next Pareto-optimal node M is the one with k_M closest to k_N satisfying the constraints $k_M < k_N$ and $\mathcal{L}_M < \mathcal{L}_N$, i.e. there is no other node M' such that $k_M < k_{M'} < k_N$ and $\mathcal{L}_{M'} < \mathcal{L}_N$. The first set of possibilities are the immediate descendants of N, i.e. $D_1 = (n_1 - 1, n_2)$ or $D_2 = (n_1, n_2 - 1)$. Other descendant nodes of N, i.e. nodes of the form $L = (l_1,l_2); 0 \le l_i < n_i$, have k values lower than $\max(k_{D_1}, k_{D_2})$ and hence do not satisfy the requirements for the next Pareto-optimal node. If one of D_1 or D_2 is Pareto-optimal then G_2 or G_1 respectively is a ground node for it.

If none of D_1 and D_2 is the next Pareto-optimal node then nodes of the form (l_1,h_2) or (h_1,l_2), where $n_i < h_i \le N_i$, must dominate them (after satisfying the required constraints) and are likely candidates. Since G_1 is a ground node for any node of the form (h_1,l_2) and G_2 for nodes of the form (l_1,h_2), the result still holds. $\square$

Theorem 3.2 Ground Node Property. Let $\mathcal{A} = \{A_1,\ldots,A_s\}$ be a set of attributes with N_i as the DGH length of A_i. If $N = (n_1,\ldots,n_s); 0 \le n_i \le N_i$ is a Pareto-optimal node, then one or more nodes in the set $\mathcal{G}_{best} \cup \mathcal{G}_{worst}$, where $\mathcal{G}_{best} = \{(n_1,\ldots,n_{j-1},0,n_{j+1},\ldots,n_s)|1 \le j \le s\}$ and $\mathcal{G}_{worst} = \{(0,\ldots,n_j,\ldots,0)|1 \le j \le s\}$, are ground nodes of the next Pareto-optimal node.

Proof The proof follows from the observation that the problem of finding Pareto-optimal nodes for s properties can be transformed to the case of finding Pareto-optimal nodes for two properties. To do so, we map the attribute set $\mathcal{A}$ to two attributes $\mathcal{A}_1$ and $\mathcal{A}_2$ such that $\mathcal{A}_1 = A_j$ and $\mathcal{A}_2 = A_1 \times \ldots \times A_{j-1} \times A_{j+1} \times \ldots \times A_s$. A tuple $\langle v_1, \ldots, v_s \rangle$ in the original data set is transformed to the form $\langle v_j, v_1; \ldots; v_{j-1}; v_{j+1}; \ldots; v_s \rangle$. Any anonymized version of a tuple is mapped in a similar manner. By Lemma 3.1, both data sets (generalized or not) will induce the same value of k. The loss metric can be modified so that the loss associated with a tuple in the data set with s attributes is proportional to that of the tuple in the data set with two attributes.

The step that remains is a specification for the DGH of $\mathcal{A}_2$. Nodes in the DGH of $\mathcal{A}_2$ are formed by taking every possible combination of nodes from the DGHs of the $s-1$ attributes. A node is therefore of the form $\mathcal{N}_j = (n_1; \ldots; n_{j-1}; n_{j+1}; \ldots; n_s); 0 \leq n_i \leq N_i$. The total ordering of these nodes is obtained by first sorting them in ascending order of the k value they induce and then in ascending order of the loss (if k is same for two nodes). This follows the general notion of a DGH where k (and then loss) increases as we step from one node to the next. The domain generalization lattice for two attributes contains nodes of the form $(n_j, \mathcal{N}_j)$.

Therefore, there is a bijective mapping between the set of nodes in the domain generalization lattice with s attributes and the set of nodes in the domain generalization lattice with two attributes. Since, the k and loss values of two corresponding nodes are also same, any Pareto-optimal node in the lattice with s attributes will also be Pareto-optimal in the lattice with two attributes, and vice versa.

Hence by Lemma 3.2, given N and a particular value for j, the ground nodes for the next Pareto-optimal node are one or both of $(0, \mathcal{N}_j)$ or $(n_j, 0)$. This translates to nodes of the form $(n_1, \ldots, n_{j-1}, 0, n_{j+1}, \ldots, n_s)$ and $(0, \ldots, n_j, \ldots, 0)$ in the lattice for s attributes. Since the transformation into the case with two attributes can be performed in s different ways ($1 \leq j \leq s$), the set of such possible ground nodes is given as $\mathcal{G}_{best} = \{(n_1, \ldots, n_{j-1}, 0, n_{j+1}, \ldots, n_s) | 1 \leq j \leq s\}$ and $\mathcal{G}_{worst} = \{(0, \ldots, n_j, \ldots, 0) | 1 \leq j \leq s\}$. $\square$

Assuring that nodes in $\mathcal{G}_{worst}$ are covered while performing a depth search, i.e. $d = \max_j (\sum_i n_i - n_j)$, ensures all possible Pareto-optimal nodes before N (ones with k and loss lower than that of N) will be discovered. This is achieved by allowing the discovery of nodes that require specialization in all (but one) attributes. However, this is often not required if only the immediately next Pareto-optimal node has to be found. Covering nodes in $\mathcal{G}_{best}$, i.e. $d = \max_i (n_i)$, is sufficient for this purpose. Depth search that covers nodes in $\mathcal{G}_{best}$ allows the discovery of nodes that may require full specialization in at most one attribute. To be precise, it allows finding nodes that have full specialization in the attribute with the longest DGH length. A longer DGH is typically specified for an attribute with a bigger domain size. Hence, full specialization in such an attribute has the tendency to induce small equivalence classes, thereby a small value of k. The immediately next Pareto-optimal node is more likely to have a k value closer to k_N.

A good strategy to adopt here is to ensure that attributes which are close to full specialization in N get a chance to become so while attributes that are far away from being fully specialized are explored in less depth. Consider the nature of nodes that get covered in a depth search of $d_{avg} = \lceil \frac{\sum_i N_i}{s} \rceil$ depth, d_{avg} being the average number of steps required for an attribute to become fully specialized from a fully generalized state. First, depth search to d_{avg} depth ensures that half of the number of attributes will have a chance to become fully specialized. Second, in d_{avg} number of steps, the attributes which are closer to being fully specialized stand a higher chance of becoming so. Third, taking d_{avg} steps allows combination of different levels of specialization for different attributes without making any of them fully specialized (if not already). Performance analysis in Section 3.4 corroborates that this strategy indeed provides the best balance between exploration of nodes and discovery of Pareto-optimal ones.

3.3 POkA Algorithm

The Pareto-Optimal k-Anonymization (POkA) algorithm is an iterative search method to identify the Pareto-optimal generalizations for a given data set. The attributes in the data set that are subjected to generalization (conventionally called quasi-identifiers) are specified and a DGH is defined for every such attribute. The algorithm is iterative be-

---

Procedure 3.1 HeightSearch(Node P, boolean *useNode*)

Input: A node P and a boolean value (*true* or *false*) *useNode*. N is the base node.

Output: $(M, k_M, \mathcal{L}_M)$: M is a node in $\mathcal{GG}(P)$ with the highest k value such that $k_M < k_N$ and $\mathcal{L}_M < \mathcal{L}_N$, or *NULL* if no such node exists.

 1: **if** (*useNode* $=$ *true*) **then**
 2:     $N_o = (P, k_P, \mathcal{L}_P)$
 3: **else**
 4:     $N_o = (NULL, 0, 0)$
 5: **end if**
 6: **for all** (child node C of P in $\mathcal{GG}(P)$) **do**
 7:     **if** ($C \in \mathcal{SG}(N)$ **and** $\sum_i |n_i - c_i| > 1$) **then**
 8:         $Q = HeightSearch(C, false)$
 9:     **else**
10:         $Evaluate(C)$
11:         **if** ($k_C < k_N$ **and** $\mathcal{L}_C < \mathcal{L}_N$) **then**
12:             $Q = HeightSearch(C, true)$
13:         **else**
14:             $Q = (NULL, 0, 0)$
15:         **end if**
16:     **end if**
17:     **if** ($k_Q > k_{N_o}$ **or** ($k_Q = k_{N_o}$ **and** $\mathcal{L}_Q < \mathcal{L}_{N_o}$)) **then**
18:         $N_o = (Q, k_Q, \mathcal{L}_Q)$
19:     **end if**
20: **end for**
21: **return** N_o

---

cause one combination of depth search and height search, and a base node, is required to identify one Pareto-optimal node. The process is applied repeatedly to identify subsequent nodes. We implicitly assume that outliers are handled using the maximum allowed suppression method as described in Section 2.4.3.2.

3.3.1 POkA

Height search and depth search are the two crucial components of POkA. We use the notation $\mathcal{SG}(P)$ and $\mathcal{GG}(P)$ to signify the set of nodes in the specialization graph and generalization graph of a node P respectively. Let N be the base node. k_P and $\mathcal{L}_P$ signify the k and information loss value associated with node P. Further, we assume the existence of a function *Evaluate* which takes as input a node P in the DGL and returns k_P and $\mathcal{L}_P$. These returned values are computed after using the suppression strategy mentioned earlier.

Procedure 3.2 DepthSearch(Node P)

Input: A node P in $\mathcal{SG}(N)$, N being the base node.

Output: $(M, k_M, \mathcal{L}_M)$: M is a node reachable by height search of some node in $\mathcal{SG}(P)$ and with the highest k value such that $k_M < k_N$ and $\mathcal{L}_M < \mathcal{L}_N$.

1:   $N_o = (NULL, 0, 0)$
2: **for all** (child node C of P in $\mathcal{SG}(P)$) **do**
3:     **if** $(\sum_i |n_i - c_i| = d)$ **then**
4:       **if** $(d = 1)$ **then**
5:         $Evaluate(C)$
6:         $Q = HeightSearch(C, true)$
7:       **else**
8:         $Q = HeightSearch(C, false)$
9:       **end if**
10:     **else**
11:       $Q = DepthSearch(C)$
12:     **end if**
13:     **if** $(k_Q > k_{N_o}$ **or** $(k_Q = k_{N_o}$ **and** $\mathcal{L}_Q < \mathcal{L}_{N_o}))$ **then**
14:       $N_o = (Q, k_Q, \mathcal{L}_Q)$
15:     **end if**
16: **end for**
17: **return** N_o

Procedure 3.1 presents the pseudo-code for a height search implementation. Height search initiated at a node therefore returns the node in its generalization graph with the highest k and one which satisfies the constraints on the k and loss. Any node that belongs to $\mathcal{SG}(N)$ and is not an immediate neighbor of N is not evaluated and height search proceeds without considering the k and loss of such a node. Otherwise, the node is evaluated to determine if further search is required as determined by the height boundary property. The method is initiated at a candidate ground node decided in the depth search. d signifies the depth to search in the following.

Procedure 3.2 shows the pseudo-code for a depth search implementation. The implementation is a simple DFS traversal with a height search being initiated when nodes at depth d are encountered. The best M found in these height searches is translated upwards towards the root of the specialization graph. Hence, the node M returned from a call to *DepthSearch*(N) is the next identified Pareto-optimal node.

POkA starts by a call to *DepthSearch* with the fully generalized node. For every new Pareto-optimal node identified, *DepthSearch* is iteratively called until the Pareto-optimal node with $k \leq 2$ is found. Procedure 3.3 shows the pseudo-code of this process.

Procedure 3.3 POkA()

Output: The set $\mathcal{P}$ of Pareto-optimal nodes in the DGL.

1:   $N = (N_1, \ldots, N_s)$
2:   $\mathcal{P} = \{N\}$
3:   **while** $(k_N > 2)$ **do**
4:     $(N, k_N, \mathcal{L}_N) = DepthSearch(N)$
5:     $\mathcal{P} = \mathcal{P} \cup \{N\}$
6:   **end while**
7:   **return** $\mathcal{P}$

3.3.2   Improvements

Node traversal in *DepthSearch* and *HeightSearch* can be further reduced by taking into account the structure of the DGL. Since the structure is that of a graph, nodes in the specialization graph and generalization graph will share nodes as children. This structure results in repeated visits to a node during a depth/height search initiated by multiple parent nodes that share the node as a child. Although repeated visits to the same node do not increase the number of unique node evaluations required, there is redundancy involved as the results from searching the node further have already been taken into account. We therefore perform some bookkeeping at every visited node to prevent repeated visits.

The output from every node visited during a height search, i.e. the return values, is separately stored in a list *HBest*. Whenever a height search is to be initiated at a node, the list *HBest* is first checked to find if an entry corresponding to the node exists. If it does, then the node (and subsequent nodes) has already been searched and the stored values are returned. If not, the height search is done as usual. Similar to *HBest*, a list *DBest* is created for every node visited during a depth search. Depth search at a node is not performed if results for the node already exist in the list. Both lists are emptied before calling *DepthSearch* in Procedure 3.3. Procedures 3.1 and 3.2 can be easily modified to maintain and use these lists.

3.4   Performance Analysis

We applied our methodology to the "adult.data" benchmark data set. The attributes used in this study along with their DGH lengths are listed in Table 3.2. Attributes with

Table 3.2: Attributes and DGH lengths used from the *adult census* data set.

Attribute	*Distinct values*	*DGH length*
Age	74	6
Work Class	7	3
Education	16	3
Marital Status	7	3
Race	5	1
Gender	2	1
Native Country	41	4
Salary Class	2	1

larger domains have been assigned a longer DGH. The total number of nodes in the lattice is 17920. The suppression limit η is set at 1% of the data set size, i.e. $\eta = 301$. Experiments are performed with three different loss metrics – namely general loss metric (GLM) [92], discernibility (DCN) [16] and classification error (CE) [92]. The attribute "Salary Class" is used as the class label while performing experiments with the CE metric. The lattice size in this case is 8960. Solutions reported by POkA are compared with those obtained by an exhaustive search of the entire DGL. Note that the number of nodes evaluated in the exhaustive search is equal to the size of the DGL, while that used by POkA is much less. Nonetheless, the exhaustive search provides us a definitive platform to judge the efficiency of POkA in finding true Pareto-optimal nodes. The depth d used in the experiment is set at $d_{avg} = \lceil \frac{\sum_i Ni}{s} \rceil = 3$, unless otherwise stated.

3.4.1 Convergence

Pareto-optimal nodes identified by POkA for the three different loss metrics are shown in Fig. 3.4. The top row highlights the nature of the search space while using different loss metrics and the true Pareto-optimal nodes. All plots are in log scale. An interesting observation is that, for all three loss metrics, the search space is more dense towards lower values of k. This means as POkA proceeds towards finding the Pareto-optimal nodes in these regions, the number of nodes in the specialization graph of the base node decreases. Further, the Pareto-optimal nodes follow varied trends in the three metrics - concavity, convexity and disconnectedness.

The bottom row in Fig. 3.4 compares the nodes identified by POkA with those ob-

Figure 3.4: Top row shows the search space and true Pareto-optimal nodes while using different loss metrics. Bottom row shows the Pareto-optimal nodes found by exhaustive search and the nodes obtained by POkA.

Table 3.3: Number of nodes evaluated when using different depth limits. Results are generated by using the GLM metric. The total number of nodes is 17920. Number of true Pareto-optimal nodes is 45.

Depth d	Nodes evaluated	True optima
1	502 (2.8%)	22 (48.8%)
2	1945 (10.9%)	41 (91.1%)
3 ($= d_{avg}$)	**4033 (22.5%)**	**45 (100%)**
4	6544 (36.5%)	45 (100%)
5	9205 (51.4%)	45 (100%)
6	11751 (65.6%)	45 (100%)

tained from an exhaustive search. POkA demonstrates noteworthy convergence to the true Pareto-optimal nodes across the different loss metrics. It manages to overcome the limitations that may be posed due to the arrangement of the Pareto-optimal nodes in the search space. While all solutions identified with GLM and CE are true Pareto-optimal nodes, one or two cases of sub-optimal or no identification is observed for DCN (notice the center of the plot). Nonetheless, identification of a sub-optimal node did not affect any subsequent searches. The requirement that the base node is a Pareto-optimal one is therefore not a strict one. POkA can very well be started from a sub-optimal node in the lattice and Pareto-optimal nodes with a k value lower than the starting node can still be discovered.

3.4.2   Impact of the depth parameter d

The depth d used in a depth search plays a crucial role in the identification of Pareto-optimal nodes. Table 3.3 shows the number of nodes evaluated in the lattice when using the GLM metric and varying depths. The maximum depth experimented with is 6, which is equal to $\max(N_i)$, and ensures that nodes in $\mathcal{G}_{best}$ will be reached for any base node. However, using such a value results in the evaluation of more than 60% of the nodes. As discussed in Section 3.2.3, coverage of all nodes in $\mathcal{G}_{best}$ should not be required. The guiding principle derived is to search a depth of at least $d_{avg} = \lceil \frac{\sum_i N_i}{s} \rceil$ (which is equal to 3 with the DGH lengths used). Fig. 3.5 shows the Pareto-optimal nodes identified by using a depth of d_{avg} or less. All Pareto-optimal nodes have been identified by using a depth of d_{avg} ($= 3$). Using a depth of 2 resulted in some misidentification while a depth of 1

Figure 3.5: Comparison of Pareto-optimal nodes identified using depths of 1, 2 and 3 ($=d_{avg}$).

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Figure 3.6: Percentage of nodes evaluated for varying lattice sizes. Results are generated by using the DCN metric. Varying lattice sizes are generated by considering varying number of attributes to anonymize.

missed a number of the Pareto-optimal nodes. Using a depth higher than 3 did not prove to be of any advantage, less the number of nodes evaluated increased without necessity.

3.4.3 Node pruning efficiency

We found that the node pruning efficiency of POkA is much better in domain generalization lattices of bigger sizes. Bigger lattices may be formed either when the DGH lengths of the attributes considered are sufficiently long or when the number of attributes to anonymize is large. We experimented with the latter possibility and found that the percentage of nodes evaluated dropped exponentially with increasing lattice size. Fig. 3.6 shows the percentage of nodes evaluated when using 2,3,...,8 attributes for anonymization in the census data set. The depth to search is set to d_{avg} in each case. The higher number of evaluations for smaller lattices can be attributed to the fact that the observed high concentration of solutions in certain regions of the search space no longer holds. As nodes are spread out in the search space, potential number of Pareto-optimal nodes are also high, thereby resulting in the evaluation of a higher fraction of the nodes. On an average, node evaluations are observed to be around 20% across the three metrics when anonymizing for all attributes.

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3.4.4 Summary

To summarize the results, POkA can identify true Pareto-optimal nodes for a wide range of loss metrics that structure the search space in different ways. The experimental results corroborate the theoretical motivation behind using the average number of steps for full specialization of an attribute as the depth to search. The performance of POkA does not deteriorate even if certain nodes identified by it are not Pareto-optimal and used as base nodes. Finally, the number of nodes evaluated is a small percentage of the total number of nodes when the lattice is significantly bigger than the number of Pareto-optimal nodes it contain.

3.5 Conclusions

Privacy preserving data dissemination has to minimize the information loss in the anonymized data set while protecting the identity of underlying individuals to the maximum extent possible. In the context of k-anonymity, existing approaches address these aspects only partially by concentrating only on the issue of minimum information loss. Specifically, these approaches do not provide any information on the trade-off behavior between privacy and data utility.

In this chapter, we propose the POkA algorithm to find generalization schemes that are Pareto-optimal with respect to k-anonymity and a loss metric. By identifying Pareto-optimal nodes in a domain generalization lattice we can guarantee that no other generalization can improve on the privacy aspect without deteriorating data utility. POkA uses a combination of depth first traversals of the lattice to efficiently find the Pareto-optimal nodes. Theoretical groundwork behind efficiently performing these traversals is presented. Results on a benchmark data set show that POkA has the potential to identify all Pareto-optimal nodes with a small percentage of node evaluations. They also demonstrate that the algorithm is applicable for a number of commonly used loss metrics.

Node evaluation can be further reduced if a better heuristic to stop the depth search can be found. An initial step in this direction is to investigate more stringent properties for the ground node. Another research direction is to extend the algorithm to other

models of privacy. Pareto-optimality is used here as a two dimensional concept between privacy and data utility, while there exists privacy models that require the specification of more than a single parameter. Investigating Pareto-optimal anonymization with such models is a challenging area as well.

CHAPTER 4

Incorporating Data Publisher Preferences

A standard approach in microdata anonymization is to progressively generalize the data until it is k–anonymous. However, such an approach cannot guarantee optimality if different attributes carry different levels of significance. For example, in a medical data set, attributes such as age and disease are more important than the ZIP code of the underlying patient. This opens up the possibility that a minimum information loss can be sustained even for higher values of k, thereby providing better privacy than specified. Searching for higher privacy generalizations is also fruitful if the data publisher can tolerate an information loss higher than the minimum possible. Existing optimization attempts do not embrace such preference criteria.

Further, k–anonymity is only a minimalistic measure of the privacy level. The actual privacy levels of two individuals in a k–anonymous data set can be very different. For example, consider a 3–anonymous data set. If record A is same as 2 other records while record B is same as 9 other records, the privacy level of individual B is much higher (9/10) than that of individual A (2/3). This characteristic, which we call the *anonymization bias* [58], is induced by the nature of the k–anonymity model since it only helps to identify the worst case privacy level. Given the subjective nature of information loss, we cannot ignore the possibility of a reciprocal relationship between privacy bias and information loss.

In this chapter, we first propose an adaptation of a goal programming based interactive procedure to resolve the problem of choosing a generalization scheme that meets a privacy property along with minimum bias and information loss. We build on the idea of anonymization bias to provide a quantitative measurement of the feature. This enables us to define a precise vector optimization problem for minimizing the privacy bias and information loss. We provide a formal characterization of an optimal solution to the problem in terms of solutions to an *achievement-scalarizing function* that is based on tolerable bias and loss values specified by the data publisher. Next, a possible formulation of a scalarizing function to minimize bias and information loss is presented. Finally, an interactive procedure is discussed to help explore the set of optimal solutions based on feedback received from the data publisher. Such feedback-based search is necessary when the solution generated with specified preferences is not acceptable. The procedure employs a *reference direction approach* in order to generate multiple solutions in the neighborhood of the data publisher's preferences.

The second contribution is an approach to obtain data generalizations satisfying the k–anonymity property given preference values on the information loss and privacy bias. As part of the reference direction approach, the scalarizing function formulated in the first contribution is subjected to a constrained minimization. We show how the proposed evolutionary multi-objective approach solves the minimization problem and helps resolve the issue of finding better privacy levels than specified (by the parameter k), in the presence of varying data attribute significance and data publisher preferences.

The remainder of the chapter is organized as follows. Section 4.1 provides a preliminary background on the problem. Section 4.2 provides a formal definition of an efficient solution to the problem and presents the requisite properties that a scalarizing function must possess to guarantee such efficiency. A possible scalarizing function for our problem is discussed in Section 4.3. Section 4.4 discusses the interactive reference direction approach designed on top of a minimax problem. Section 4.5 presents our multi-objective algorithm to solve the minimax problem. Empirical results on a benchmark data set are presented in Section 4.6. Finally, Section 4.7 summarizes and concludes the chapter.

Table 4.1: Example data set and its 2–anonymous generalized version.

Employee Code (emp)	Salary Class (sal)		emp	sal
81521	C1		8152*	C1
81522	C1		8152*	C1
81523	C1		8152*	C1
82635	C2		8263*	C2
82636	C2		8263*	C2
82647	C2		8264*	C2
82648	C2		8264*	C2
81634	C3		8163*	C3
81631	C3		8163*	C3
81632	C3		8163*	C3
81639	C3		8163*	C3
81630	C3		8163*	C3

4.1 Preliminaries

We shall use Iyengar's bit vector representation of a generalization (Section 2.5.1) to demonstrate the concepts in this chapter. We further associate a value ec_i to each tuple in the anonymized data set D′ to signify the size of the equivalence class to which the tuple belongs. k-anonymity then requires that $\min(EC_{D'}) \geq k$, where $EC_{D'}$ is the vector $(ec_1, \ldots, ec_{n_{row}})$ for D′. Consider the data set in Table 4.1 (left). The data set has 12 entries of 5-digit employee codes and the corresponding salary class. The right table is a generalized version of this data set where the last digit of the employee code is removed. As a result, the entries can be grouped together into equivalence classes and the corresponding equivalence class vector is $(3,3,3,2,2,2,2,5,5,5,5,5)$. The data set then becomes 2–anonymous since the minimum value in this vector is 2. Since, k–anonymity satisfies the *monotonicity* property, i.e. a k–anonymous data set is also $(k-1)$–anonymous, we shall refer to the parameter k in k–anonymity as k_{pref} and $\min(EC_{D'})$ as the effective k resulting from the generalizations. In addition, we define the following loss and bias metrics for our experimental analysis.

4.1.1 Normalized weighted penalty

Let $(w_1, \ldots, w_{n_{col}})$ be a vector of weights where weight $0 \leq w_i \leq 1$ reflects the importance of the attribute a_i. The sum of weights is fixed at 1.0. The penalty for information

loss associated with a value $v_{i,j}$ is then given as follows.

$$penalty(v_{i,j}) = w_j loss(v_{i,j}), \tag{4.1}$$

where *loss* refers to the generalization loss defined in Eq. (2.4). The penalty attains a maximum value (equal to the weight of the attribute) when the number of partitions of an attribute's domain is one. An entire tuple can thus have a penalty of at most 1.0. The *normalized weighted penalty* in D$'$ is then obtained as the fractional penalty over all tuples in the data set.

$$NWP(\text{D}') = \frac{\sum_{i=1}^{n_{row}} \sum_{j=1}^{n_{col}} penalty(v_{ij})}{n_{row}} \tag{4.2}$$

4.1.2 Normalized equivalence class dispersion

The k–anonymity model is only representative of the worst case privacy measurement. As a result, it is possible that two anonymized versions of a data set, both satisfying k–anonymity, result in very different equivalence class sizes for the tuples. The privacy level of a tuple is directly related to its ec_i value – the higher the value, lower is the probability of privacy breach. Since the k–anonymity definition does not enforce any requirement on how ec_i values should be distributed, it is often possible that an anonymization is biased towards a set of tuples ($ec_i \gg k_{pref}$) while providing minimalistic privacy ($ec_i = k_{pref}$) for others. Our attempt here is to control the occurrence of such biased privacy within acceptable limits.

The value of ec_i for a tuple can range from 1 to the number of tuples in the data set, i.e. n_{row}. This range reflects the maximum bias that can be present in the anonymized data set. The *normalized equivalence class dispersion* measures the bias as the maximum dispersion present in the ec_i values relative to the maximum possible dispersion.

$$NECD(\text{D}') = \frac{\max(EC_{\text{D}'}) - \min(EC_{\text{D}'})}{n_{row} - 1} \tag{4.3}$$

Note that tighter privacy constraints can implicitly satisfy relaxed ones owing to the monotonicity property. Hence, if the privacy constraint is $2-$anonymity, then any generalization that achieves $k-$anonymity with $k \geq 2$ is a privacy preserving generalization.

In this case, a 2−anonymous and a 3−anonymous table are both privacy preserving but with the possibility that the latter induces lower bias than the former. Under such conditions, trade-offs can exist between the amount of bias and the utility of the data.

4.2   Objective Scalarization

A typical vector optimization problem involves decision making under the presence of multiple conflicting objectives. The most important characteristic of these problems is the non-existence of a single optima, but rather a set of "incomparable" solutions with respect to the objectives. In the context of data privacy, these solutions embody the trade-off characteristics in the two objectives – minimum bias (for e.g., minimum NECD) and minimum information loss (for e.g., minimum NWP) – and are the points for analysis by a data publisher. Further, while multiple such trade-off solutions may exist, a data publisher is only interested in those that induce NECD and NWP values close to some preference levels. We ask the following questions in this regard.

1. What is an efficient solution in the multi-objective minimization of NECD and NWP?

2. Can such a solution be obtained by minimizing a scalar function that also incorporates preferences on NECD and NWP?

3. What guarantees that a minimum of the scalar function will be an efficient solution of the multi-objective problem?

4. Will it be possible to generate different efficient solutions by minimizing the scalar function?

5. Can we find a scalar function whose minimum is an efficient solution in the neighborhood of the data publisher's preferred NECD and NWP values?

The answer to the first question is grounded in the dominance based comparison of points in a multi-objective space. We provide the definition of a trade-off solution in this space using the principle of dominance, also called an *efficient point*. The second question is answered by introducing the concept of *scalar achievement functions* that combine the

two objectives into one and take the data publisher preferences as one of its parameters. The issues raised in the third and fourth questions are resolved by enforcing the *strictly order preserving* and *strictly order representing* properties in the scalar function. Finally, such a function will be formulated in the next section as an answer to the fifth question.

4.2.1 Efficient points

Let $\mathcal{F}$ be the set of privacy preserving generalizations given the privacy constraint $\mathcal{P}_{\mathcal{CON}}$. In other words, all generalizations in $\mathcal{F}$ satisfy the privacy constraint $\mathcal{P}_{\mathcal{CON}}$. A generic privacy constraint is considered in order to emphasize that this approach is not limited to k–anonymity alone. Other models such as ℓ–diversity or t–closeness may as well be used to specify $\mathcal{P}_{\mathcal{CON}}$. Further, the following discussion is free from any intrinsic characteristic of the privacy constraint, other than the fact that the set of generalizations considered (the set $\mathcal{F}$) satisfy the constraint. We shall later see in Section 4.5 how the search algorithm stays focused on this set. Due to the same reason, the privacy guarantees provided by a resulting generalization will also be same as that provided by the underlying privacy model.

Let $\Delta : \mathcal{F} \to \mathbb{R}$ be a privacy bias function that assigns a privacy preserving generalization a real number signifying the privacy bias induced by it. Similarly, let $\Pi : \mathcal{F} \to \mathbb{R}$ be an information loss function signifying the amount of information lost due to a privacy preserving generalization. NECD and NWP are examples of Δ and Π respectively.

Consider the set of points in $Q = \{(\delta, \pi) | \delta = \Delta(F), \pi = \Pi(F), F \in \mathcal{F}\}$. The set Q contains the points signifying the bias and information loss for each possible privacy preserving generalization and shall be called the *efficiency space*. Hence, each point in Q can be associated with a privacy preserving generalization in $\mathcal{F}$. The following discussion is presented in terms of the points in Q given the understanding that a solution is actually represented by the associated point in $\mathcal{F}$. A partial order can be imposed on the points in Q as follows.

Definition 4.1 DOMINANCE. $q_1 = (\delta_1, \pi_1)$ *weakly dominates* $q_2 = (\delta_2, \pi_2)$, denoted by $q_1 \prec_W q_2$, iff $\delta_1 < \delta_2$ and $\pi_1 < \pi_2$. Further, $q_1 = (\delta_1, \pi_1)$ *strongly dominates* $q_2 = (\delta_2, \pi_2) \neq q_1$, denoted by $q_1 \prec_S q_2$, iff $\delta_1 \leq \delta_2$ and $\pi_1 \leq \pi_2$.

Figure 4.1: Strong and weak efficiency. Weak efficiency implies one of the objective values may be improved without changing the other.

A privacy preserving generalization can then be characterized in terms of its ability to dominate other generalizations.

Definition 4.2 EFFICIENT POINT. $q^* \in Q$ is called a *strongly efficient* point of Q iff $\nexists q_0 \neq q^* \in Q$ such that $q_0 \prec_S q^*$. q^* is called *weakly efficient* iff $\nexists q_0 \in Q$ such that $q_0 \prec_W q^*$.

The set of all strongly and weakly efficient points of Q is denoted by $\mathcal{E}_S$ and $\mathcal{E}_W$ respectively. In the context of the optimization problem at hand, efficient points correspond to privacy preserving generalizations that induce a level of bias and information loss that cannot be reduced simultaneously by another generalization. Using weak efficiency can result in points that are equal in at least one objective compared to other weakly efficient points. Strongly efficient points demonstrate a trade-off in both objectives. Fig. 4.1 illustrates these concepts. Strong efficiency implicitly implies weak efficiency.

Refer to the data set shown in Table 4.1 (left). Assume that the employee code (denoted as 'nnnnn') can be generalized progressively by removing the last four digits from right to left one at a time. The removed digits are denoted by an asterisk. The only generalization allowed for the salary class is to merge class 1 and 2 (denoted by 'C12/3'), otherwise it must stay in an ungeneralized form (denoted by 'C1/2/3'). Let $\mathcal{P}_{\mathcal{CON}}$ be 2–anonymity. Table 4.2 shows the NECD and NWP values corresponding to the eight possible generalizations satisfying 2–anonymity. These eight generalizations form the set $\mathcal{F}$ and the corresponding NECD and NWP pairs form the set Q. $\mathcal{G}_1$ and $\mathcal{G}_2$ are the two strongly efficient points in this case. $\mathcal{G}_3$, $\mathcal{G}_4$, $\mathcal{G}_6$, $\mathcal{G}_7$ and $\mathcal{G}_8$ are weakly efficient. $\mathcal{G}_5$ is nei-

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Table 4.2: Efficiency and **ach** values for different generalizations of the data set in Table 4.1. $\overline{q_1} = (0.3, 0.08); \overline{q_2} = (0.2, 0.15); \overline{q_3} = (0.15, 0.05); \overline{q_4} = (0.25, 0.05); \overline{q_5} = (0.1, 0.5)$. $w_{emp} = 0.3; w_{sal} = 0.7$.

	Generalization	NECD	NWP	k	Efficiency	$s(\cdot, \overline{q_1})$	$s(\cdot, \overline{q_2})$	$s(\cdot, \overline{q_3})$	$s(\cdot, \overline{q_4})$	$s(\cdot, \overline{q_5})$	$pref_{dev}$ from $\overline{q_5}$
$\mathcal{G}_1$	nnnn* + C1/2/3	0.27	0.07	2	strong	**0.06**	0.12	0.07	**0.06**	0.23	-0.26
$\mathcal{G}_2$	nnn** + C1/2/3	0.18	0.09	3	strong	0.07	**0.08**	**0.06**	0.07	**0.15**	**-0.33**
$\mathcal{G}_3$	nn*** + C1/2/3	0.18	0.15	3	weak	0.12	0.09	0.12	0.13	**0.15**	-0.26
$\mathcal{G}_4$	n**** + C1/2/3	0.18	0.30	3	weak	0.24	0.17	0.23	0.25	**0.15**	-0.12
$\mathcal{G}_5$	nnnn* + C12/3	0.27	0.27	2	-	0.22	0.16	0.20	0.23	0.23	-0.05
$\mathcal{G}_6$	nnn** + C12/3	0.18	0.29	3	weak	0.23	0.17	0.22	0.24	**0.15**	-0.13
$\mathcal{G}_7$	nn*** + C12/3	0.18	0.36	3	weak	0.28	0.20	0.27	0.30	**0.15**	-0.06
$\mathcal{G}_8$	n**** + C12/3	0.18	0.50	5	weak	0.40	0.29	0.38	0.42	**0.15**	0.09

ther weakly nor strongly efficient since $\mathcal{G}_2$ weakly dominates and $\mathcal{G}_1$ strongly dominates $\mathcal{G}_5$.

4.2.2 Necessary and sufficient conditions

Multiple techniques exist to solve a vector optimization problem. These techniques may focus on approximating the set of efficient points using ϵ–*dominance* [3, 66, 105, 174] or concentrate on parts of the set using *reference points* [49, 119, 170, 177, 192]. A standard approach in the latter category is to associate a scalar problem to the original vector problem and generate efficient points by a single objective optimization of the scalar function [129, 130]. A typical scalarizing function $s : \mathbb{R}^2 \to \mathbb{R}$ defined over $\mathbb{R}^2$ depends on an arbitrary parameter $\bar{q} \in \mathbb{R}^2$, the functional form being denoted as $s(\cdot, \bar{q})$. Such functions must possess certain properties so that a minimum of the function can imply an efficient point of Q and vice versa.

Definition 4.3 STRICTLY ORDER PRESERVING. Let $s : \mathbb{R}^2 \to \mathbb{R}$ be a scalarizing function with respect to $\bar{q} \in \mathbb{R}^2$. s is strictly order preserving in Q iff

P1. $\forall q_1, q_2 \in Q, q_1 \prec_S q_2 \Rightarrow s(q_1, \bar{q}) \leq s(q_2, \bar{q})$, and

P2. $\forall q_1, q_2 \in Q, q_1 \prec_W q_2 \Rightarrow s(q_1, \bar{q}) < s(q_2, \bar{q})$.

These two properties enforce the monotonicity requirements on s so that a minimum of the function is an efficient point of Q. This provides a sufficient optimality condition. The function s must satisfy another property so that any efficient solution of Q can be obtained by minimization of s parametrized by $\bar{q}$.

Definition 4.4 STRICTLY ORDER REPRESENTING. Let $s : \mathbb{R}^2 \to \mathbb{R}$ be a scalarizing function with respect to $\bar{q} \in \mathbb{R}^2$. s is strictly order representing in Q iff

R1. $\forall q_0 \in S_0 = \{q \in Q | s(q, \bar{q}) \leq w_c\}$, we also have $q_0 \prec_S \bar{q}$, where $w_c = s(\bar{q}, \bar{q})$, and

R2. $\forall q_0 \in S_0' = \{q \in Q | s(q, \bar{q}) < w_c\}$, we also have $q_0 \prec_W \bar{q}$.

The strictly order representing properties help us prove that each efficient solution of Q has a corresponding value of the parameter $\bar{q}$ so that the minimum of s is the efficient solution. The scalarization thus provides the necessary optimality conditions.

The following propositions provide a characterization of the efficient points of Q in terms of the solutions to a strictly order preserving and strictly order representing function s.

Proposition 4.1 If q^* is a global minimum point of $s(q, \bar{q})$ on Q, then $q^* \in \mathcal{E}_W$.

Proposition 4.2 If q^* is a unique global minimum point of $s(q, \bar{q})$ on Q, then $q^* \in \mathcal{E}_S$.

Proposition 4.3 If $q^* \in \mathcal{E}_W$, then q^* is a global minimum of $s(q, \bar{q})$ on Q for $\bar{q} = q^*$.

Proposition 4.4 If $q^* \in \mathcal{E}_S$, then q^* is a unique global minimum of $s(q, \bar{q})$ on Q for $\bar{q} = q^*$.

A global minimum of the scalar achievement function should not be confused as a global minimum in the NWP vs. NECD objective space. In fact, a global minimum is not defined in a multi-objective case. Following Propositions 4.1 and 4.2, a global minimum of $s(\cdot, \bar{q})$ (a single objective function) only implies that the resulting solution is an efficient point of the two-objective space. Further, by Propositions 4.3 and 4.4, one can arrive at different efficient points by modifying the parameter $\bar{q}$ in the achievement function.

With reference to Table 4.2, Propositions 4.1 and 4.2 guarantee that the global minimum of a strictly order preserving and strictly order representing function will never appear at generalization $\mathcal{G}_5$. Further, if the parameter $\bar{q}$ of the function is set to the corresponding NECD and NWP values of a generalization, then the global minimum is guaranteed to appear at that generalization (Propositions 4.3 and 4.4). Hence, every efficient point is reachable by the function.

Note that minimal points of a scalarizing function does not correspond to the final solution of choice by the data publisher. They are rather used to locally approximate the preferences of the data publisher with the guarantee that the generated solutions are efficient. Data publisher preferences are typically embodied in the parameter $\bar{q}$ of the function (also called a *preference point, reference objective* or *aspiration level*) with the idea that an efficient solution minimizing some sort of distance from $\bar{q}$ is sought. This is followed by an interaction with the data publisher to inquire if the reported solution is satisfactory. Order preserving and order representing properties guarantee that the reported solution (generated by a minimization of the scalarizing function) is efficient, or is the best one possible within the specification of the aspiration levels. Therefore, in

this framework, an optimal solution from the perspective of the data publisher is not the generalization that results in minimum information loss (as assumed in existing works), but the generalization that meets the preference levels in the best possible way. The next section dwells into the formulation of a scalar achievement function that integrates these requirements.

4.3 Scalarizing Bias and Loss

The typical constrained optimization problem explored in disclosure control is to find an anonymized version of a data set, or effectively a set of generalizations resulting in the anonymized version, that induce minimum information loss subject to the constraint that the anonymized data set is k–anonymous. Given the NP-hard nature of the problem [127], heuristic based approaches in this context progressively increase the amount of generalization for the attributes until the k–anonymity constraint is satisfied [16, 72, 176]. The anonymized data set at this point is assumed to incur minimum information loss. These approaches have two major drawbacks.

First, the information loss metric is assumed to have a monotonic relationship with the amount of generalization. In other words, as more generalization is performed (no matter for which attribute), the information loss increases. Only under this assumption can one claim that by performing generalization only to the extent necessary to satisfy the k–anonymity constraint, we shall also be minimizing the information loss. However, the assumption is not valid when all attributes do not carry the same significance.

Second, these approaches do not provide the framework to explore the possibility of attaining higher effective k values. Owing to the monotonicity property, an effective k value higher than k_{pref} will also satisfy the anonymity constraint, but comes with the added advantage of better privacy. Further, existing approaches do not take into account any preference specified on information loss. There are some successful attempts to obtain all possible k–anonymized versions of a data set [109, 150], out of which the optimal one can be chosen based on preference criteria. Nonetheless, the set of solutions obtained with those approaches still remains exponentially large, making the search for an optimal choice equally difficult to perform. The issue of privacy bias remains unexplored in all

these attempts.

The objective behind the scalarization of bias and utility is to arrive at privacy preserving generalizations that correspond to efficient solutions that are close to a reference objective. The reference objective is a point that signifies a tolerable level of bias and information loss to the data publisher. Depending on whether the reference objective lies inside or outside the efficiency space, the specified bias and information loss constraint may or may not be satisfied. If solutions better than the reference objective exist, then an efficient solution as far as possible from the reference point will provide the best possible improvements beyond the aspirations of the data publisher. On the other hand, if the reference objective is unachievable, then the data publisher can at best have a solution which is efficient and closest to the reference point.

We seek a scalarizing function that embeds these two requirements. In addition, the strictly order preserving and strictly order representing properties must be satisfied so that efficient points can be generated by a minimization of the scalar function. We present here one possible formulation for such a function.

Let $q_{ideal} = (\delta_{ideal}, \pi_{ideal})$ denote the ideal point in $\mathbb{R}^2$, the components of which are obtained by individual minimization of the bias and loss functions, i.e. $\delta_{ideal} = \min \Delta(\cdot)$ and $\pi_{ideal} = \min \Pi(\cdot)$. If a unique $F \in \mathcal{F}$ minimizes both the bias and loss functions, then q_{ideal} is the optimal solution. However, under the presence of trade-off behavior in the two functions, such a generalization will not exist. Hence, the ideal point is of theoretical importance only. For most cases in data privacy, the ideal point is the point $(0,0)$. Next, let $q_{utp} = (\delta_{utp}, \pi_{utp})$ be an utopian point computed as $\delta_{utp} = \delta_{ideal} - \epsilon_\delta$ and $\pi_{utp} = \pi_{ideal} - \epsilon_\pi$ where ϵ_δ and ϵ_π are small positive numbers. The scalarizing function **ach** $:= s(q, \bar{q})$ for $q = (\delta, \pi) \in \mathbb{R}^2$ and $\bar{q} = (\bar{\delta}, \bar{\pi}) \in \mathbb{R}^2$ is then formulated as

$$s(q, \bar{q}) = \max \ \left[w(\delta - \delta_{utp}), (1 - w)(\pi - \pi_{utp}) \right], \tag{4.4}$$

$$\text{where } w = \left[\frac{1}{\bar{\delta} - \delta_{utp}} \right] \bigg/ \left[\frac{1}{\bar{\delta} - \delta_{utp}} + \frac{1}{\bar{\pi} - \pi_{utp}} \right].$$

This scalarization of bias and loss provides a maximal over-achievement of the objectives if the reference point is feasible. Otherwise, the function provides a minimal underachievement. The parameter w allows us to vary the weights on the two objectives,

thereby providing a mechanism to explore the neighborhood of a solution. The precise impact of w is discussed in Section 4.4.

Consider the preference point $\bar{q}_1 = (0.3, 0.08)$ in Table 4.2. The minimum of the **ach** function in this case appears at $\mathcal{G}_1$ since the corresponding NECD and NWP values of 0.27 and 0.07 can over-achieve the preferred ones. However, $\mathcal{G}_1$ cannot provide the same improvements for the preference point $\bar{q}_2 = (0.2, 0.15)$. The minimum in this case appears at $\mathcal{G}_2$ since it can provide the best over-achievement in the two objectives. $\bar{q}_3 = (0.15, 0.05)$ and $\bar{q}_4 = (0.25, 0.05)$ are infeasible preferences. The minima here are obtained at the generalizations that produce minimal underachievement, i.e. $\mathcal{G}_2$ and $\mathcal{G}_1$ respectively. Further, the minimum point in all cases is also an efficient point in the NECD vs. NWP objective space. This is due to the strictly order preserving property of **ach**.

Theorem 4.1 The scalarizing function **ach** is strictly order preserving.

Proof To prove property P1, consider the distinct points $q_1 = (\delta_1, \pi_1), q_2 = (\delta_2, \pi_2) \in Q$ such that $q_1 \prec_S q_2$.

Hence, we have $\delta_1 \leq \delta_2$ and $\pi_1 \leq \pi_2$. Assuming that $\bar{q} > q_{utp}$ (a valid assumption since the reference point will at best be q_{ideal}), we have $0 < w < 1$, giving us $(1 - w) > 0$. We can thus obtain the following two relations.

$$
\begin{aligned}
w(\delta_1 - \delta_{utp}) &\leq w(\delta_2 - \delta_{utp}) \\
(1 - w)(\pi_1 - \pi_{utp}) &\leq (1 - w)(\pi_2 - \pi_{utp})
\end{aligned}
$$

Using the observation that $a \leq b$ and $c \leq d$ implies $\max(a, c) \leq \max(b, d)$, we obtain $s(q_1, \bar{q}) \leq s(q_2, \bar{q})$, thus proving property P1.

To prove property P2, let $q_1 \prec_W q_2$. We then have $\delta_1 < \delta_2$ and $\pi_1 < \pi_2$. The remainder of the proof follows in a manner similar to above, giving us $s(q_1, \bar{q}) < s(q_2, \bar{q})$. Thus **ach** is strictly order preserving. $\square$

Theorem 4.2 The scalarizing function **ach** is strictly order representing for

$$
w_c = 1 / \left[\frac{1}{\bar{\delta} - \delta_{utp}} + \frac{1}{\bar{\pi} - \pi_{utp}} \right]. \tag{4.5}
$$

Proof To prove property R1, let $q_1 = (\delta_1, \pi_1) \in S_0$. Hence $s(q_1, \bar{q}) \le w_c = s(\bar{q}, \bar{q})$, i.e.

$$\max \left[w(\delta_1 - \delta_{utp}), (1-w)(\pi_1 - \pi_{utp}) \right] \le w_c.$$

Rewriting the expression in terms of w_c we get

$$\max \left[w_c \left(\frac{\delta_1 - \delta_{utp}}{\bar{\delta} - \delta_{utp}} \right), w_c \left(\frac{\pi_1 - \pi_{utp}}{\bar{\pi} - \pi_{utp}} \right) \right] \le w_c$$

$$\text{i.e. } w_c \left(\frac{\delta_1 - \delta_{utp}}{\bar{\delta} - \delta_{utp}} \right) \le w_c \text{ and } w_c \left(\frac{\pi_1 - \pi_{utp}}{\bar{\pi} - \pi_{utp}} \right) \le w_c.$$

After simplification we get, $\bar{\delta} - \delta_1 \ge 0$ and $\bar{\pi} - \pi_1 \ge 0$. Hence, $q_1 \prec_S \bar{q}$. This proves property R1.

To prove property R2, let $q_1 \in S_0'$. Hence $s(q_1, \bar{q}) < w_c$. By proceeding in a manner as above we get $\bar{\delta} - \delta_1 > 0$ and $\bar{\pi} - \pi_1 > 0$. Therefore, $q_1 \prec_W \bar{q}$. Thus **ach** is strictly order representing. $\square$

Based on Propositions 4.1 and 4.2, minimization of **ach** (a minimax problem) over Q will therefore result in an efficient privacy preserving generalization. However, solutions to the minimax problem can return either weakly or strongly efficient solutions. Ideally, a strongly efficient solution is more desirable since weakly efficient solutions cannot guarantee that one of the objective values cannot be reduced by keeping the other constant. Strongly efficient solutions are indicated by a unique global minimum of **ach**. For the case when multiple global minima exists, inference of strong efficiency is difficult. As a resolution to this problem, we choose a solution that provides the maximum over-achievement or minimum underachievement with respect to the reference objective, i.e. choose the solution $q = (\delta, \pi) \in Q_g$ with minimum *preference deviation, $pref_{dev} = (\delta + \pi - \bar{\delta} - \bar{\pi})$*, where $Q_g \subseteq Q$ is the set of global minima points of **ach**. Thus, if an unachievable reference objective is specified by the data publisher, an efficient solution as close as possible to the reference point will be returned. However, if the reference objective is suboptimal (lower bias and loss possible), then an efficient solution as far away as possible from the reference point is returned. Further, Propositions 4.3 and 4.4 guarantee that if the reference objective chosen by the data publisher is an efficient point in itself, then a minimum of **ach** will exist exactly at the desired solution.

Consider the preference point $\bar{q}_5 = (0.1, 0.5)$ in Table 4.2. Multiple minima points exist for **ach** in this case. These points provide the least underachievement in NECD while satisfying the NWP preference of 0.5. The minimum preference deviation is obtained in $\mathcal{G}_2$ as it best over-achieves the NWP preference. $\mathcal{G}_2$ is also the strongly efficient point in the set of minima.

Incorporating data publisher preferences in the optimization procedure can also potentially hinder *minimality attacks* [180]. Minimality attacks exploit the knowledge that most data generalization algorithms attempt to enforce the privacy requirement with as less generalization as possible (minimum information loss). Hence, most existing methods are prone to this attack. However, the **ach** function subjected to minimization in this work is parametrized by the data publisher preferences. Therefore, a minimum of this function is not necessarily the generalization that induces the minimal information loss. In other words, the extent of generalization performed is not guided by the minimality principle (as assumed in minimality attacks), but is rather dependent on what the data publisher specifies as tolerable information loss. This can void the efficacy of minimality attacks since the preferences of the data publisher are most likely unknown to the attacker. However, a more extensive analysis is required to validate this claim.

4.4 Reference Direction Approach

While a solution obtained by minimizing the **ach** function is efficient, a data publisher may not find the reported solution satisfactory enough with respect to the reference objectives in mind. This is because the initial knowledge of the data publisher on the feasibility of a solution inducing the aspired bias and loss is limited. Once a solution is generated, the data publisher obtains additional knowledge on what levels of bias and loss are possible around the neighborhood of the preference point. This prompts for an interactive method that allows the data publisher to progressively explore efficient solutions until a satisfactory one is found. We adapt a method based on *reference directions* to facilitate this exploration [47, 128].

Definition 4.5 REFERENCE DIRECTION. Let q be a point in $\mathbb{R}^2$ and $\bar{q} \in \mathbb{R}^2$ be a prefer-

Procedure 4.1 ReferenceDirectionApproach(Point q_1, Point $\bar{q}$, integer n)

Input: Initial point $q_1 \in \mathbb{R}^2$ and a preference point $\bar{q} > q_{utp}$. n is the number of efficient solutions to generate along the reference direction.

1: $t = 1$
2: **repeat**
3:     $Q_{eff} = \phi$
4:     $d_t = \bar{q} - q_t$ {*vector arithmetic*}
5:     **for all** (m in $\{1,\dots,n\}$) **do**
6:         $\bar{q}_m = q_t + (m/n)d_t$
7:         Obtain q_m^* as the solution to the problem of minimizing **ach** $:= s(q, \bar{q}_m)$
8:         $Q_{eff} = Q_{eff} \cup \{q_m^*\}$
9:     **end for**
10:    **if** (no solution in Q_{eff} is satisfactory) **then**
11:        Choose a point in Q_{eff} as q_{t+1} to define a new reference direction
12:        $t = t + 1$
13:    **end if**
14: **until** (a satisfactory solution is found)

ence point. The reference direction d is then defined as the vector $d = \bar{q} - q$.

The basic approach here is to generate a number of efficient points along the projection of a reference direction on the efficiency space. This gives the data publisher an idea of the trade-off characteristics of solutions in the neighborhood where the data publisher's interest lies in the first place. If a satisfactory solution is found, then the process stops. Otherwise, one of the generated solutions is chosen to define a new reference direction for the next iteration of the process. This procedure of specifying a new search direction enables a data publisher to control how much deviation from the aspiration levels is tolerable in a certain objective. The reference direction approach computes what parameter value needs to be passed to the scalar achievement function to generate solutions in this new direction of interest. A new set of solutions is then generated and the process continues. By doing so, the data publisher can extensively explore the neighborhood surrounding the preference point until a solution is in agreement with the publisher's requirements. We shall first provide the basic steps of the method (Procedure 4.1) and then provide a geometrical perspective of the procedure.

Refer to the data set in Table 4.1. Assume that the data publisher starts with a preference point $\bar{q} = (0.1, 0.1)$. Based on the extent of NECD and NWP values to explore,

Figure 4.2: (a) Geometrical interpretation of solution procedure to Chebyshev min-max problem. (b) Geometrical interpretation of reference direction approach.

the publisher sets q_1 to $(1.0, 0.2)$ and $n = 10$. Multiple points will then be generated on the line segment joining $\bar{q}$ and q_1 at intervals of $(0.1, 0.01)$. Each such point is used as the parameter while minimizing the **ach** function, resulting in $Q_{eff} = \{\mathcal{G}_1, \mathcal{G}_2\}$. The data publisher at this point can choose either of $\mathcal{G}_1$ or $\mathcal{G}_2$ as the solution, or move ahead to the second iteration by selecting one of these points as q_2. However, no new solution will be found in this case.

In order to understand the method from a geometrical perspective, we rewrite the minimax problem in the so-called *Chebyshev min-max* form. In this notation, $\min_{q \in Q} \mathbf{ach}$ is written as a constrained minimization problem, given as

$$
\text{minimize } \lambda \text{ subject to } \begin{cases} w(\delta - \delta_{utp}) & \leq \lambda \\ (1 - w)(\pi - \pi_{utp}) & \leq \lambda \, . \\ (\delta, \pi) & \in Q \end{cases} \tag{4.6}
$$

The Chebyshev problem has the geometrical interpretation of a directional search starting at q_{utp} and progressing along the direction $z = (\frac{1}{w}, \frac{1}{1-w})$, i.e. the search takes place on the straight line $q_{utp} + tz$ where t is a real positive parameter. Note that the reference point $\bar{q}$ lies on this straight line, as given by the point when $t = w_c$. Since a point on this line moves away from q_{utp} as t increases, thereby increasing λ, minimum value of λ is achieved with the lowest value of t that gives a point in Q. Refer to Fig. 4.2a. For the unattainable reference objective $\bar{q}_1$, this gives the point where the shifted reference point along the search direction first touches the efficiency space. On the other hand, for the reference objective $\bar{q}_2$, the search along the direction encounters a point in

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the efficiency space before encountering the reference point, i.e. $t < w_c$, thereby providing an over-achievement.

Note that the direction of search is decided by w, which in turn is parametrized by the reference objective. It is therefore possible to change the direction of search by providing different reference objectives. The interactive procedure does so by generating intermediate reference objectives $\bar{q}_m$ on the reference direction. Refer to Fig. 4.2b. At iteration t, a current solution q_t and the reference objective $\bar{q}$ defines the reference direction. Intermediate reference points are then generated along this direction. For each such point, a search is performed along the straight line joining q_{utp} and the reference point. In other words, solutions to the Chebyshev problem is found taking different search directions, each returning an efficient solution in the neighborhood of q_t and $\bar{q}$. For example, the solution q_m^* is found as the shifted intersection of the reference direction and straight line defined by the vector $(\frac{1}{w}, \frac{1}{1-w})$ with w being computed using the reference objective $\bar{q}_m$.

Recall that since **ach** is strictly order representing, all efficient solutions are minima of some representation of the function (parametrized by the reference objective). This in turn implies that there exists some value of w (again, parametrized by the reference objective) corresponding to every efficient point, such that the solution to **ach** is the efficient point.

Theorem 4.3 Let $q^* \in Q$ be an efficient solution. There exists a value of w such that $0 < w < 1$ and q^* is a unique global minimum of **ach**.

Proof Let $q^* = (\delta^*, \pi^*) \in Q$ be an efficient solution (weakly or strongly). To the contrary, let us assume that there exists no positive value of w such than q^* is a unique global minimum of **ach**.

Let us set $\bar{q} = q^*$. This gives us

$$
w = \left[\frac{1}{\delta^* - \delta_{utp}} \right] / \left[\frac{1}{\delta^* - \delta_{utp}} + \frac{1}{\pi^* - \pi_{utp}} \right].
$$

Note that w is greater than zero (since for any $q \in Q$, $q > q_{utp}$) and less than one. In the Chebyshev formulation of **ach**, q^* will be a feasible solution. In other words, we can find a λ^* such that $w(\delta^* - \delta_{utp}) \leq \lambda^*$ and $(1 - w)(\pi^* - \pi_{utp}) \leq \lambda^*$. After simplification, the least value of such a λ^* is found to be w_c.

However, since q^* is not a unique global minimum, there must exist another point $q_0 = (\delta_0, \pi_0) \in Q$ with a corresponding $\lambda = \lambda_0$ such that $\lambda^* \geq \lambda_0 \geq w(\delta_0 - \delta_{utp})$ and $\lambda^* \geq \lambda_0 \geq (1 - w)(\pi_0 - \pi_{utp})$. By substituting λ^* with w_c, and the value of w in the two relations, we arrive at the following two expressions: $\delta^* \geq \delta_0$ and $\pi^* \geq \pi_0$, or in other words, $q_0 \prec_S q^*$. Hence, q^* cannot be an efficient solution unless $q_0 = q^*$. Hence, q^* is a unique global minimum of the Chebyshev problem. Therefore, there exists a value $0 < w < 1$ such that q^* is a unique global minimum of **ach**. $\square$

Thus, **ach** can be used to generate any efficient point by varying its parameters. However, unlike a typical weighted sum approach, parameter specification in this approach is more intuitive to the data publisher, namely the aspiration levels of the publisher. Artificial parameters like weights are often difficult to specify, with the drawback that they can sometimes be inconsistent with intuition. Further, convergence in the interactive procedure is completely guided by the data publisher. Hence, although the finally chosen solution may be very different from the initial aspiration levels, it is guaranteed to be a satisfactory one.

4.5   Minimizing the Achievement Function

A crucial step in the reference direction approach is finding a privacy preserving generalization that minimizes **ach** (a solution q_m^* in Step 7). This optimization problem is repeatedly solved as part of the approach. With respect to $k-$anonymity and the metrics NWP and NECD, the problem can be stated in the following manner. Here, $\bar{\pi} = NWP_{pref}$ and $\bar{\delta} = NECD_{pref}$.

Problem 4.1 Optimization Problem (OP). Given a data set D, k_{pref}, NWP_{pref} and $NECD_{pref}$, find the anonymized data set D$'$ that minimizes the achievement function **ach** subject to the constraint $k_{pref} - min(EC_{D'}) \leq 0$.

The optimization problem at hand is a constrained single objective problem. In this section we propose an approach based on evolutionary multi-objective optimization to find a solution to the problem. The method involves transforming the constraint into a

separate objective giving us a bi-objective vector optimization problem [40]. The multi-objective variant of OP is formulated as follows.

Problem 4.2 Multi-Objective Optimization Problem (MOOP). Given a data set D, k_{pref}, NWP_{pref} and $NECD_{pref}$, find the anonymized data set D′ that minimizes the achievement function $f_1(D') : \mathbf{ach}(D')$ and the function $f_2(D') : k_{pref} - min(EC_{D'})$.

Solutions to the MOOP are characterized by the Pareto-dominance concept. Under such a characterization, an anonymized data set D′ found by the solution methodology is a non-dominated solution to the MOOP if it cannot find another solution D″ such that

- $f_1(D'') \leq f_1(D')$ and $f_2(D'') < f_2(D')$, or

- $f_1(D'') < f_1(D')$ and $f_2(D'') \leq f_2(D')$.

A direct and positive consequence of using this formulation is the exposure of higher effective k solutions, if any. Note that a solution to OP only needs to satisfy the constraint $k_{pref} - min(EC_{D'}) \leq 0$. In the multi-objective formulation, the solutions undergo further filtering based on non-dominance – for two solutions with equal value of **ach**, the one with higher effective k (lower f_2) gets preference. Thus, if multiple solutions to OP exists at different effective k values, the multi-objective approach directs the search towards the one providing the highest level of privacy.

Recall the case when multiple minima are obtained with $\bar{q_5} = (0.1, 0.5)$ as the preference point in Table 4.2. All points except $\mathcal{G}_8$ will be filtered out by the multi-objective optimizer since it provides a comparatively higher value for k. Given that a NWP value of 0.5 can be tolerated, $\mathcal{G}_8$ leverages it to improve the privacy level. Note that if the NWP preference is changed to say 0.2, then $\mathcal{G}_8$ will no longer be a minimum point of **ach**. In that case, there will be multiple solutions with same k and **ach** values. The preference deviation metric will then be used to choose a solution.

4.5.1 A modified selection operator

We use the Non-dominated Sorting Genetic Algorithm-II (NSGA-II) with Iyengar's bit vector representation to obtain solutions to the MOOP. While most components of

NSGA-II are retained in the original form, a modification is proposed for the selection procedure in order to direct solutions towards the feasible region of OP [65]. NSGA-II employs a *crowded comparison operator* as part of its binary tournament selection scheme. This operator gives preference to solutions with lower ranks (in accordance with the non-dominated ranking scheme of the algorithm). For the case when the compared solutions are of the same rank, the crowding distance metric is used to choose the solution with least diversity. Our modification involves distinguishing between feasible $(f_2(D') \leq 0)$ and infeasible $(f_2(D') > 0)$ solutions of OP during the selection procedure. The procedure is outlined as follows for two solutions x and y.

1. If both x and y are feasible, select based on crowded comparison operator.

2. If x is feasible and y is not, or vice versa, select the feasible one.

3. If both x and y are infeasible:

   (a) select one with minimum f_2.

   (b) if f_2 is equal, select one with minimum f_1.

   (c) if f_1 is also equal, use crowding distance metric.

4. Use arbitrary selection for any unresolved case.

Using this selection procedure, we can initially direct the search towards the feasible region of OP and thereafter concentrate on exploring the trade-off characteristics. Step 3 of the procedure uses a lexicographic approach to selection. Note that the space of possible solutions to the problem is increasingly dense as the effective k approaches 1, i.e. a relatively higher number of solutions are feasible for lower k_{pref}. Step 3b and 3c are particularly useful in such settings. Also, the feasibility check here involves determining the size of the smallest equivalence class and whether it is at least k_{pref}. This check must be appropriately modified when considering a privacy model different from k–anonymity. This involves determining the value of the privacy parameter (e.g. k in k–anonymity or ℓ in ℓ–diversity) resulting from using a generalization, and checking whether it satisfies the minimum threshold set by the data publisher.

Table 4.3: Attributes and domain size from the *adult census* data set.

Attribute	Domain Size
Age (*age*)	20 (granularity=5)
Work Class (*wkc*)	7
Education (*edc*)	16
Marital Status (*mst*)	7
Occupation (*occ*)	14
Race (*rac*)	5
Gender (*gen*)	2
Native Country (*ncy*)	41
Salary Class (*slc*)	2

4.5.2 Solution to OP

Once the final non-dominated solution set $\mathcal{ND}$ to MOOP is obtained, the solution to OP is chosen as the point D' such that $D' = \underset{D'' \in \mathcal{ND}_f}{\operatorname{argmin}} f_1(D'')$, where $\mathcal{ND}_f = \{D_i \in \mathcal{ND} | f_2(D_i) \leq 0\}$. The case of multiple such solutions is resolved using the preference deviation metric. Since the minimum of **ach** obtained in this manner is only justifiable w.r.t. ND_f, we shall say that D' is an efficient solution only w.r.t. the non-dominated solutions generated by NSGA-II. Hence, although a global minimum of **ach** is guaranteed to be an efficient solution (w.r.t. the entire efficiency space) in theory, the NP-hard nature of the problem prevents us from claiming that the solution found is indeed a true minimum. The statement is just a cautious side note to indicate that the effectiveness of the approach is as good as the convergence and diversity preservation abilities of the multi-objective optimizer.

4.6 Empirical Results

We applied the NSGA-II methodology to the "adult.data" benchmark data set. The attributes used in this study along with their domain size are listed in Table 4.3. Recall that each attribute requires (*domain size - 1*) bits to represent all possible generalizations. This gives us a chromosome of length 105 bits representing a solution. For NSGA-II, the population size N_{pop} is set at 100 with a maximum of 50,000 function evaluations. Binary crossover is performed on the entire chromosome with rate 0.8. Mutation is performed on the individual encodings of every attribute with a rate of 0.001. The modified selection

Figure 4.3: NWP and NECD values of non-dominated solutions returned by NSGA-II for different k_{pref} (effective k value obtained is highlighted).

operator is used for binary tournament selection. Weights on the attributes are assigned equally (1/9), unless otherwise stated.

4.6.1 Solution efficiency

Fig. 4.3 illustrates the NWP and NECD values of the non-dominated solutions returned by NSGA-II for different values of k_{pref}. The preference point of $NWP_{pref} = 0.2$ and $NECD_{pref} = 1.0$ is used in these experiments. Choosing a NECD preference of 1.0 effectively allows NSGA-II to look for low NWP solutions irrespective of the privacy bias they induce. As higher values of k_{pref} are used, the number of feasible solutions obtained decreases. This is likely to happen since the search space is known to be very dense for low values of k_{pref}, while solutions become rare as higher privacy requirements are enforced. Consequently, while reported solutions for $k_{pref} = 2, 5$ and 10 have an effective k close to k_{pref}, higher values are obtained for $k_{pref} = 25, 50$ and 100. However, higher information loss has to be sustained for stronger privacy requirements. A unique feasible minimum of **ach** is obtained in all the cases. In confirmation to our theoretical observation, the minimum point is a non-dominated point in the NWP vs. NECD objective space w.r.t other feasible solutions returned by NSGA-II. Further, the existence of solutions at effective k values higher than k_{pref} (for example $k = 3$ for $k_{pref} = 2$) strengthens our claim that the optimal solution need not always have effective k value equal to k_{pref}. Once again, the concept of Pareto-dominance helps here in discovering these solutions.

4.6.2 Impact of population size

Fig. 4.4 shows the reported solutions for three different settings of population size, $N_{pop} = 100, 250$ and 500. Notice that increasing the population size, while keeping the number of function evaluations fixed, seems to have only marginal impact on the overall solution quality. Solutions are slightly less effective in terms of the preference deviation metric for larger population size, albeit there is no logical pattern in the behavior. Larger populations typically have the potential to explore more parts of the search space. However, the absence of a uniform distribution of solutions in the search space makes this exploration difficult. We also notice that the number of unique solutions obtained is sim-

Figure 4.4: Impact of population size (N_{pop}) on solution quality for $k_{pref} = 5$ and preference point $NWP_{pref} = 0.2; NECD_{pref} = 1.0$.

ilar irrespective of the population size used. Large populations have more duplicates that affect the convergence rate of the population. This is primarily due to the higher selective pressure of duplicate solutions, which limits the exploratory capabilities of the population. Small populations and higher number of iterations is a key element in solving this problem.

4.6.3 Effect of weight vector

Fig. 4.5 illustrates the solutions obtained for different assignment of weights to the attributes. As is evident from the solutions, the assignment of equal weights (wv_1) in this problem results in a much higher NWP and NECD. Weight assignments impact the amount of generalization that may be performed for an attribute, which in turn influence the information content of the anonymized data set. Even when all attributes are equally important, higher weights can be assigned to attributes with larger domain sizes to retain as much information as possible. For example, while most solutions in the figure completely suppress (number of partitions = 1) the "native country" attribute, assigning a higher weight to the attribute (as in wv_2 and wv_5) return solutions with more number of partitions. In general, NSGA-II is seemingly effective in generating solutions with higher number of partitions in accordance with the weight assignments. The "age" attribute seems to have some correlation with the other attributes as the generalization performed

Figure 4.5: Effect of the weight vector on attainable NWP and NECD. Also shown are the corresponding number of groupings in the attribute domains.

on it is low even if most of the weight is distributed on other attributes. This experiment with weight vectors provides us with another example (wv_4) demonstrating that the optimal solution need not always be present at $k = k_{pref}$.

4.6.4 Impact of bias preference

Fig. 4.6 illustrates the impact of setting the NECD preference value. A typical preference of 1.0 effectively means that any level of bias is acceptable. As a result, a solution only needs to perform as much generalization as is necessary to meet the feasibility constraint, assuming that the minimum value of NWP is attained at $k = k_{pref}$. Such a case happens with the weight vector wv_2. However, when the bias preference is dropped below 0.1, solutions are generated with higher NWP (although within the preference value of 0.2) and higher effective k. This happens because the method is now forced to explore solutions with more generalization in order to better meet the low bias preference. More generalization typically yield higher effective k. Notice that as the bias preference is lowered, the effective k increases. It is imperative to ask at this point why a bias preference of 1.0 should not be set for this problem since the best solution (with $k = 5$) is obtained with

Figure 4.6: Impact of bias preference on solutions obtained with weights wv_2.

this setting. The answer lies in the trade-off characteristic of the solutions between the level of privacy and NWP. Note that the $k = 6$ solution has higher NWP at the expense of slightly higher privacy level than the $k = 5$ solution. Since both solutions meet the NWP preference, the $k = 6$ solution is more preferable. In fact, given the four solutions in the figure and the NWP preference of 0.2, the $k = 7$ solution (one marked with a circle) is the solution of choice. This solution overachieves the preference criteria and provides better privacy than the $k = 5$ and $k = 6$ solutions. In general, specifying a very high bias preference may prohibit the method from exploring the trade-off characteristics between privacy level and NWP.

4.6.5 Efficiency

Evolutionary algorithms often receive criticism for their high running time. NSGA-II takes around 15 minutes to complete the **ach** minimization problem on the test dataset. This can be reduced by temporarily storing evaluated points. The reference direction procedure could require multiple calls to this optimization routine depending on how well a solution meets the data publisher preferences, or how extensively the data publisher explores the neighborhood solutions. However, this is typically an offline problem. Further, evolutionary algorithms are inherently parallel and can easily be adapted to utilize the processing power of today's massively parallel systems [9, 126, 121], thereby signifi-

cantly improving the run time. As a rough estimate, evolutionary algorithms in today's multi-core systems can perform 5000 function evaluations in the same amount of time it took to perform one evaluation on a uni-core system ten years ago. The proliferation of grid-based systems can increase this factor by twenty folds in a short span of time. Nonetheless, the evolutionary algorithm is just one technique to minimize the **ach** function. It is worth investigating if existing search methods proposed for loss minimization in a generalization lattice can be adapted to minimize an achievement function instead. This would make existing methods equally suitable for finding privacy preserving generalizations that also adhere to data publisher preferences.

4.7 Conclusions

In this chapter, we explore the problem of privacy bias minimization along with data utility maximization in the space of data generalizations that satisfy a particular privacy property. In addition, we also emphasize that data publisher preferences are an important component in this optimization problem. As a possible solution methodology, we propose using scalarizing functions based on preferences of the data publisher to transform the vector optimization problem to a scalar one. Strictly order preserving and strictly order representing properties in such a function guarantee that a solution obtained by minimizing the scalar function is also efficient in terms of the vector problem.

Minimization of the scalar function is performed by transforming the privacy constraint into a second objective, thereby providing a method to improve upon the specified privacy levels, if possible. The multi-objective problem is solved using an evolutionary algorithm. Moreover, a reference direction based interactive procedure iteratively uses this algorithm to help a data publisher explore efficient solutions until a satisfactory one is found. Results on a benchmark data set demonstrate the effectiveness of the evolutionary algorithm in finding solutions that best achieve the preferences of the data publisher. The method is also able to find higher effective k values depending on the weights assigned to different attributes.

It would be interesting to see how different notions of homogeneity in privacy levels can be used to define bias metrics, and what impact they have on the information preser-

vation efficiency of a generalization. The efficacy of minimality attacks when generalizations are based on preferences is also a direction worth exploring. Given the advantages of the proposed concepts, it will be important to reduce the runtime complexity of the approach. Currently, most of the overhead is imposed by the evolutionary algorithm. It is not yet clear if a more guided heuristic search can be performed on the space of generalizations.

CHAPTER 5

Comparing Disclosure Control Algorithms

A typical disclosure control algorithm searches over the space of anonymizations satisfying a particular privacy model, seeking the one with highest utility. An implicit assumption in such optimization attempts is that all anonymizations satisfying a particular privacy property fare equally well in preserving the privacy of individuals. For example, in the *k*-anonymity model where the measure of privacy is given by the size of the minimum equivalence class in the anonymized data set, two anonymizations of the same data set achieving the same value of *k* will be considered equally good with respect to privacy protection. Comparative studies based on such an assumption ignore the fact that an anonymization can introduce unwanted bias towards a certain fraction of the individuals represented in the data set. This bias stems from the fact that current privacy models offer only a collective representation of the level of privacy, resulting in higher privacy for some individuals and minimalistic for others. Under such a scenario, the results of comparing the effectiveness of various anonymizations can be misleading.

Let us consider the data set T_1 shown in Table 5.1. The data set contains 10 tuples with 3 attributes in each. Table 5.2 show two possible 3-anonymous (T_{3a} and T_{3b}) generalizations of T_1. The level of privacy inferred from T_{3a} and T_{3b} is based on 3-anonymity which is essentially the minimum size of an equivalence class. The notion of privacy assumed here is in a minimalistic sense, meaning that every tuple has at most a $\frac{1}{3}$ probability of

Table 5.1: Hypothetical microdata.

	Zip Code	Age	Marital Status
1	13053	28	CF-Spouse
2	13268	41	Separated
3	13268	39	Never Married
4	13053	26	CF-Spouse
5	13253	50	Divorced
6	13253	55	Spouse Absent
7	13250	49	Divorced
8	13052	31	Spouse Present
9	13269	42	Separated
10	13250	47	Separated

T_1 :

privacy breach. Thus, both T_{3a} and T_{3b} are considered to have the same level of privacy. However, we argue that T_{3b} should rightfully be evaluated as providing better privacy. This is because tuples {2,3,5,6,7,9,10} in T_{3b} has $\frac{1}{7}$ probability of breach, lower than their counterparts in T_{3a}. In general, with models like *k*-anonymity and others based on equivalence class sizes, such subtle information is likely to be lost. This is because these privacy measurements are based on an aggregate property of the anonymization, namely the minimum equivalence class size in this case. What differentiates T_{3a} and T_{3b} is the data utility factor which in some sense is orthogonal to the privacy requirement. From a data utility perspective T_{3a} is perhaps better since the attributes "Zip code" and "age" are less generalized in T_{3a} than in T_{3b}. Thus, either of T_{3a} or T_{3b} can be preferable depending on a higher utility or a better privacy requirement.

In this chapter, we focus on identifying ways of comparing anonymizations when such bias is known to exist. We reject a categorical statement such as "4-anonymity is better than 3-anonymity," but seek alternative ways of comparing anonymizations. Towards this end, we introduce a vector-based representation of privacy to address the problem of bias that is induced by the scalar representation. Each property, such as privacy or utility, is associated with a *property vector*, where each element gives a measure of the property for an individual anonymized tuple. Such a representation would signify, for example, the privacy level present for every individual in the data set under a particular privacy model. This will not only allow one to capture the anonymization bias introduced by existing privacy models, but also enable one to perform comparisons between various

Table 5.2: Two 3-anonymous generalizations of T$_1$. Real values of marital status are shown in italics. Left table is denoted as T$_{3a}$ and right table as T$_{3b}$.

	Zip Code	Age	Marital Status
1	1305*	[25,35]	Married (*CF-Spouse*)
4	1305*	[25,35]	Married (*CF-Spouse*)
8	1305*	[25,35]	Married (*Spouse Present*)
2	1326*	[35,45]	Not Married (*Separated*)
3	1326*	[35,45]	Not Married (*Never Married*)
9	1326*	[35,45]	Not Married (*Separated*)
5	1325*	[45,55]	Not Married (*Divorced*)
6	1325*	[45,55]	Not Married (*Spouse Absent*)
7	1325*	[45,55]	Not Married (*Divorced*)
10	1325*	[45,55]	Not Married (*Separated*)

	Zip Code	Age	Marital Status
1	130**	[15,35]	Married (*CF-Spouse*)
4	130**	[15,35]	Married (*CF-Spouse*)
8	130**	[15,35]	Married (*Spouse Present*)
2	132**	[35,55]	Not Married (*Separated*)
3	132**	[35,55]	Not Married (*Never Married*)
5	132**	[35,55]	Not Married (*Divorced*)
6	132**	[35,55]	Not Married (*Spouse Absent*)
7	132**	[35,55]	Not Married (*Divorced*)
9	132**	[35,55]	Not Married (*Separated*)
10	132**	[35,55]	Not Married (*Separated*)

anonymizations based on the difference in distribution of the privacy levels.

We propose the notion of *quality index functions* that can be used to evaluate the effectiveness of an anonymization and then formally analyze the characteristics of such functions. An *m*-ary quality index function assigns a real number to a combination of *m* property vectors. This quantitative estimate is useful in measuring the quality of the property vector. If the quality index value for one instance of a property vector produced by an anonymization is better than another instance produced by a different anonymization, we will say that the first anonymization is preferred over the second. Unary quality index functions are limited in their ability in performing comparisons and can measure only the aggregate properties of the anonymizations. Towards this end, we explore other methods of comparison that allows one to quantify the differences in the values of the property measured across the tuples instead of a minimalistic estimate. We also present various preference based techniques when comparisons are to be made across multiple properties.

Problems similar to the ones encountered here are known to exist in the multi-objective optimization community as well. Quality assessment of solution sets in this community is often difficult because of the existence of non-dominance relationships between one or more members of two different sets. Hansen and Jaszkiewicz propose the use of quality measures that induce a linear ordering on the space of all possible solution sets [81]. Knowles and Corne [101] provide a critical overview of existing quality measures and show that most existing measures do not cater to the "ordering" requirement proposed by Hanse and Jaszkiewicz. Later work presented by Zitzler et al. explore the limitations of a comparative study done under the light of *quality indicators* [196]. We have found that a principle theorem proved in their work is equally applicable in the case of anonymization comparisons. The analysis presented in their work serves as a backbone for this study.

The remainder of the chapter is organized as follows. The idea of anonymization bias is elaborated upon in Section 5.1. Section 5.2 defines the concepts pertinent to the remaining discussion. Section 5.3 explores the limitations of performing a comparative study using strict comparisons. The requirement for other methods of comparison follows from this in Section 5.4. A number of comparators are suggested in this section. Finally,

Section 5.5 concludes the chapter with a discussion on future extensions.

5.1 Anonymization Bias

Let us revisit the data in Table 5.1. Table 5.3 shows a 4-anonymous generalization of the table. We note that further discrepancy beyond those discussed earlier is evident when we start looking at the anonymizations from a user's perspective. Typically, a 4-anonymous generalization is considered to provide higher privacy than a 3-anonymous one. Again, this idea is based on a minimalistic notion of privacy, keeping in mind the entire data set and a certain property satisfied by the tuples in it.

Table 5.3: A 4-anonymous generalization of T_1.

		Zip Code	Age	Marital Status
	1	13***	(20,40]	*
	3	13***	(20,40]	*
	4	13***	(20,40]	*
	8	13***	(20,40]	*
T_4 :	2	13***	(40,60]	*
	5	13***	(40,60]	*
	6	13***	(40,60]	*
	7	13***	(40,60]	*
	9	13***	(40,60]	*
	10	13***	(40,60]	*

However, it is worth noting that attacks on the anonymized data sets could be targeted towards a particular subset of the individuals represented in the data set. In such a situation, a user needs to be concerned about her own level of privacy, rather than that maintained collectively. For example, if user 8 is to choose between the anonymizations T_{3b} and T_4 , the choice would be the latter which conforms to our understanding that 4-anonymity is better. However, if user 3 is in question then the 3-anonymous generalization T_{3b} is in fact better than T_4. Fig. 5.1 plots the size of the equivalence class for each tuple in the three different generalizations. The plot tells us that different anonymizations can in fact be better for different individuals. This in some way disrupts our understanding of "better" and "poor" privacy.

These fundamental problems are often ignored while performing comparative studies. The notion of privacy assumed in most studies is limited to some overall measure,

Figure 5.1: The size of equivalence class to which a tuple in T_1 belongs to for different anonymizations. Two different anonymizations with the same collective privacy level can have different privacy levels for individual tuples.

which can result in anonymizations being biased towards some fraction of the data set. Although removing this bias can be a difficult task, no attempt is known to have been made to measure it, less provide privacy measures keeping the bias in consideration. Comparative studies typically assume that if parameters are set similarly in a privacy model, for example k in k-anonymity, then the resulting level of privacy would also be similar. Thereafter, most of the focus during optimization is directed towards obtaining higher utility. With the anonymization bias in picture, the very assumption in the first step of an optimization procedure does not hold any longer. Our work in this chapter is not targeted towards defining a new privacy model that overcomes this bias, but to find ways of comparing anonymizations when the bias is known to exist.

Note that the appearance of bias is not limited to k-anonymity alone. When individual measures of privacy are not considered, such bias can appear in any privacy model. The bias can be present even in a personalized privacy setting, such as in the model presented by Xiao and Tao [185]. Personalized privacy in such a model is achieved by constraining the probability of privacy breach for an individual, depending on personal preferences of a breach, to an upper bound. Nonetheless, the individual probabilities need not be same for all tuples, thereby biasing a generalization scheme in more favor towards some tuples

than others.

It is imperative to ask if an anonymization can be strictly better than another. It is known that privacy and utility are two conflicting facets of an anonymization, indicating that an anonymization better in one aspect (privacy or utility) is likely to suffer on the other. However, even if utility is not considered as a criteria for obtaining a better generalization, how easy is it to establish that one anonymization is better than another? The answer could be subjective depending on how one defines a "better" anonymization. We shall explore the limitations and alternatives to establish the superiority of an anonymization for various definitions in this context.

5.2 Preliminaries

Let $\Phi_1, \Phi_2, \ldots, \Phi_a$ represent the domains of a attributes. A data set of size $\mathcal{N}$ defined over these attributes is a collection of $\mathcal{N}$ tuples of the form $(\phi_1, \phi_2, \ldots, \phi_a) \in \Phi_1 \times \Phi_2 \times \ldots \times \Phi_a$. An anonymization of a data set is achieved by performing generalization/suppression of the tuples, resulting in an anonymized data set unidentifiable from the original one. Since suppression of tuples can be represented as a special case of generalization, we adhere to the term "generalization" to mean both. Further, although tuples suppressed during an anonymization are usually eliminated, we assume that they still exist in the anonymized data set in an overly generalized form. This enables us to say that both the original data set and the anonymized one are of the same size. An anonymized data set is then subjected to a variety of property measurements. A *property* here refers to a scalar quantity signifying a privacy, utility or any other measurable feature of a tuple in the anonymized data set. This gives us a vector of values representing the property value for every tuple of the anonymized data set.

Definition 5.1 PROPERTY VECTOR. A property vector $\mathcal{D}$ for a data set of size $\mathcal{N}$ is an $\mathcal{N}$-dimensional vector $(d_1, d_2, \ldots, d_{\mathcal{N}})$ with $d_i \in \mathbb{R}; 1 \leq i \leq \mathcal{N}$ specifying a measure of a property for the ith tuple of the data set.

A property signifies the grounds under which a comparison is made between two anonymizations. Consider that we are performing k-anonymization on a data set. A

generalization scheme to do so results in multiple equivalence classes, the desired property being that all such equivalence classes are of size at least k. If we pick our privacy property to be the *"size of the equivalence class to which a tuple belongs,"* then each tuple will have an associated integer. This results in a property vector $\mathcal{D} = (d_1, d_2, \ldots, d_{\mathcal{N}})$ for a data set of size $\mathcal{N}$. For example, the equivalence class property vector induced in T_{3a} is $(3,3,3,3,4,4,4,3,3,4)$. Another example of a property could be the contribution made by a tuple to the total information loss. If measurements on the diversity of sensitive attributes in an equivalence class is desired, then the property can be the number of times the sensitive attribute value of a tuple appears in its equivalence class. Considering "marital status" as a sensitive attribute, such a property vector for T_{3a} will be $(2,2,1,2,2,1,2,1,2,1)$.

Ideally, any number of properties can be studied on a given anonymization. An anonymization is only a representation of the generalization scheme, inducing different property vectors for different properties. For example, one may be interested in analyzing an anonymization w.r.t. both k-anonymity and ℓ-diversity. In such a case, the property vectors to consider are the ones generated by the properties - *size of equivalence class of a tuple* and *count of the sensitive attribute value of a tuple in its equivalence class*. For an anonymization, a property vector due to the first property relates to k-anonymity, while that from the latter one relates to ℓ-diversity. We thus use the notion of *r-property anonymization* to indicate that the set of properties decided upon for a comparison process is restricted to a pre-specified set of r properties. The objective of a comparison is to decide if one anonymization is better than another w.r.t. the specified properties.

Definition 5.2 *r*–PROPERTY ANONYMIZATION. Let Δ be the set of all data sets of size $\mathcal{N}$. Let Y be the set of all elements $v \in 2^{\mathbb{R}^{\mathcal{N}}}$ such that $|v| = r$. A r–property anonymization $\mathcal{G}$ is a function $\mathcal{G} : \Delta \rightarrow Y$ which induces a set of r $\mathcal{N}$-dimensional property vectors $\mathcal{G}(\delta)$ on a data set $\delta \in \Delta$.

Note that an *r*-property anonymization does not mean that there are restrictions on how an anonymization is done. It only indicates that, for a given anonymization, r different property vectors are chosen to proceed with the comparison. The idea is to project an anonymized data set into a set of $\mathcal{N}$-dimensional vectors with regard to r proper-

ties, and then compare the resulting vectors for different anonymization schemes. For example, the aforementioned example of analyzing the size of equivalence class and the number of sensitive attribute values of a tuple in its equivalence class will be referred to as 2–property anonymization.

Comparisons between anonymizations is done by defining a *comparator*, denoted by $\triangleright$. A comparator is an ordering operation defined on sets of property vectors. An example is the dominance-based comparator $\succeq$ – under a 1-property anonymization, say $Y_1 = \{(d_1^1, \ldots, d_{\mathcal{N}}^1)\}$ and $Y_2 = \{(d_1^2, \ldots, d_{\mathcal{N}}^2)\}$, then $Y_1 \succeq Y_2$ iff $\forall 1 \leq i \leq \mathcal{N}, d_i^1 \geq d_i^2$. In other words, comparators are user-defined ways of evaluating the superiority of a property vector. An anonymization is better than another only w.r.t. the comparator used in comparing the induced set of property vectors. Therefore, given a comparator $\triangleright$, the relation $\mathcal{G}_1 \triangleright \mathcal{G}_2 \iff Y_1 \triangleright Y_2$ is implicitly assumed to be true. Note that the definition of a comparator need not always be explicitly made in terms of the values in a property vector. For example, a comparator may be defined just to check if more values in one property vector are higher than the corresponding values in another vector. Hence, we use quality index functions on property vectors to quantitatively measure the competence of a set of property vectors.

Definition 5.3 *m*-ARY QUALITY INDEX. Let Π be the set of all property vectors w.r.t a particular property. An *m*-ary quality index $\mathcal{I}$ is a function $\mathcal{I} : \Pi^m \to \mathbb{R}$ which assigns a combination of *m* property vectors $\mathcal{D}_1, \ldots, \mathcal{D}_m$ a real value $\mathcal{I}(\mathcal{D}_1, \ldots, \mathcal{D}_m)$.

Since comparisons are usually performed by applying an anonymization on the same data set, we shall restrict Π to be the set of all property vectors of the same size, i.e. the size $\mathcal{N}$ of the data set. Based on the same reasoning, a quality index may also use the original data set while mapping property vectors to real numbers.

A commonly used method of performing comparisons is through unary quality index functions (1-ary). Unary quality indices are functions applied independently on anonymizations. They measure one or more feature (privacy/utility) of an anonymization, and the quantitative value is considered representative of the measured feature of the anonymization. For example, *k*-anonymity is an unary quality index on the equiv-

alence class size property vector, given as $\mathcal{I}_{k-anon}(\mathcal{D}) = \min_{d_i}(\mathcal{D})$ ($= 3$ for T_{3a}). Another possible quality index could be the average size of the equivalence class maintained in the anonymized data set, i.e. $\mathcal{I}_{avg}(\mathcal{D}) = \sum d_i / \mathcal{N}$ ($= 3 \times 3 + 3 \times 3 + 4 \times 4 / 10 = 3.4$ for T_{3a}). Other models like ℓ-diversity and t-closeness result in other property vectors depending on the property being measured, for example ℓ-diversity uses a count of the number of times the sensitive attribute value of a tuple is represented in an equivalence class. With this property, we shall have a unary quality index $\ell = \mathcal{I}_{\ell-div}((2,2,1,2,2,1,2,1,2,1))$ value of 1 for T_{3a} (considering "marital status" as the sensitive attribute). Once again, the ℓ or t values for an anonymization is a quality measurement on the property vectors, the minimum values in these models. Certain forms of utility measurements can also be captured from property vectors. A loss measurement, such as the general loss metric [92], computes a normalized loss quantity for every tuple of the data set. For such metrics, a property vector specifies the loss resulting from each tuple in the data set. Thereafter, the quality index is some form of aggregation of the individual losses.

Note that unary indices only allow the measurement of an aggregate property of an anonymization. This limits any kind of comparison against the bias that may be present across anonymizations. More specifically, comparisons are not possible across the property values maintained by a tuple in two different anonymizations. This problem is eliminated by binary indices (2-ary) since both anonymizations are now available to allow comparison of individual components of the induced property vectors. A binary quality index has two anonymizations as input and the real-valued output signifies a relative measure of one anonymization's effectiveness over another. For example, a binary quality index such as $\mathcal{I}_{binary}(s,t) = |\{s_i | s_i > t_i\}|$ counts the number of entries in the property vector s that has higher property values than the corresponding entries in t. Note that s and t in this case are property vectors measuring the same property in two different anonymizations. For the size of equivalence class property in T_{3a}, with property vector $s = (3,3,3,3,4,4,4,3,3,4)$, and T_{3b}, with property vector $t = (3,7,7,3,7,7,7,3,7,7)$, we have $\mathcal{I}_{binary}(s,t) = 0$ and $\mathcal{I}_{binary}(t,s) = 7$. These index values indicate that if an 1-property anonymization is analyzed w.r.t. the size of equivalence class property, then anonymization T_{3b} inducing the property vector t is preferable over T_{3a}.

Quality estimation based on index functions stresses on the fact that a particular anonymization can be interpreted with respect to many different privacy properties and utility measurements. The superiority of one anonymization to another is thus dependent on what privacy properties are taken into consideration while performing the comparison. If quality indices are used to establish this superiority, then our concern is how many of them are needed to do so.

5.3   Strict Comparisons

Most algorithms in disclosure control are designed to obtain anonymizations that can maximize the utility of the anonymized data while satisfying a pre-specified privacy property. An anonymization is considered to be better than another if it provides higher utility within the constraints of the privacy requirement. Privacy, as given by a model, and utility are two properties induced by an anonymization. A disclosure control algorithm scans through the space of anonymizations satisfying a privacy property to find the one with maximum utility. Performance of one algorithm is said to be better than another if it is able to find an anonymization with higher utility.

This form of comparison suffers from the fact that the privacy level measured from an anonymization is a scalar quantity. It is known that maximum privacy and maximum utility are orthogonal objectives that cannot be achieved at the same time. Hence, it is imperative that when an anonymization with a better utility is found by an algorithm, the privacy factor must suffer. However, this facet may not be exposed if scalar measures are used to represent privacy. The anonymization bias plays an important role here in explaining a degraded performance in privacy from high utility anonymizations. Further, optimization attempts are also rare where emphasis is laid on obtaining anonymizations that satisfy more than one privacy property.

Discrepancies of the above nature prompts us to consider vector based measurements of the properties induced by an anonymization. Our perspective of an anonymization is that of a source that induces various properties, both in terms of privacy and utility, and more importantly, the properties can be measured on each tuple in the data set. Thereafter, methods to compare these property vectors (one or more) are devised to evaluate

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Table 5.4: Strict comparators based on dominance relationships.

Comparator ($\triangle$)	$\mathcal{D}_1 \triangle \mathcal{D}_2$	$Y_1 \triangle Y_2$	$\mathcal{G}_1 \triangle \mathcal{G}_2$
Weak dominance ($\succeq$)	$\forall i; d_i^1 \geq d_i^2$	$\forall \mathcal{D}_i \in Y_1, \mathcal{D}_i' \in Y_2; \mathcal{D}_i \succeq \mathcal{D}_i'$	$\mathcal{G}_1$ is not worse than $\mathcal{G}_2$
Strong dominance ($\succ$)	$\forall i; d_i^1 \geq d_i^2$ $\land \exists j; d_j^1 > d_j^2$	$\forall \mathcal{D}_i \in Y_1, \mathcal{D}_i' \in Y_2; \mathcal{D}_i \succeq \mathcal{D}_i'$ $\land \forall \exists \mathcal{D}_j \in Y_1, \mathcal{D}_j' \in Y_2; \mathcal{D}_j \succ \mathcal{D}_j'$	$\mathcal{G}_1$ is better than $\mathcal{G}_2$
Non-dominance ($\parallel$)	$\exists i,j; d_i^1 < d_i^2 \land d_j^1 > d_j^2$	$\exists \mathcal{D}_i \in Y_1, \mathcal{D}_i' \in Y_2; \mathcal{D}_i \succ \mathcal{D}_i'$ $\land \exists \forall \mathcal{D}_j \in Y_1, \mathcal{D}_j' \in Y_2; \mathcal{D}_j' \succ \mathcal{D}_j$	$\mathcal{G}_1$ and $\mathcal{G}_2$ are incomparable
User defined ($\blacktriangleright$-better)	$\mathcal{D}_1$ is $\blacktriangleright$-better than $\mathcal{D}_2$	Y_1 is $\blacktriangleright$-better than Y_2	$\mathcal{G}_1$ is $\blacktriangleright$-better than $\mathcal{G}_2$

the competence of an anonymization.

The first question we ask is the feasibility of performing *strict comparisons*. A strict comparison between property vectors follows from the concept of dominance, widely used in the multi-objective optimization community. The notions of *weak* and *strong* dominance enables us to make strong statements about the superiority of an anonymization. Weak dominance says that every measured value of a property on every tuple of the data set after an anonymization must be at least as good as the value measured from the corresponding tuples with another anonymization. This establishes a "not worse than" relationship between vectors. Strong dominance offers a stricter notion and establishes the "better than" relationship (Table 5.4). The non-dominance relationship signifies incomparable vectors. We are interested in strict comparisons based on dominance because they provide a framework to undoubtedly justify why an anonymization is better than another. It can potentially eliminate the effects of anonymization bias during a comparative study. However, the following discussion shows that adopting a dominance based comparative method could be rather impractical.

Theorem 5.1 Let $\mathcal{D}_1$ and $\mathcal{D}_2$ be two property vectors measuring the same property and induced by the 1–property anonymizations $\mathcal{G}_1$ and $\mathcal{G}_2$ respectively on a data set of size $\mathcal{N}$. If $(\mathcal{I}_1, \mathcal{I}_2, \ldots, \mathcal{I}_n)$ be a vector of n unique unary quality indices such that

$$\forall 1 \leq i \leq n : \mathcal{I}_i(\mathcal{D}_1) \geq \mathcal{I}_i(\mathcal{D}_2) \iff \mathcal{D}_1 \succeq \mathcal{D}_2, \tag{5.1}$$

then the number of indices is at least equal to the size of the data set, i.e. $n \geq \mathcal{N}$.

Proof The proof follows from the fact that the number of open hypercubes required to cover $\mathbb{R}^{\mathcal{N}}$ is finite. We shall show that if $n < \mathcal{N}$, then infinite such open hypercubes can be defined. The proof is by induction. Consider the two non-comparable property vectors $\mathcal{D}_1 = (a,b)$ and $\mathcal{D}_2 = (b,a)$ with $a,b \in \mathbb{R}$ and $n = 1$. Then either $\mathcal{I}_1(\mathcal{D}_1) \geq \mathcal{I}_1(\mathcal{D}_2)$ or $\mathcal{I}_1(\mathcal{D}_1) < \mathcal{I}_1(\mathcal{D}_2)$. This implies that either $\mathcal{D}_1 \succeq \mathcal{D}_2$ or $\mathcal{D}_2 \succeq \mathcal{D}_1$, which leads to a contradiction since $\mathcal{D}_1 \parallel \mathcal{D}_2$. Hence the theorem holds for $\mathcal{N} = 2$.

Let us suppose that the theorem holds for data sets of size $\mathcal{N} - 1$. Consider the two property vectors $\mathcal{D}_1 = (a,a,\ldots,a,c)$ and $\mathcal{D}_2 = (b,b,\ldots,b,c)$ with $a,b \in \mathbb{R}$ and $a < b$ on a

data set of size $\mathcal{N}$. Further, let us assume that there exists a combination of $n < \mathcal{N}$ unary quality indices satisfying the relation in the theorem.

We first show that $\forall 1 \leq i \leq n : \mathcal{I}_i(\mathcal{D}_1) < \mathcal{I}_i(\mathcal{D}_2)$.

If $\exists j$ such that $\mathcal{I}_j(\mathcal{D}_1) > \mathcal{I}_j(\mathcal{D}_2)$ then $\mathcal{D}_2 \not\succeq \mathcal{D}_1$ which results in a contradiction.

Next, consider a property vector $\mathcal{D} \in \mathcal{D}_c = \{(d_1, d_2, \ldots, d_{\mathcal{N}-1}, c) | \forall 1 \leq i \leq \mathcal{N} - 1 : a < d_i < b\}; c \in \mathbb{R}$. We then have $\mathcal{D}_2 \succeq \mathcal{D} \succeq \mathcal{D}_1$ leading to $\forall 1 \leq i \leq n : \mathcal{I}_i(\mathcal{D}_2) \geq \mathcal{I}_i(\mathcal{D}) \geq \mathcal{I}_i(\mathcal{D}_1)$. If $\exists j$ such that $\mathcal{I}_j(\mathcal{D}_2) = \mathcal{I}_j(\mathcal{D}_1)$, then $\mathcal{I}_j(\mathcal{D}_2) = \mathcal{I}_j(\mathcal{D}) = \mathcal{I}_j(\mathcal{D}_1)$. With this we can now prove that the number of indices required for data sets of size $\mathcal{N} - 1$ can be less than $\mathcal{N} - 1$, contrary to the hypothesis assumed. To do so, we consider two property vectors $\mathcal{D}_p = (p_1, p_2, \ldots, p_{\mathcal{N}-1})$ and $\mathcal{D}_q = (q_1, q_2, \ldots, q_{\mathcal{N}-1})$ on a data set of size $\mathcal{N} - 1$, such that $\forall 1 \leq i \leq \mathcal{N} - 1 : a < p_i \leq q_i < b$. Hence, $\mathcal{D}_q \succeq \mathcal{D}_p$. Next, we expand the two vectors by concatenating the same arbitrary element $c \in \mathbb{R}$, the concatenated vectors being denoted by $\mathcal{D}_{p|c}$ and $\mathcal{D}_{q|c}$ respectively. Since $\forall 1 \leq i \leq n : \mathcal{I}_i(\mathcal{D}_{q|c}) \geq \mathcal{I}_i(\mathcal{D}_{p|c}) \iff \mathcal{D}_{q|c} \succeq \mathcal{D}_{p|c}$ and $(\mathcal{D}_q \succeq \mathcal{D}_p) \iff (\mathcal{D}_{q|c} \succeq \mathcal{D}_{p|c})$, we have $\forall 1 \leq i \leq n : \mathcal{I}_i(\mathcal{D}_{q|c}) \geq \mathcal{I}_i(\mathcal{D}_{p|c}) \iff \mathcal{D}_q \succeq \mathcal{D}_p$. Also, using the result that $\mathcal{I}_j(\mathcal{D}_2) = \mathcal{I}_j(\mathcal{D}) = \mathcal{I}_j(\mathcal{D}_1)$ for any $\mathcal{D} \in \mathcal{D}_c$, we have $\mathcal{I}_j(\mathcal{D}_{q|c}) = \mathcal{I}_j(\mathcal{D}_{p|c})$. Hence the result of $\mathcal{I}_j$ can be ignored and one can write $\forall 1 \leq i \neq j \leq n : \mathcal{I}_i(\mathcal{D}_{q|c}) \geq \mathcal{I}_i(\mathcal{D}_{p|c}) \iff \mathcal{D}_q \succeq \mathcal{D}_p$, which means that less than $\mathcal{N} - 1$ (recall $n < \mathcal{N}$) quality indices can be used to compare $\mathcal{D}_p$ and $\mathcal{D}_q$. Since this is contrary to the assumed hypothesis, the existence of such j is not possible.

Thus, $\forall 1 \leq i \leq n : \mathcal{I}_i(\mathcal{D}_1) < \mathcal{I}_i(\mathcal{D}_2)$.

Let us now consider the open hyperrectangle $\mathcal{I}_c = \{(r_1, r_2, \ldots, r_n) \in \mathbb{R}^n | \forall 1 \leq i \leq n : \mathcal{I}_i(\mathcal{D}_1) < r_i < \mathcal{I}_i(\mathcal{D}_2)\}$. Also, for $f > c$, $\mathcal{I}_c \cap \mathcal{I}_f = \phi$; if $\exists (r_1, r_2, \ldots, r_n) \in \mathcal{I}_c \cap \mathcal{I}_f$ then $\forall 1 \leq i \leq n : [\mathcal{I}_i((a, a, \ldots, a, c)) < r_i < \mathcal{I}_i((b, b, \ldots, b, c))] \wedge [\mathcal{I}_i((a, a, \ldots, a, f)) < r_i < \mathcal{I}_i((b, b, \ldots, b, f))]$, giving $\forall 1 \leq i \leq n : \mathcal{I}((a, a, \ldots, a, f)) < \mathcal{I}_i((b, b, \ldots, b, c))$, and implying that $(b, b, \ldots, b, c) \succeq (a, a, \ldots, a, f)$ which is absurd. Hence, since $\mathbb{R}$ is uncountable and c is chosen arbitrarily in $\mathbb{R}$, there are uncountably many disjoint open hyperrectangles in the n-dimensional quality index space $\mathbb{R}^n$. This is in contradiction to the fact that $\mathbb{R}^n$ contains countably many disjoint open hyperrectangles. Therefore, $n \not< \mathcal{N}$ which implies $n \geq \mathcal{N}$. $\square$

A similar proof can be given for the case when strong dominance has to be inferred between two property vectors, i.e. for the equivalence relation $[\forall 1 \leq i \leq n : \mathcal{I}_i(\mathcal{D}_1) \geq$

$\mathcal{I}_i(\mathcal{D}_2) \wedge \exists j \in \{1,\ldots,n\} | \mathcal{I}_j(\mathcal{D}_1) > \mathcal{I}_j(\mathcal{D}_2)] \iff \mathcal{D}_1 \succ \mathcal{D}_2$ to hold, n must be at least $\mathcal{N}$.

An important question here is whether all possible $\mathcal{N}$-dimensional property vectors are valid given a specific privacy measurement and a specific data set. The answer is a strict no. For example, if we consider the size of the equivalence class as the privacy property, one commonly used in models like k-anonymity and ℓ-diversity, we would find that the measurements are dependent. In other words, if a tuple belongs to an equivalence class of size s, then there exists $s-1$ other tuples that belong to the same equivalence class. Hence the measurement (size of equivalence class) will be the same for all s tuples. This restricts us from attaining all possible property vectors. This then raises the question if Theorem 5.1 is valid when the set of possible property vectors is actually a subset of the set of all possible property vectors. We show that the theorem in fact holds for such a subset as well.

Corollary 5.1 Let Π be the set of all property vectors that can be defined for a data set of size $\mathcal{N}$. Let $\mathcal{D} \subseteq \Pi$ be a set of property vectors measuring a particular property such that $\forall \mathcal{D}_1, \mathcal{D}_2 \in \mathcal{D}$,

$$\forall 1 \le i \le n : \mathcal{I}_i(\mathcal{D}_1) \ge \mathcal{I}_i(\mathcal{D}_2) \iff \mathcal{D}_1 \succeq \mathcal{D}_2. \tag{5.2}$$

Then, $n \ge \mathcal{N}$.

Proof Let us assume that $n < \mathcal{N}$. The proof is given by constructing the maximal superset D_M of $\mathcal{D}$ on which the relationship holds. Let $a = (a_1, a_2, \ldots, a_\mathcal{N}), b = (b_1, b_2, \ldots, b_\mathcal{N}) \in \mathcal{D}$. Hence we can say, $\forall 1 \le i \le n : \mathcal{I}_i(a) \ge \mathcal{I}_i(b) \iff a \succeq b$. Also then $a, b \in D_M$. Now consider the following vectors.

- $x \in \mathcal{X} = \{(a_1 c_1, a_2 c_2, \ldots, a_\mathcal{N} c_\mathcal{N}) | \forall 1 \le i \le \mathcal{N}; c_i \ge 1\}$

- $y \in \mathcal{Y} = \{(b_1 + (a_1 - b_1)e_1, \ldots, b_\mathcal{N} + (a_\mathcal{N} - b_\mathcal{N})e_\mathcal{N}) | \forall 1 \le i \le \mathcal{N}; 0 \le e_i \le 1\}$

- $z \in \mathcal{Z} = \{(b_1/d_1, b_2/d_2, \ldots, b_\mathcal{N}/d_\mathcal{N}) | \forall 1 \le i \le \mathcal{N}; d_i \ge 1\}$

We have the following relation on these vectors: $a \succeq b \iff x \succeq a \succeq y \succeq b \succeq z$. Hence, by applying the quality indices $\mathcal{I}_i$'s on a and b one can compare two property vectors belonging to two different sets from $\mathcal{X}$, $\mathcal{Y}$ and $\mathcal{Z}$. Note that one can still not assert that

two vectors belonging to the same set ($\mathcal{X}, \mathcal{Y}$ or $\mathcal{Z}$) can be compared in the same manner. Thus, we arbitrarily choose three vectors, one each from $\mathcal{X}, \mathcal{Y}$ and $\mathcal{Z}$, and include it in D_M, i.e. $D_M = \{x, a, y, b, z\}$. Hence, given two vectors in D_M, we can find at least three other vectors that satisfy the relationship, thereby increasing the cardinality of D_M. Since the choice of the new vectors is arbitrary, we can continue the process as many times as possible, every time increasing the cardinality of D_M. Then, in the limit, D_M will become equal to Π. This contradicts the result from Theorem 5.1 saying that $n \geq \mathcal{N}$ quality indices are required to compare two property vectors in Π. Therefore, the assumed hypothesis $n < \mathcal{N}$ is incorrect, implying $n \geq \mathcal{N}$. $\square$

The situation becomes worse if comparisons are to be made against multiple properties. On first thought, it is possible to map a set of property vectors to an unique property vector of size $\mathcal{N}$ and Theorem 5.1 could suggest that the lower bound on n is therefore $\mathcal{N}$. Let $Y_p = \{\mathcal{D}_{p1}, \mathcal{D}_{p2}, \ldots, \mathcal{D}_{pr}\}$ be a set of r property vectors, where $\mathcal{D}_{pi} = (d_{i1}^p, d_{i2}^p, \ldots, d_{i\mathcal{N}}^p)$; $1 \leq i \leq r$. Consider the vector $d_j \in D_j = \{(d_{1j}^p, d_{2j}^p, \ldots, d_{rj}^p) \in \mathbb{R}^r\}$ for some $j \in \{1, \ldots, \mathcal{N}\}$. Note that $|D_j| = \mathbb{R}^r$ and since the cardinality of $\mathbb{R}^r$ and $\mathbb{R}$ is the same, there exists a bijective function $f_j : D_j \to \mathbb{R}$. Now define the function $\mathcal{F} : Y \to \mathbb{R}^{\mathcal{N}}$ which maps each set of r property vectors to a vector of size $\mathcal{N}$ as $\mathcal{F}(Y_p \in Y) = (f_1(d_1), f_2(d_2), \ldots, f_{\mathcal{N}}(d_{\mathcal{N}}))$. Since every f_j is a bijective function, $\mathcal{F}$ is bijective too. However, a bijective mapping is not sufficient to imply the equivalence relation $\mathcal{F}(Y_1) \succeq \mathcal{F}(Y_2) \iff Y_1 \succeq Y_2$ and hence the lower bound ($n \geq \mathcal{N}$) given by Theorem 5.1 would be incorrect. It can be shown that for the relation to hold, n should at least be equal to $r\mathcal{N}$. The following corollary gives this lower bound on the number of quality indices required to compare two sets of property vectors. Note that the corollary uses notions of quality index functions extended to sets of property vectors.

Corollary 5.2 Let $Y_1 = \{\mathcal{D}_{11}, \mathcal{D}_{12}, \ldots, \mathcal{D}_{1r}\}$ and $Y_2 = \{\mathcal{D}_{21}, \mathcal{D}_{22}, \ldots, \mathcal{D}_{2r}\}$ be two sets of property vectors induced by the r–property anonymizations $\mathcal{G}_1$ and $\mathcal{G}_2$ respectively on a data set of size $\mathcal{N}$. If $(\mathcal{I}_1, \mathcal{I}_2, \ldots, \mathcal{I}_n)$ be a vector of n unary quality indices such that

$$\forall 1 \leq i \leq n : \mathcal{I}_i(Y_1) \geq \mathcal{I}_i(Y_2) \iff Y_1 \succeq Y_2, \tag{5.3}$$

then $n \geq r\mathcal{N}$.

Proof Let us assume that there exists quality indices $\mathcal{I}_i; 1 \leq i < r\mathcal{N}$ such that the equivalence relation holds. Now, $Y_1 \succeq Y_2 \iff \forall 1 \leq j \leq r; \mathcal{D}_{1j} \succeq \mathcal{D}_{2j}$. If $\mathcal{D}_{pj} = (d_{j1}^p, d_{j2}^p, \ldots, d_{j\mathcal{N}}^p)$ then $\mathcal{D}_{1j} \succeq \mathcal{D}_{2j} \iff \forall 1 \leq z \leq \mathcal{N}; d_{jz}^1 \geq d_{jz}^2$. Therefore, $Y_1 \succeq Y_2 \iff \forall 1 \leq j \leq r; \forall 1 \leq z \leq \mathcal{N}; d_{jz}^1 \geq d_{jz}^2$. By transitivity, we then have $\forall 1 \leq i < r\mathcal{N} : \mathcal{I}_i(Y_1) \geq \mathcal{I}_i(Y_2) \iff \forall 1 \leq j \leq r; \forall 1 \leq z \leq \mathcal{N}; d_{jz}^1 \geq d_{jz}^2$.

Let us now consider a data set of size $r\mathcal{N}$. For two property vectors $\mathcal{D}_1$ and $\mathcal{D}_2$ defined on this data set, $\mathcal{D}_1 \succeq \mathcal{D}_2$ if and only if every component of $\mathcal{D}_1$ is greater than or equal to the corresponding component of $\mathcal{D}_2$. We first divide a property vector on this data set into r equal sections, thereby resulting in r $\mathcal{N}$-dimensional vectors. The quality indices $\mathcal{I}_i$ can then be applied to these resulting vectors. The equivalence relation $\forall 1 \leq i < r\mathcal{N} : \mathcal{I}_i(Y_1) \geq \mathcal{I}_i(Y_2) \iff \forall 1 \leq j \leq r; \forall 1 \leq z \leq \mathcal{N}; d_{jz}^1 \geq d_{jz}^2$ can then be used to state that less than $r\mathcal{N}$ quality indices can be used to establish a dominance relation between $\mathcal{D}_1$ and $\mathcal{D}_2$. This contradicts the result from Theorem 5.1 which states that we would require at least $r\mathcal{N}$ quality indices to compare any two property vectors on a data set of size $r\mathcal{N}$. Hence, no such combination of quality indices with $n < r\mathcal{N}$ can exist. $\square$

The results till this point indicate that it is rather impractical to determine the superiority of an anonymization based on notions of weak or strong dominance, and with unary quality indices. This prompts us to consider other methods of representing the quality of anonymizations relative to one another by defining $\blacktriangleright$-better (read as *metric better*) comparators. For example, we can define a $\blacktriangleright_{cov}$-better comparator such that, given two equivalence size property vectors $\mathcal{D}_1$ and $\mathcal{D}_2$, $\mathcal{D}_1 \blacktriangleright_{cov} \mathcal{D}_2$ if more tuples in $\mathcal{D}_1$ have a higher equivalence class size than the corresponding number in $\mathcal{D}_2$. Another example is the $\blacktriangleright_{min}$-better comparator typically used in models such as k-anonymity – $\mathcal{D}_1 \blacktriangleright_{min} \mathcal{D}_2$ if $\min_{d_i^1}(\mathcal{D}_1) > \min_{d_i^2}(\mathcal{D}_2)$. In fact, current methods of comparison are all based on some $\blacktriangleright$-better comparator.

$\blacktriangleright$-better comparators may naturally induce the quality indices to be used to infer the relationship, for example $\mathcal{I}_{k-anon}$ is the quality index to be used to infer $\blacktriangleright_{min}$-better. However, as mentioned earlier, comparators such as $\blacktriangleright_{min}$ are limited by the fact that they are based on some aggregate property of the vectors, ignoring the anonymization bias altogether. On the other hand, $\blacktriangleright_{cov}$ is a rational candidate since the relationship is based

on the output from comparing multiple tuples in the two property vectors. Note that $\blacktriangleright_{cov}$ in fact induces a binary quality index as presented in the next section.

5.4 ▶-better Comparators

Comparisons with dominance based comparators suffer from the drawback that the number of unary quality indices required is impractically large. This also follows from our intuition that comparison of two $\mathcal{N}$-dimensional vectors cannot be accomplished with less than $\mathcal{N}$ scalar quantities without losing information. Besides this difficulty, dominance based comparison is a rather strict way of evaluating the quality of an anonymization. It is not unlikely that a property vector is not able to dominate another vector because of low property values for a minor fraction of the tuples. This effectuates to saying that the anonymization bias could be present negligibly resulting in a non-dominance relationship.

Although removing the effects of anonymization bias is hard (or perhaps impossible), methods can be devised to keep it in consideration during a comparative study. We therefore seek other comparators, called $\blacktriangleright$-*better comparators,* that can capture the quality of anonymizations together with the bias they introduce. $\blacktriangleright$-better comparators provide a weaker notion of superiority than dominance-based ones, nonetheless their objective is also targeted towards identifying anonymizations with better utilization of the bias.

In the following discussion, we introduce a number of $\blacktriangleright$-better comparators and the corresponding unary/binary quality index they induce. All expressions below are in terms of two property vectors $\mathcal{D}_1 = (d_1^1, d_2^1, \ldots, d_{\mathcal{N}}^1)$ and $\mathcal{D}_2 = (d_1^2, d_2^2, \ldots, d_{\mathcal{N}}^2)$ measuring a given property when induced by two different anonymizations on a data set of size $\mathcal{N}$. Without loss of generality, we assume that a higher value of a property measurement for a tuple is better.

5.4.1 $\blacktriangleright_{rank}$-better

Consider the $\mathcal{N}$-dimensional space of all property vectors for a given property. Let $\mathcal{D}_{max}$ be a point of interest and all points are assigned a *rank* based on their distance from $\mathcal{D}_{max}$. We then say $\mathcal{D}_1 \blacktriangleright_{rank} \mathcal{D}_2$ if the rank of $\mathcal{D}_1$ is lower than the rank of $\mathcal{D}_2$. A

Figure 5.2: The rank-based $\blacktriangleright_{rank}$-better comparator assigns ranks to property vectors based on the distance from a point of interest $\mathcal{D}_{max}$.

visualization of this on a 2-dimensional space is depicted in Fig. 5.2. The point $\mathcal{D}_{max}$ here is the property vector that is the most desired one, quite often the property vector that offers the maximum measure of the property for every tuple in the data set. The arcs surrounding $\mathcal{D}_{max}$ are the locus of points at the same distance from $\mathcal{D}_{max}$. Note that any two points on the same arc are incomparable vectors and are assigned the same rank. Points on two different arcs are compared based on how close they are to achieving the most desired property vector. The *rank-based* unary quality index

$$\mathcal{I}_{rank}(\mathcal{D}_1) = \| \mathcal{D}_1 - \mathcal{D}_{max} \| \tag{5.4}$$

can then be used to infer the relationship $\mathcal{I}_{rank}(\mathcal{D}_1) < \mathcal{I}_{rank}(\mathcal{D}_2) \iff \mathcal{D}_1 \blacktriangleright_{rank} \mathcal{D}_2$. It is also possible to associate a tolerance level ϵ to the rank such that two property vectors differing in rank by ϵ or less are considered equally good.

The direct implication of equi-ranked property vectors is that two different anonymizations are equivalent in terms of their ability to pursue the most desirable levels of the property being measured. In other words, the amount of bias each has to overcome (or introduce) to reach the desirable level is equivalent. In a comparative setting, the rank of a property vector can be viewed as an estimate of the bias present in an anonymization w.r.t. a particular property.

Figure 5.3: Computation of the quality index functions based on the $\blacktriangleright_{cov}$ and $\blacktriangleright_{spr}$ comparators. $\mathcal{I}_{cov}$ is based on the number of tuples with better property value while $\mathcal{I}_{spr}$ is based on the actual difference in magnitude of the measured property value.

5.4.2 $\blacktriangleright_{cov}$-better

The *coverage* comparator $\blacktriangleright_{cov}$ compares two property vectors based on the fraction of tuples in one that has a better measurement of the property than in the other. This comparator follows from the intuition that an anonymization better than another should be able to retain higher values of the measured property for more individuals represented in the data set. With this comparator and the equivalence class size property, the aforementioned anonymization T_4 is $\blacktriangleright_{cov}$-better than T_{3a}, and T_{3b} is $\blacktriangleright_{cov}$-better than T_4. The quality index induced from this comparator is binary in nature, given as

$$\mathcal{I}_{cov}(\mathcal{D}_1, \mathcal{D}_2) = \frac{|\{d_i^1 | d_i^1 \geq d_i^2\}|}{\mathcal{N}} \tag{5.5}$$

and satisfying $\mathcal{I}_{cov}(\mathcal{D}_1, \mathcal{D}_2) > \mathcal{I}_{cov}(\mathcal{D}_2, \mathcal{D}_1) \iff \mathcal{D}_1 \blacktriangleright_{cov} \mathcal{D}_2$. Note that if $\mathcal{I}_{cov}(\mathcal{D}_1, \mathcal{D}_2) = 1$ and $\mathcal{I}_{cov}(\mathcal{D}_2, \mathcal{D}_1) = 0$, then $\mathcal{D}_1 \succ \mathcal{D}_2$, and vice versa. Fig. 5.3 shows the computation of the coverage-based quality index for two hypothetical property vectors.

The coverage comparator is useful when two anonymizations demonstrate similar levels of collective privacy, for example both are *k*-anonymous for some given *k*. The comparator then identifies what fraction of the tuples is favored by the skewness in the distribution of privacy levels. A higher value of $\mathcal{I}_{cov}(\mathcal{D}_1, \mathcal{D}_2)$, compared to $\mathcal{I}_{cov}(\mathcal{D}_2, \mathcal{D}_1)$, implies that more tuples benefit from the skewed distribution of property values in one

anonymization than in the other. Such a comparison helps justify that the bias introduced by an anonymization can be useful in providing better property values for a larger fraction of the data set.

5.4.3 ▶$_{spr}$-better

The coverage comparator does not take into consideration the difference in magnitude of a measured property for a given tuple when comparing it across two different property vectors. It is possible that for two vectors $\mathcal{D}_1$ and $\mathcal{D}_2$ with $\mathcal{I}_{cov}(\mathcal{D}_1,\mathcal{D}_2) \approx \mathcal{I}_{cov}(\mathcal{D}_2,\mathcal{D}_1)$ (the quality index values are close), $\mathcal{D}_1$ maintains much better values in the property for the tuples on which it is superior than $\mathcal{D}_2$, compared to those on which $\mathcal{D}_2$ is superior. For example, consider the hypothetical property vectors $\mathcal{D}_1 = (2,2,3,4,5)$ and $\mathcal{D}_2 = (3,2,4,2,3)$. In this case, $\mathcal{I}_{cov}(\mathcal{D}_1,\mathcal{D}_2) = \mathcal{I}_{cov}(\mathcal{D}_2,\mathcal{D}_1) = \frac{3}{5}$. However, one can argue that $\mathcal{D}_1$ is a superior vector since the difference in magnitude of the measured property is higher (a value of 2) in the two tuples where it is better than $\mathcal{D}_2$. This difference is only 1 in the two tuples where $\mathcal{D}_2$ is better. The *spread* comparator ▶$_{spr}$ is based on the total amount of variation (or spread) present between tuples w.r.t. a property. The quality index based on this comparator is given as

$$\mathcal{I}_{spr}(\mathcal{D}_1,\mathcal{D}_2) = \sum_{i=1}^{\mathcal{N}} \max(d_i^1 - d_i^2, 0) \tag{5.6}$$

and measures the total difference in magnitude of the measured property for the tuples on which $\mathcal{D}_1$ performs better than $\mathcal{D}_2$. We then say $\mathcal{I}_{spr}(\mathcal{D}_1,\mathcal{D}_2) > \mathcal{I}_{spr}(\mathcal{D}_2,\mathcal{D}_1) \iff \mathcal{D}_1 ▶_{spr} \mathcal{D}_2$. We also have $\mathcal{I}_{spr}(\mathcal{D}_1,\mathcal{D}_2) = 0 \iff \mathcal{D}_2 \succeq \mathcal{D}_1$. Fig. 5.3 shows the difference in computation of the coverage-based and spread-based quality indices.

The spread comparator uses a quantification of the leverage availed by individual tuples from the skewed distribution of property values. This is crucial when the fraction of tuples benefited, as given by the $\mathcal{I}_{cov}$ quality index, are equivalent. The $\mathcal{I}_{spr}$ quality index quantifies the utilization of the bias in terms of differences in observed property values. Even for the case when two anonymizations have different collective privacy levels, this quantification differentiates them further, often counter to established preferential norms. For example, consider the equivalence class size property vector $(3,3,3,5,5,5,5,5,3,3,3,4,4,4,4)$ from a 3-anonymous generalization and $(2,2,6,6,6,6,6,6,3,$

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3,3,4,4,4,4) from a 2-anonymous generalization. Both anonymizations show signs of bias towards a certain fraction of the tuples. The 3-anonymous generalization will be a typical choice when pre-defined notions of "better privacy" is used. However, the 2-anonymous generalization achieves better privacy for 6 more tuples (tuples 3 to 8) at the expense of reducing the privacy levels of tuple 1 and 2. This is a reasonable justification to choosing the 2-anonymous generalization instead. The $\mathcal{I}_{spr}$ quality index values compare at 2 and 8, thereby revealing this reasoning. In fact, the $\mathcal{I}_{cov}$ index also points at the same.

5.4.4 $\blacktriangleright_{hv}$-better

Another way of measuring the superiority of an anonymization is to ask how good it is against other possible anonymizations apart from the one it is compared to. Such a method refrains itself from performing relative comparisons and instead adopts a "tournament" style mechanism. In a tournament mechanism, a candidate a is preferred over candidate b not because a performs better than b, but because a performs better than a larger number of other candidates than the number of candidates over which b performs better.

For a given property vector $\mathcal{D}_i$, we consider the set of property vectors $\Psi_i = \{\mathcal{D}_j | \mathcal{D}_i \succeq \mathcal{D}_j\}$. Fig. 5.4 depicts this set, for a 2-dimensional space, as the volume enclosed by all property vectors which are not better than the vector in question under the $\succeq$ comparator. When performing comparisons for two property vectors, the *hypervolume* comparator $\blacktriangleright_{hv}$ assigns superiority based on the hypervolume enclosed by points which are not superior to both property vectors under the $\succeq$ comparator. In Fig. 5.4, $\mathcal{D}_1$ is $\succeq$-better than all points in region A, none of which appear in Ψ_2. Similarly, region B has all points that do not appear in Ψ_1 and region C has all points that appear in $\Psi_1 \cap \Psi_2$. Thus, the quality index

$$\mathcal{I}_{hv}(\mathcal{D}_1, \mathcal{D}_2) = \prod_{i=1}^{\mathcal{N}} d_i^1 - \prod_{i=1}^{\mathcal{N}} \min(d_i^1, d_i^2) \tag{5.7}$$

is a measurement of the hypervolume on which $\mathcal{D}_1$ is solely $\succeq$-better, giving us the relationship $\mathcal{I}_{hv}(\mathcal{D}_1, \mathcal{D}_2) > \mathcal{I}_{hv}(\mathcal{D}_2, \mathcal{D}_1) \iff \mathcal{D}_1 \blacktriangleright_{hv} \mathcal{D}_2$. Further, if $\mathcal{I}_{hv}(\mathcal{D}_1, \mathcal{D}_2) = 0$ then $\mathcal{D}_2 \succeq \mathcal{D}_1$, and vice versa. Note that the subtraction of the commonly dominated hyper-

Figure 5.4: The hypervolume comparator $\blacktriangleright_{hv}$ gives preference to property vectors that solely outperform more property vectors – $\mathcal{D}_2$ in this case since volume of region B > volume of region A.

volume is only used to signify what the quality index wishes to compute, but is otherwise not required during a comparison.

The hypervolume comparator checks the effectiveness of an anonymization w.r.t. possibly unseen anonymizations. Such anonymizations are captured in the dominated hypervolume. This expands the comparison beyond the anonymizations considered in the comparative process. A higher hypervolume for an anonymization indicates that a larger number of other anonymizations (possibly generated by other algorithms) will induce worse property values than it.

Let us consider the property vector $s = (3,3,3,5,5,5,5,5)$ induced by an anonymization S for a particular property. Let $t = (4,4,4,4,4,4,4,4)$ be the property vector induced by anonymization T for the same property. Any anonymization U inducing the property vector $u = (u_1, u_2, \ldots, u_8)$ is worse than S if $u_i < 3; i = 1,2,3$ and $u_i < 5; i = 4, \ldots, 8$. The hypervolume is a measure of such anonymizations. Similarly, U is worse than T if $u_i < 4; i = 1, \ldots, 8$. In this case, $\mathcal{I}_{hv}(s,t) > \mathcal{I}_{hv}(t,s)$ indicating that the number of possible anonymizations that is worse than S is more than that is worse than T. If the volume enclosed by all property vectors is finite, then one can also say that the number of possible anonymizations better than S is less than that is better than T. This method of looking into the effectiveness of an anonymization is complementary to the method behind $\mathcal{I}_{cov}$

and $\mathcal{I}_{spr}$. While the latter two facilitates a comparison on a one-to-one basis, comparisons via $\mathcal{I}_{hv}$ involves a broader extent of the space of property vectors.

These four quality indices can be used to compare property vectors measuring a single property only. It is true that an anonymization evaluated as being better w.r.t. a particular property can be worse in the context of another. Hence, for the case when more than one property is being measured on an anonymization, other mechanisms are required to weigh the importance of the different properties when making comparisons. We now consider sets of property vectors, instead of a single one, and suggest three indices to compare two such sets. Let us assume that comparisons are to be made across the sets $Y_1 = \{\mathcal{D}_{11}, \mathcal{D}_{12}, \ldots, \mathcal{D}_{1r}\}$ and $Y_2 = \{\mathcal{D}_{21}, \mathcal{D}_{22}, \ldots, \mathcal{D}_{2r}\}$ induced by two r–property anonymizations $\mathcal{G}_1$ and $\mathcal{G}_2$ respectively. In the following expressions, we use the notation $\mathcal{I}(X,Y)$ to indicate a binary quality index defined to compare two property vectors X and Y. Note that different quality indices can be used to compare different properties. Also, without any loss of generality, we assume that a higher value of the quality index is desirable; otherwise we can negate the index value.

5.4.5 $\blacktriangleright_{WTD}$-better

The first method to compare sets of property vectors is based on the classical weighted sum approach. Weight assignments are useful when the properties analyzed are orthogonal in nature, such as privacy and utility. The *weight-based* comparator $\blacktriangleright_{WTD}$ requires a vector $W = (w_1, \ldots w_r)$ such that the weight w_i signifies the importance of the ith property being measured. Typically the weights are chosen such that, for $1 \leq i \leq r$, $0 < w_i < 1$ and $\sum_{i=1}^{r} w_i = 1$. The quality index is given as

$$\mathcal{I}_{WTD}(Y_1, Y_2) = \sum_{i=1}^{r} [w_i \cdot \mathcal{I}(\mathcal{D}_{1i}, \mathcal{D}_{2i})] \tag{5.8}$$

and comparisons are made using the relationship $\mathcal{I}_{WTD}(Y_1, Y_2) > \mathcal{I}_{WTD}(Y_2, Y_1) \iff Y_1 \blacktriangleright_{WTD} Y_2$. It is advisable to normalize the $\mathcal{I}$ values before computing the weighted sum.

Consider the 3-anonymous generalizations in T_{3a} and T_{3b}. The size of equivalence class property vectors for the two generalizations are $p_a = (3,3,3,3,4,4,4,3,3,4)$ and $p_b = (3,7,7,3,7,7,7,3,7,7)$ respectively. Using Iyengar's data utility metric [92], the utility

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property vectors for them are $u_a = (2.03, 1.7, 1.7, 2.03, 1.6, 1.6, 1.6, 2.03, 1.7, 1.6)$ and $u_b = (2.03, 0.97, 0.97, 2.03, 0.97, 0.97, 0.97, 2.03, 0.97, 0.97)$ respectively. Using the coverage comparator we have, $\mathcal{I}_{cov}(p_a, p_b) = 0.3 < 1 = \mathcal{I}_{cov}(p_b, p_a)$ and $\mathcal{I}_{cov}(u_a, u_b) = 1 > 0.3 = \mathcal{I}_{cov}(u_b, u_a)$. Thus, if equal weights are assigned to both privacy and utility, then generalizations T_{3a} and T_{3b} are equally good.

5.4.6 $\blacktriangleright_{LEX}$-better

The weight-based comparator suffers from the drawback that it may sometimes be difficult to assign weight values specifying the preference of a property. An alternative to this is to instead specify a lexicographic ordering of the different properties. For example, when privacy levels depicted by different privacy models are to be used in conjunction to decide on an anonymization, the property vectors induced from different privacy properties can be ordered in descending order of relevance. Such a method orders the elements of a set of property vectors such that the most desirable property is the first element, the second most desirable property as the second element, and so on. With this ordering in place, we define the following quality index.

$$\mathcal{I}_{LEX}(Y_1, Y_2) = \min_i \{1 \leq i \leq r | \mathcal{I}(\mathcal{D}_{1i}, \mathcal{D}_{2i}) - \mathcal{I}(\mathcal{D}_{2i}, \mathcal{D}_{1i}) > \epsilon_i\} \tag{5.9}$$

Here, $\epsilon = (\epsilon_1, \ldots, \epsilon_r)$ is a significance vector where ϵ_i gives the maximum tolerable difference in the $\mathcal{I}$ values for the ith property. Thus, two property vectors $\mathcal{D}_{1i}$ and $\mathcal{D}_{2i}$ are considered to be equally good if $|\mathcal{I}(\mathcal{D}_{1i}, \mathcal{D}_{2i}) - \mathcal{I}(\mathcal{D}_{2i}, \mathcal{D}_{1i})| \leq \epsilon_i$. The ϵ-lexicographic comparator $\blacktriangleright_{LEX}$ assigns superiority to a set of property vectors based on the property on which it is more superior, given the ordering on the properties and the significance vector. $\mathcal{I}_{LEX}(Y_1, Y_2)$ thereby computes the first property on the ordering where Y_1 is superior. If $\mathcal{I}_{LEX}(Y_1, Y_2) < \mathcal{I}_{LEX}(Y_2, Y_1)$, then Y_1 has a more desirable property where it is superior to Y_2, i.e. $\mathcal{I}_{LEX}(Y_1, Y_2) < \mathcal{I}_{LEX}(Y_2, Y_1) \iff Y_1 \blacktriangleright_{LEX} Y_2$.

5.4.7 $\blacktriangleright_{GOAL}$-better

The *goal-based* comparator $\blacktriangleright_{GOAL}$ is useful when the competence of an anonymization can be measured in terms of its closeness to a desirable level (or goal). In such a situation,

a goal vector $G = (g_1, \ldots, g_r)$ specifies the values desired in the quality indices used – the $\mathcal{I}$ functions. The quality index

$$\mathcal{I}_{GOAL}(Y_1, Y_2) = \sum_{i=1}^{r} [\mathcal{I}(\mathcal{D}_{1i}, \mathcal{D}_{2i}) - g_i]^2 \tag{5.10}$$

then computes the sum-of-squares error of the quality indices from the goal values. The comparison is performed with the relationship $\mathcal{I}_{GOAL}(Y_1, Y_2) < \mathcal{I}_{GOAL}(Y_2, Y_1) \iff Y_1 \blacktriangleright_{GOAL} Y_2$. We can also use unary performance indices as replacements for the binary functions $\mathcal{I}$. The use of unary indices simplifies the specification of the goal vector in terms of the goal property vectors instead. If $\mathcal{D}_{g_1}, \ldots, \mathcal{D}_{g_r}$ are the goal property vectors, then the goal vector can be formulated as $G = (\mathcal{I}_1(\mathcal{D}_{g_1}), \ldots, \mathcal{I}_r(\mathcal{D}_{g_r}))$, where $\mathcal{I}_i$s are unary quality indices.

5.5   Conclusions

In this chapter, we explore an often ignored factor in the comparison of anonymizations reported by microdata disclosure control algorithms. This factor, which we call the anonymization bias, results in anonymizations being skewed towards a fraction of the data set w.r.t. a privacy property. Hence, the attainment of a collective privacy level is not sufficient to conclude that two anonymizations offer the same level of privacy. We therefore introduce property vectors as an alternate representation of a property (privacy/utility) measured on an anonymization. This representation helps indicate the privacy level of every individual in the data set separately. Further, the issue of comparing anonymizations is addressed by the usage of quality index functions that give an absolute or a relative quantitative estimate of the quality of property vectors. Our initial conclusion on such a comparative method is that unary quality indices are limited in their ability to perform comparisons, specifically when strict inferences like "not worse than" or "better than" are to be made between anonymizations. As a result, we explore alternative methods of comparison, keeping in mind that comparators defined on such grounds should make adequate effort to quantify the differences in values of the property measured across the tuples of the entire data set instead of a minimalistic estimate. Estimates based on rank, spread and hypervolume are thus suggested. We also present

various preference based techniques when comparisons are to be made across multiple properties induced by anonymizations.

An immediate extension of this work is the identification of privacy measures that address the anonymization bias. Moreover, vector-based representations of privacy would require a rethinking of the utility optimization problem since trade-offs between privacy and utility now becomes more apparent. The currently adopted optimization framework in disclosure control is single objective in nature. The scalar quantification of privacy enables the framework to direct its search for higher utility anonymization while satisfying a privacy constraint. If vector representations of privacy are adopted, then the framework has to undergo changes. This is primarily because the vector representation allows one to distinguish between anonymizations even when a typical privacy constraint is satisfied by both. Note that the current framework only makes this distinction based on utility, whereas the vector representation enables the distinction to be made at the privacy front as well. Finding "good" anonymizations thus converts into a multi-objective problem. Although the multi-objective nature of the privacy versus utility problem is well understood in the community, it has remained irrelevant under the pretext of scalar privacy representations. However, under the light of vector representations, privacy should no longer be imposed only as a constraint in the framework but rather handled directly as an objective to maximize.

CHAPTER 6

Data Privacy Through Property Based Generalizations

$\mathbf{M}$ost algorithms in disclosure control are designed to obtain anonymizations (generalization with or without suppression) that can maximize the utility of the anonymized data while satisfying a pre-specified privacy property. A specific generalization is therefore considered to be better than another if it provides higher utility within the constraints of the privacy requirement. The adoption of such an optimization framework brings forth pertinent practical issues that have been ignored for long. We have independently identified and sought resolution to three such issues in the previous chapters.

(1) *The data publisher's dilemma:* ([55, 88]) Data utility and respondent privacy are two equally important facets of data publishing. Proper anonymization thus involves weighing the risk of publicly disseminated information against the statistical utility of the content. In such a situation, it is imperative that the data publisher understands the implications of setting a parameter in a privacy model to a particular value. There is clearly a trade-off involved. Setting the parameter to a "very low" value impacts the privacy of individuals. Picking a "very high" value disrupts the inference of any significant statistical information.

(2) *Anonymity model correlations:* The k–anonymity model is prone to other forms of attacks on privacy. As a result, a multitude of other privacy models have been proposed over time [112, 120, 173, 191], quite often followed by newer forms of privacy attacks.

The inclusion of multiple models in the anonymization process is desirable since a single comprehensive model is yet to be developed. However, the correlation between these models is not well studied making the task of parameter selection a difficult one. Data generalization algorithms will also find it difficult to satisfy multiple privacy constraints when existing correlations are inverse in nature.

(3) *Biased privacy:* ([58]) Consider the $k-$anonymity model where the measure of privacy (the value of k) is given by the minimum size of an equivalence class. Thus, two anonymizations inducing the same value of k will be considered equally good with respect to privacy protection. However, it is quite possible that for one of the anonymizations a majority of the individual tuples have lesser probabilities of privacy breaches than their counterparts in the other anonymization. Individual privacy levels as depicted by such a model can therefore be misleading – higher for some, minimalistic for others.

Resolution of the first issue is a difficult task within the existing optimization framework. Any attempt to find the "best level of privacy" will require an enumeration across different parameter values to determine what is suitable. The second issue makes parameter selection further difficult owing to a larger combination of possible choices. The decision maker is also required to have an understanding of the feasibility of combining different models. The third issue is a resultant of the worst case representation of privacy. Such a representation identifies the minimum privacy present for all individuals but is inadequate to capture the amount of skewness (homogeneity or heterogeneity) introduced in individual privacy levels. Thus, the level of privacy guaranteed by a generalization can become questionable.

In this chapter, we propose resolutions to these issues using the notion of *property based generalizations*. First, inclusion of multiple objectives in the anonymization process is captured using *properties* as anonymization objectives. A generalization here is viewed as a source which can be independently subjected to evaluation in terms of different privacy or utility properties. Second, evaluation of a generalization with respect to a privacy property is performed using both worst case and vector based measurements. The overall effectiveness of a generalization is then measured with a dominance operator. The dominance relationship analyzes a generalization in terms of its achievement and

trade-offs in the different properties. The concept of a single optimal solution is therefore discarded and a representative subset of the minimal solution set is sought. Towards this end, our third contribution is in terms of an evolutionary algorithm that can be used to efficiently search the domain generalization lattice to identify such representative solutions. The algorithm uses a modified dominance operator to select these solutions.

The remainder of the chapter is organized as follows. Section 6.1 introduces property based generalizations, followed by a description of the modified dominance operator in Section 6.2. The evolutionary algorithm is presented in Section 6.3. Section 6.4 discusses the results of applying the algorithm on a benchmark data set. Finally, Section 6.5 concludes the chapter.

6.1 Property Based Generalization

Multiple objectives to meet during data anonymization are captured in the form of properties [58]. Formally, a property is defined as follows.

Definition 6.1 PROPERTY. A property is a function $\mathcal{P}$ that maps a table T to a vector of size equal to the number of tuples in the table. The vector is called a property vector and denoted by $\mathcal{P}(\mathsf{T})$.

A property refers to a privacy, utility or any other measurable feature of a tuple. It signifies the grounds under which a comparison is made between two nodes in the domain generalization lattice. Our perspective of a node is that of a source inducing multiple properties that can be measured on each tuple in the data set. For example, applying the generalization levels corresponding to a node results in multiple equivalence classes. If we pick our property to be the *"size of the equivalence class to which a tuple belongs,"* then each tuple will have an associated integer. This results in a property vector $\mathcal{P}_{equiv}(\mathsf{T}) = (k_1, k_2, \ldots, k_\mathcal{N})$ for a data set of size $\mathcal{N}$, where k_i is the equivalence class size of the ith tuple. If measurements on the diversity of sensitive attributes is required, the property can be the number of times the sensitive attribute value of a tuple appears in its equivalence class. A property is therefore a vector based measurement. The motivation behind using such vector based measurements is two folded. First, it fits the conventional

"worst case" method of measuring privacy. Second, it allows us to determine the efficiency of a node with consideration to the distribution of privacy levels across the data set. These two methods of assessing a node are jointly represented through the use of quality index functions.

6.1.1 Quality Index functions

Comparison between generalizations with respect to a single property can be done by defining an ordering operation on the co-domain of the property. Towards this end, quality index functions map a node to the set of real numbers. The underlying idea is to quantify quality differences between generalizations by applying common metrics.

Definition 6.2 QUALITY INDEX. Let $\mathcal{T}$ be the collection of all possible generalized versions of a table T. Given a property $\mathcal{P}$, a quality index $\mathcal{I}_\mathcal{P}$ is a function $\mathcal{I}_\mathcal{P} : \mathcal{T} \times \mathcal{T} \to \mathbb{R}$ which assigns an ordered pair of two tables $\mathsf{T}_l, \mathsf{T}_m \in \mathcal{T}$ a real value $\mathcal{I}_\mathcal{P}(\mathsf{T}_l, \mathsf{T}_m)$.

The value $\mathcal{I}_\mathcal{P}(\mathsf{T}_l, \mathsf{T}_m)$ signifies the quality of table T_l relative to table T_m and with respect to the property $\mathcal{P}$. We would therefore say that T_l is preferable over T_m with respect to $\mathcal{P}$ if $\mathcal{I}_\mathcal{P}(\mathsf{T}_l, \mathsf{T}_m) > \mathcal{I}_\mathcal{P}(\mathsf{T}_m, \mathsf{T}_l)$, assuming that a higher value signifies better achievement of the property. Otherwise, the relationship is $\mathcal{I}_\mathcal{P}(\mathsf{T}_l, \mathsf{T}_m) < \mathcal{I}_\mathcal{P}(\mathsf{T}_m, \mathsf{T}_l)$.

6.1.1.1 Worst case measurements

A quality index function in the definition requires two tables as input. However, a commonly used method of evaluating a generalization is through unary quality index functions. Unary quality indices are functions applied independently on generalizations, i.e. they have a single table as input. For example, the *k*-anonymity property is a unary quality index based on the equivalence class size property $\mathcal{P}_{equiv}$, given as $\mathcal{I}_{\mathcal{P}_{equiv}}(\mathsf{T}) = \min_i(\mathcal{P}_{equiv}(\mathsf{T}))$. Unary indices only allow the measurement of an aggregate property of a generalization. This prohibits any kind of comparison of individual property values maintained by tuples in a generalization with that maintained in another. Having said so, we do not specify any restriction on the formulation of a quality index function. This is because data utility functions are typically unary in nature, i.e. they are absolute estimates of the information content of the anonymized data. We keep the

generic binary formulation since unary functions are a special case of binary functions. In other words, when using worst case privacy models such as k-anonymity and ℓ-diversity, we shall assume $\mathcal{I}_{\mathcal{P}}(\mathsf{T}_l, \mathsf{T}_m) \equiv \mathcal{I}_{\mathcal{P}}(\mathsf{T}_l)$. A similar assumption holds for information loss measurements that are not done tuple wise.

6.1.1.2 Measuring quality with spread

Privacy of an anonymized table can also be quantified in terms of the differences in individual privacy levels when compared with another anonymized table. Characterizing privacy in this manner captures the changes brought forth in individual privacy levels when moving from one node to another in the generalization lattice. This helps distinguish the privacy preserving efficiency of the two nodes even when both generate the same worst case privacy. We use the spread based quality index function in this context. The function is based on the total amount of variation (or spread) present between tuples with respect to a property, given as

$$\mathcal{I}_{\mathcal{P}}^{spr}(\mathsf{T}_l, \mathsf{T}_m) = \sum_{i=1}^{\mathcal{N}} \max(p_i^l - p_i^m, 0), \tag{6.1}$$

where $(p_1^x, \ldots, p_{\mathcal{N}}^x) = \mathcal{P}(\mathsf{T}_x)$. Thus, T_l better preserves privacy than T_m if $\mathcal{I}_{\mathcal{P}}^{spr}(\mathsf{T}_l, \mathsf{T}_m) > \mathcal{I}_{\mathcal{P}}^{spr}(\mathsf{T}_m, \mathsf{T}_l)$. This characterization follows from the intuition that a generalization better than another should be able to retain higher values of the measured property for more individuals represented in the data set. The spread function identifies the fraction of tuples with higher property values relative to another generalization and quantifies the total difference in magnitude.

The spread quality index function provides a relative characterization of privacy. The function value is only representative of the quality of a node relative to another. However, absolute estimates are more preferable since a node then does not have to be evaluated repeatedly for the same property. Unary quality functions are better in this regard since the quality of the node with respect to a property does not change irrespective of the node with which it is compared to. Hence, a unary function that can provide the same information as the binary spread function is desired. Formulating such a function is not difficult as highlighted in the following observation.

Observation: Let $S_{\mathcal{P}}(\mathsf{T}_x)$ denote the sum of the property values in the property vector $\mathcal{P}(\mathsf{T}_x)$. Then $\mathcal{I}_{\mathcal{P}}^{spr}(\mathsf{T}_l,\mathsf{T}_m) > \mathcal{I}_{\mathcal{P}}^{spr}(\mathsf{T}_m,\mathsf{T}_l)$ if and only if $S_{\mathcal{P}}(\mathsf{T}_l) > S_{\mathcal{P}}(\mathsf{T}_m)$.

The observation is also valid if the "greater than" relationship is replaced with an "equal to" or "less than" relationship. Comparing nodes under the light of the spread function can therefore be performed using the sum of the property values, i.e. $\mathcal{I}_{\mathcal{P}}^{spr}(\mathsf{T}_l,\mathsf{T}_m) \equiv \mathcal{I}_{\mathcal{P}}(\mathsf{T}_l) = S_{\mathcal{P}}(\mathsf{T}_l)$. Hence, in the subsequent sections, we shall use the notation $\mathcal{I}_{\mathcal{P}}(\mathsf{T}_l)$ to denote the quality of T_l with respect to $\mathcal{P}$, keeping in mind that $\mathcal{I}_{\mathcal{P}}$ is either a unary function (as used in worst case measurements and loss assessments) or the sum function $S_{\mathcal{P}}$ (sufficient to infer the quality according to the binary spread function).

6.1.2 Anonymizing with multiple properties

Ideally, any number of properties can be studied on a generalized table. In the presence of multiple properties, quality with respect to each property must be weighed in appropriately before choosing one generalization over another. Let us consider an anonymization with respect to the set of properties $\mathbf{P} = \{\mathcal{P}_1,\ldots,\mathcal{P}_r\}$. Assessing the quality of a generalization T_l with respect to the properties $\mathbf{P}$ will result in a vector of values $\mathbf{I_P}(\mathsf{T}_l) = [\mathcal{I}_{\mathcal{P}_1}(\mathsf{T}_l),\ldots,\mathcal{I}_{\mathcal{P}_r}(\mathsf{T}_l)]$ where the ith element represents the quality of T_l with respect to the property $\mathcal{P}_i$. The dominance relation $\succeq$ is then specified over the set of such vectors to characterize the efficiency of a generalization with respect to $\mathbf{P}$, such that $\mathbf{I_P}(\mathsf{T}_l) \succeq \mathbf{I_P}(\mathsf{T}_m)$ if

1. $\forall i = 1\ldots r : \mathcal{I}_{\mathcal{P}_i}(\mathsf{T}_l) \geq \mathcal{I}_{\mathcal{P}_i}(\mathsf{T}_m)$, and

2. $\exists j \in \{1,\ldots,r\} : \mathcal{I}_{\mathcal{P}_j}(\mathsf{T}_l) > \mathcal{I}_{\mathcal{P}_j}(\mathsf{T}_m)$.

This relation states that for a table to be better than another, it must not have worse quality across all the properties while maintaining better quality with respect to at least one property. Once again, the notion of better quality can be in terms of a lower quality index value, implying the use of a "less than" relationship for comparing across such properties. Note that the dominance relation is transitive in nature, i.e if $\mathbf{I_P}(\mathsf{T}_1) \succeq \mathbf{I_P}(\mathsf{T}_2)$ and $\mathbf{I_P}(\mathsf{T}_2) \succeq \mathbf{I_P}(\mathsf{T}_3)$, then $\mathbf{I_P}(\mathsf{T}_1) \succeq \mathbf{I_P}(\mathsf{T}_3)$. Using dominance to evaluate a generalization introduces the concept of a property based generalization (PBG).

Definition 6.3 PROPERTY BASED GENERALIZATION. Let $\mathcal{T}$ be the collection of all possible generalized versions of a table T of size $\mathcal{N}$. Given the properties $\mathbf{P} = \{\mathcal{P}_1, \ldots, \mathcal{P}_r\}$ and quality index functions $\mathbf{I} : \mathcal{I}_{\mathcal{P}_1}, \ldots, \mathcal{I}_{\mathcal{P}_r}$ (not necessarily unique), $\mathsf{T}_l \in \mathcal{T}$ is a property based generalization of $\mathsf{T}_m \in \mathcal{T}$ with respect to $\mathbf{P}$, denoted as $\mathsf{T}_l \vdash_{\mathbf{P}} \mathsf{T}_m$, if and only if $\mathbf{I_P}(\mathsf{T}_l) \succeq \mathbf{I_P}(\mathsf{T}_m)$.

Use of the dominance relationship in a PBG is not strict. It can be replaced with any other operator that defines a partial order on the set of r-dimensional vectors. We looked at a few such operators in the previous chapter. The following observations summarize the literary meaning of property based generalizations.

- T_l is *better* than T_m if and only if $\mathsf{T}_l \vdash_{\mathbf{P}} \mathsf{T}_m$. Equivalently, T_m is *worse* than T_l.

- T_l and T_m are *incomparable* (or mutually non-dominated) if and only if $\mathsf{T}_l \nvdash_{\mathbf{P}} \mathsf{T}_m$, $\mathsf{T}_m \nvdash_{\mathbf{P}} \mathsf{T}_l$ and $\mathsf{T}_l \neq \mathsf{T}_m$.

Incomparable generalizations typically signify trade-offs across certain properties. The non-dominance relationship results when a generalization has better quality with respect to one property but is worse with respect to another. Therefore, it is our objective to identify such generalizations for reporting. In addition, the chosen generalizations must also be minimal.

Definition 6.4 MINIMAL PROPERTY BASED GENERALIZATION. Given a collection $\mathcal{T}$ of generalized versions of a table T and the properties $\mathbf{P} = \{\mathcal{P}_1, \ldots, \mathcal{P}_r\}$, $\mathsf{T}_w \in \mathcal{T}$ is a minimal property based generalization of $\mathcal{T}$ if $\nexists \mathsf{T}_m \in \mathcal{T} : \mathsf{T}_m \vdash_{\mathbf{P}} \mathsf{T}_w$.

Minimal property based generalizations are analogous to Pareto-optimal solutions in a multi-objective optimization problem. For an intuitive understanding, let us consider the k–anonymity model as a measure of privacy and an information loss function Π. The properties under consideration in this case are $\mathcal{P}_1$: *equivalence class size of a tuple* and $\mathcal{P}_2$: *total information loss*, as given by Π. The quality index functions are $\mathcal{I}_{\mathcal{P}_1}(\mathsf{T}) = \min_i \mathcal{P}_1(\mathsf{T})$ and $\mathcal{I}_{\mathcal{P}_2}(\mathsf{T}) = \Pi(\mathsf{T})$. A generalization T is then a minimal PBG if there is no other generalization T' such that

- $\mathcal{I}_{\mathcal{P}_1}(\mathsf{T}') \geq \mathcal{I}_{\mathcal{P}_1}(\mathsf{T})$ and $\mathcal{I}_{\mathcal{P}_2}(\mathsf{T}') < \mathcal{I}_{\mathcal{P}_2}(\mathsf{T})$, or

Figure 6.1: Schematic of an iterative search process.

- $\mathcal{I}_{\mathcal{P}_1}(\mathsf{T}') > \mathcal{I}_{\mathcal{P}_1}(\mathsf{T})$ and $\mathcal{I}_{\mathcal{P}_2}(\mathsf{T}') \leq \mathcal{I}_{\mathcal{P}_2}(\mathsf{T})$.

In other words, a minimal PBG attains a higher value of k (or a lower information loss), as compared to any other generalization, without increasing the information loss (or a decrease in the value of k). The set of all minimal PBGs is therefore the collection of minimally distorted generalizations for different values of k.

6.2 Representative PBGs

One drawback of using the dominance relation $\succeq$ is the inability to control the number of minimal PBGs to report during the search process. We assume here a search process with a finite memory to store minimal PBGs. Fig. 6.1 illustrates a typical algorithm in this context. The search process iteratively tries to converge to the set of minimal PBGs. The *generator* component is responsible for creating a new candidate generalization, preferably using the current set of generalizations in the archive. The *updator* component performs a comparison of the candidate generalization with those maintained in the archive and removes all generalizations which cannot be minimal PBGs. The purpose behind maintaining such an archive is to guide the search process towards better regions of the search space, and at the same time maintain a list of the best solutions found so far.

The issue to address is the size of the archive. With no limitation on the size, it may become impossible to store additional prospective generalizations owing to restrictions on physical memory. This is likely to happen faster when the number of properties under consideration increases, in which case satisfying a dominance relationship becomes more

and more difficult (a consequence of the curse of dimensionality). A generator using the current archive in creating new candidate generalizations will also use more computation time if the archive size is allowed to grow unbounded. Further, it is sufficient that a data disseminator obtains a finite representative subset of minimal PBGs for the purpose of decision making. The primary criteria to fulfill is that the archive maintain generalizations that are minimal PBGs and at the same time have enough diversity to represent the trade-off behavior across the multiple properties.

6.2.1 Box-dominance

Let $\mathcal{M}$ denote the set of all minimal PBGs corresponding to a given data set. The objective is to obtain a polynomially bounded sized subset of $\mathcal{M}$. Let $(\epsilon_1, \ldots, \epsilon_r); \epsilon_i > 0$ denote a *discretization vector*, r being the number of properties considered. The quality index space $\mathbb{R}^r$ is then discretized by placing a hypergrid with the co-ordinates $0, \epsilon_i, 2\epsilon_i, \ldots$ along each of the r dimensions. This divides the space into boxes with side lengths same as the discretization vector. Assuming the quality index functions are bounded on both sides, i.e. $0 < \mathcal{I}_{\mathcal{P}_i}(\mathsf{T}) \leq K_i$, the box of a generalization T_l is given by the vector

$$\mathcal{B}(\mathsf{T}_l) = \left[\left\lfloor \frac{\mathcal{I}_{\mathcal{P}_1}(\mathsf{T}_l)}{\epsilon_1} \right\rfloor, \ldots, \left\lfloor \frac{\mathcal{I}_{\mathcal{P}_r}(\mathsf{T}_l)}{\epsilon_r} \right\rfloor \right]. \tag{6.2}$$

A modified dominance relation, called *box-dominance* and denoted by $\succeq_{box}$, is then formulated as

$$\mathbf{I_P}(\mathsf{T}_l) \succeq_{box} \mathbf{I_P}(\mathsf{T}_m) \iff \begin{cases} \mathcal{B}(\mathsf{T}_l) \succeq \mathcal{B}(\mathsf{T}_m) & \text{,if } \mathcal{B}(\mathsf{T}_l) \neq \mathcal{B}(\mathsf{T}_m) \\ \mathbf{I_P}(\mathsf{T}_l) \succeq \mathbf{I_P}(\mathsf{T}_m) & \text{,otherwise} \end{cases}. \tag{6.3}$$

The box-dominance relation first places the quality index value vectors ($\mathbf{I_P}(\mathsf{T}_l)$ and $\mathbf{I_P}(\mathsf{T}_m)$) in their boxes. If the vectors are on different boxes, then T_l cannot be a PBG of T_m if the box of T_l does not dominate the box of T_m. Otherwise, for the case when boxes are the same, the dominance is checked on the quality index values. Further, every box is allowed to hold only one generalization. Choice between two incomparable generalizations belonging to the same box is made arbitrarily. Fig. 6.2 illustrates this modified dominance relation.

Non-dominated boxes signify regions where a minimal PBG exists. By allowing the existence of a single generalization per non-dominated box, the modified dominance rela-

Figure 6.2: Non-dominated boxes and representative minimal PBGs for a hypothetical two property anonymization.

tionship maintains a representative subset of the minimal PBGs. Intuitively, every generalization in a box differs by less than ϵ_i, in terms of its quality with respect to every property $\mathcal{P}_i$, when compared to every other generalization in the same box. Box-dominance does not consider such differences to be substantial enough. The discretization vector determines the size of the boxes and hence impacts the size of the representative subset. If quality index values are in the integer domain, then using a discretization vector of all ones implies using the un-modified dominance relation.

6.2.2 An updator using $\succeq_{box}$

The use of the box-dominance relation in an updator guarantees two properties – (i) the reported generalizations constitute a diverse set of PBGs across the quality index space, and (ii) the reported generalizations are minimal PBGs of all generalizations generated so far. Procedure 6.1 outlines an updator algorithm using box-dominance.

The algorithm starts with an empty archive $\mathcal{A}$. The first candidate generalization from the generator is therefore automatically inserted into the archive. For subsequent candidates, use of $\succeq_{box}$ effectuates a two level dominance check as explained earlier. First, all generalizations for which the candidate T is a PBG are removed from the archive (Steps 5 and 6). Next, two sets are computed – (i) $\mathcal{S}_{dominated}$ as the set of all generalizations which are PBGs of T, and (ii) $\mathcal{S}_{box}$ as the set of all generalizations whose boxes are same as that of T. The candidate T should not be inserted into the archive if the set $\mathcal{S}_{dominated}$ is non-

Procedure 6.1 BoxDominanceUpdator(Archive $\mathcal{A}$, Generalization T)

Input: Archive $\mathcal{A}$, candidate generalization T.
Output: Updated archive $\mathcal{A}$.
1: **if** $(\mathcal{A} = \phi)$ **then**
2:     $\mathcal{A} = \{T\}$
3:     **goto** Line 12
4: **end if**
5: $\mathcal{S}_{dominate} = \{T' \in \mathcal{A} | \mathbf{I_P}(T) \succeq_{box} \mathbf{I_P}(T')\}$
6: $\mathcal{A} = \mathcal{A} - \mathcal{S}_{dominate}$
7: $\mathcal{S}_{dominated} = \{T' \in \mathcal{A} | \mathbf{I_P}(T') \succeq_{box} \mathbf{I_P}(T)\}$
8: $\mathcal{S}_{box} = \{T' \in \mathcal{A} | \mathcal{B}(T) = \mathcal{B}(T')\}$
9: **if** $(\mathcal{S}_{dominated} = \phi$ **and** $\mathcal{S}_{box} = \phi)$ **then**
10:     $\mathcal{A} = \mathcal{A} \cup \{T\}$
11: **end if**
12: **return** $\mathcal{A}$

empty, i.e. there exists a generalization in the archive which is a PBG of T. Further, if $\mathcal{S}_{box}$ is not empty then inclusion of T in the archive will result in the presence of two different generalizations which are positioned in the same box. Step 9 checks for these two conditions, thereby guaranteeing that only non-dominated boxes contain a solution and only one solution is contained in a non-dominated box. The following two theorems prove that the archive maintained in this manner will only contain minimal PBGs from the set of candidates generated so far and will be of bounded size.

Theorem 6.1 Let $\mathcal{M}_g$ denote the set of all generalizations produced by a generator until iteration t and $\mathcal{M}_g^*$ denote the set of minimal PBGs of $\mathcal{M}_g$. Then the archive $\mathcal{A}$ as maintained by Procedure 6.1 contains only minimal PBGs of $\mathcal{M}_g$, i.e. $\mathcal{A} \subseteq \mathcal{M}_g^*$.

Proof We assume that Procedure 6.1 is incorrect, implying $\mathcal{A} \not\subseteq \mathcal{M}_g^*$. Therefore there exists $T_s \in \mathcal{A}$ generated at iteration s such that $T_s \notin \mathcal{M}_g^*$.

If $T_s \notin \mathcal{M}_g^*$ then there exists $T_q \in \mathcal{M}_g$ discovered at iteration $q \neq s$ such that $\mathbf{I_P}(T_q) \succeq \mathbf{I_P}(T_s)$. Also, either $\mathcal{B}(T_q) = \mathcal{B}(T_s)$ or $\mathcal{B}(T_q) \succeq \mathcal{B}(T_s)$. We can merge these cases and say $\mathbf{I_P}(T_q) \succeq_{box} \mathbf{I_P}(T_s)$.

Case (i): $q < s$

If T_q is present in $\mathcal{A}$ at iteration s then T_s will not be included in the archive since $\mathcal{S}_{dominated}$ for T_s contains at least T_q.

If T_q is not present in $\mathcal{A}$ at iteration s then it must have been removed by a generalization T_r in $\mathcal{A}$ such that $\mathbf{I_P}(\mathsf{T}_r) \succeq_{box} \mathbf{I_P}(\mathsf{T}_q)$. We therefore have $\mathbf{I_P}(\mathsf{T}_r) \succeq \mathbf{I_P}(\mathsf{T}_q)$ or $\mathcal{B}(\mathsf{T}_r) \succeq \mathcal{B}(\mathsf{T}_q)$. Using the transitivity of the $\succeq$ relation, we have $\mathbf{I_P}(\mathsf{T}_r) \succeq \mathbf{I_P}(\mathsf{T}_s)$ or $\mathcal{B}(\mathsf{T}_r) \succeq \mathcal{B}(\mathsf{T}_s)$, which implies $\mathbf{I_P}(\mathsf{T}_r) \succeq_{box} \mathbf{I_P}(\mathsf{T}_s)$. Hence in this case as well $\mathcal{S}_{dominated} \neq \phi$ for T_s. Note that T_r itself might have got removed from the archive between iteration q and iteration s. However, owing to the transitivity, the generalization which removes it will instead appear in $\mathcal{S}_{dominated}$ for T_s. Hence T_s will never appear in $\mathcal{A}$, i.e. $\mathsf{T}_s \notin \mathcal{A}$, which is a contradiction.

Case (ii): $q > s$

In this case, if T_s exists in $\mathcal{A}$ at iteration q then it would be removed from the archive as it belongs to the set $\mathcal{S}_{dominate}$ of T_q. Further, if T_q gets removed and T_s gets re-generated at a later iteration, the transitivity property would assure that T_s does not get re-inserted into the archive. Thus, $\mathsf{T}_s \notin \mathcal{A}$ which is again a contradiction.

Therefore, T_s can never be a member of the archive at iteration t if it is not a minimal PBG. We can therefore say Procedure 6.1 is correct and the archive $\mathcal{A}$ contains only minimal PBGs of $\mathcal{M}_g$. $\square$

Theorem 6.2 The archive $\mathcal{A}$ as maintained by Procedure 6.1 is of bounded size, given as

$$|\mathcal{A}| \leq \prod_{i=1}^{r-1} b_i, \tag{6.4}$$

where b_i is the ith largest element of the vector $(\frac{K_1}{\epsilon_1}, \ldots, \frac{K_r}{\epsilon_r})$.

Proof Recall that $K_1, \ldots, K_r$ are the upper bounds of the quality index functions for r properties. These values can very well be equal. By using box co-ordinates at $0, \epsilon_i, 2\epsilon_i, \ldots$ along each dimension i, we have divided the quality index value space into $\prod_{i=1}^{r} \frac{K_i}{\epsilon_i}$ boxes and only one node in each box can be included in $\mathcal{A}$. We now cluster these boxes into groups of b_r boxes, giving us a total of $\prod_{i=1}^{r-1} b_i$ clusters. A cluster is formed by grouping together boxes that have the same co-ordinates in all but one dimension. Note that choosing b_r as the parameter to decide the number of boxes in a cluster gives us the smallest possible cluster size and hence the largest number of clusters. This is required if an upper bound on the archive size is to be computed. Next, in a cluster, the box having the maximum

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Procedure 6.2 PBG-EA()

Output: Archive $\mathcal{A}$ of representative minimal PBGs.

1: $\mathcal{A} = \phi; t = 0$
2: Initialize population P_t
3: Evaluate P_t
4: Update $\mathcal{A}$ with nodes in P_t
5: **repeat**
6:     Assign fitness to nodes in P_t and $\mathcal{A}$
7:     Perform selection in $P_t \cup \mathcal{A}$
8:     Generate P_{t+1} by performing recombination on selected nodes
9:     Update $\mathcal{A}$ with nodes in P_{t+1}
10:    $t = t + 1$
11: **until** ($t \geq$ maximum number of iterations allowed)
12: **return** $\mathcal{A}$

---

co-ordinate value in the differing dimension will dominate all other boxes in the cluster. Therefore, only such a box will contain a minimal PBG. Each cluster can therefore contribute only one minimal PBG, bounding the archive size to the number of such clusters, i.e. $|\mathcal{A}| \leq \prod_{i=1}^{r-1} b_i$. $\square$

Note that the specific upper bounds on the quality index functions are not used by the updator but is only required to show that the size of the archive depends on these bounds and the chosen discretization vector. Given a specific size for the representative subset of minimal PBGs, the discretization vector can be adjusted accordingly if the upper bounds on the quality index functions are known.

6.3 An Evolutionary Generator

An efficient generator is required not only to find new candidate PBGs, but also to minimize the number of node evaluations performed during the search process. The generator evaluates each node that it explores and provides it to the updator. Discovering nodes that can converge quickly (without involving too many intermediate node evaluations) to a minimal PBG is therefore the primary objective of the generator. We propose here an evolutionary algorithm for this purpose, henceforth called *PBG-EA*. The algorithm follows the structure described in Procedure 6.2.

The update method from Procedure 6.1 is used iteratively in steps 4 and 9. Specifics of the other steps are described next.

6.3.1 Population initialization and evaluation

A population P_t is a collection of N_{pop} nodes in the lattice and undergoes changes as the algorithm progresses. Recall that every node is a vector of s dimensions where s is the number of quasi-identifiers. The population P_0 is created by randomly selecting nodes in the lattice. The fully generalized and fully specialized nodes are always inserted into this initial population as they are trivially minimal PBGs. Evaluation of a population means computing the quality index values for each node in the population. Outliers in the data set must be handled in this step and is done using a maximum allowed suppression scheme.

6.3.2 Fitness assignment

Fitness signifies the potential of a node to be a minimal PBG relative to the current population and archive. The fitness assignment we use is adapted from the one used in the SPEA2 algorithm [195]. Let dom_P be the number of nodes in $P_t \cup \mathcal{A}$ dominated by $P \in P_t \cup \mathcal{A}$. The fitness of a node P is then computed as the sum of the dominance counts of the nodes which dominate P, or

$$Fitness_P = \sum_{P' \in P_t \cup \mathcal{A} \text{ and } P' \succeq P} dom_{P'}. \tag{6.5}$$

All non-dominated generalizations will therefore have a fitness of zero. Hence, lower fitness implies better generalizations. The fitness of a node not only reflects the number of nodes that dominate it but also takes into account the dominance power of those nodes.

6.3.3 Selection

Selection of nodes for recombination is performed by using a binary tournament strategy in $P_t \cup \mathcal{A}$. Under this strategy, two nodes are randomly chosen from $P_t \cup \mathcal{A}$ and the one with the lower fitness is selected. The process is repeated for N_{pop} times, giving a selected population of size N_{pop}.

6.3.4 Recombination

Recombination is a two step process involving the crossover and mutation operators, the resulting nodes from which are used as the next population P_{t+1}. A single

Figure 6.3: Single point crossover and single-step mutation in the recombination process.

point crossover is started by first choosing two nodes (without replacement) from the selected population. Parts of the vectors representing the two nodes are then swapped at a randomly chosen crossover point. The swapping procedure is performed with a probability of p_{cross}; otherwise chosen nodes move unchanged into the next population. Each crossover operation results in two nodes for the next population. Performing the operation on the entire selected population creates N_{pop} nodes for inclusion in P_{t+1}. An intermediate single-step mutation is performed on these nodes — with a probability p_{mut}, each attribute's generalization level is either increased or decreased by one using appropriate rounding so that generalization levels are between zero and the DGH lengths. Fig. 6.3 illustrates the recombination procedure. Note that similar recombination procedures can be designed when the representation of a generalization is finer grained than the one used here [16, 92, 110].

6.4   Performance Analysis

We applied our methodology to the "adult.data" benchmark data set. The attributes used in this study along with their DGH lengths are listed in Table 4.3. In addition, we consider another attribute "occupation" having 14 distinct values as the sensitive attribute wherever required. The total number of nodes in the lattice is 17920. Using the maximum allowed suppression scheme (Section 2.4.3.2), the suppression limit η is set at 1% of the data set size.

Experiments are performed using worst case as well as spread based privacy measure-

ments. k-anonymity and ℓ-diversity are used as the privacy objectives for experiments using worst case privacy. For experiments with spread based measurements, we consider the two properties $\mathcal{P}_1$: *size of equivalence class of a tuple* and $\mathcal{P}_2$: *count of sensitive attribute value of a tuple in its equivalence class.* Sum of the property values in the respective property vectors are denoted by S_k and S_ℓ respectively in the plots.

Information loss estimates are obtained using the general loss metric (GLM) and classification error (CM) [92]. GLM computes a normalized information loss using the sizes of the generalized domains to which values of an attribute belong. The attribute "salary class" is used as the class label while performing experiments with the CM metric. The metric assigns zero error to a tuple if its class label is same as the majority class label; otherwise one. The lattice size in this case is 8960. Solutions reported by PBG-EA are compared with those obtained by an exhaustive search of the entire generalization lattice. Note that the number of nodes evaluated in the exhaustive search is equal to the size of the lattice, while that used by PBG-EA is much less. Further, an exhaustive search may not always be computationally feasible. Nonetheless, an exhaustive search in this study provides us a definitive platform to judge the efficiency of PBG-EA in finding true minimal PBGs.

An instance of PBG-EA is run with a population size $N_{pop} = 25$ and for 100 iterations, $p_{cross} = 0.8$ and $p_{mut} = 1/$number of quasi-identifiers $= 0.125$. Each experiment is run 20 times to compute the mean and variance of the performance metrics (discussed below). The discretization vector is set to all ones, unless otherwise indicated.

6.4.1 Performance metrics

Efficiency of PBG-EA is measured in terms of its ability to converge to the true minimal PBGs (as found by the exhaustive search) and how well the solutions represent the set of all minimal PBGs. Two metrics are used to quantify these two aspects.

6.4.1.1 Convergence Error (CE)

Let $\mathcal{M}$ be the set of all minimal PBGs for a data set and $\mathcal{M}'$ be the solutions in the archive at the end of the final iteration. Quality index values of all nodes in $\mathcal{M}$ and

Table 6.1: CE and RR values in PBG-EA anonymization with different sets of properties. Values are shown as $\frac{\text{mean}}{\text{variance}}$ from the 20 runs.

Objectives	*CE*	*RR*
k, GLM	$\frac{3.7\times10^{-4}}{6.5\times10^{-9}}$	$\frac{0.94}{8\times10^{-4}}$
k, ℓ, GLM	$\frac{3.3\times10^{-4}}{1.1\times10^{-7}}$	$\frac{0.93}{1.1\times10^{-3}}$
S_k, GLM	$\frac{5.7\times10^{-4}}{2.5\times10^{-7}}$	$\frac{0.84}{1.7\times10^{-3}}$
S_k, S_ℓ, GLM	$\frac{6.6\times10^{-4}}{2.0\times10^{-7}}$	$\frac{0.83}{1.4\times10^{-3}}$

$\mathcal{M}'$ are normalized by dividing the values by the corresponding maximum in $\mathcal{M}$. The convergence error (CE) is then given as

$$CE = \sum_{M' \in \mathcal{M}'} \min_{M \in \mathcal{M}} \left[dist\left(\mathbf{I_P}(M), \mathbf{I_P}(M') \right) \right], \tag{6.6}$$

where *dist* is the euclidean distance between two vectors. A CE value of zero means all solutions in the archive have converged to some minimal PBG.

6.4.1.2   Representation Ratio (RR)

The representation ratio is the fraction of non-dominated boxes in $\mathcal{M}$ that are occupied by a solution in $\mathcal{M}'$. Given a discretization vector, solutions in $\mathcal{M}$ are assigned their respective boxes and the non-dominated boxes are marked. RR signifies how many of these marked boxes are occupied by a solution in the archive. A value of one signifies that a solution in each non-dominated box exists in the archive.

6.4.2   Worst case privacy

Fig. 6.4 compares the PBG-EA solutions with those obtained from an exhaustive search for worst case privacy measurements as in k-anonymity and ℓ-diversity. Trade-offs between the k value and the loss are evident from the two property (k-GLM) solutions. The convergence efficiency of PBG-EA is worth mentioning as 94% of all minimal PBGs are discovered by the algorithm (Table 6.1). Although the algorithm utilizes random numbers in a number of places, this performance of the algorithm is more or less persistent (low variance across the 20 runs). The trade-offs in the three property (k-ℓ-GLM) seem to be more in terms of loss, rather than between k and ℓ. Nonetheless, a few points do exist that demonstrate that alternative ℓ values can be obtained for a fixed value of k at the

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Figure 6.4: PBG-EA solutions for the two property (k-GLM) and three property (k-ℓ-GLM) problems.

Figure 6.5: PBG-EA solutions for the two property (S_k-GLM) and three property (S_k-S_ℓ-GLM) problems using spread based quality.

expense of more information loss. The convergence of the solutions are equally good in this case as well.

6.4.3 Measuring privacy with spread function

Fig. 6.5 shows the PBG-EA solutions when the spread function is used to measure privacy. The RR is slightly lower in this case. Nonetheless, the convergence error is still low. Using the spread function induces a higher number of minimal PBGs whose discovery typically requires more number of iterations. Given that the same population size and number of iterations are used here as for the worst case privacy, PBG-EA demonstrates quick convergence to a number of the minimal PBGs. We observe that increasing

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Figure 6.6: PBG-EA solutions with a single privacy property (*k*-anonimity) and two loss metrics (GLM-CM).

the number of properties from two to three has very little influence on the RR. A good fitness assignment strategy is required for early determination of solution efficiency. The dominance count based fitness assignment is ideally suited here since the efficiency of the dominating solutions is also taken into account while assessing a solution. Typically, a solution is not worth exploring if it is dominated by a large fraction of the nodes in the lattice. The fitness scheme takes this a step further to also consider the quality of the solutions that dominate it.

An imperative question to ask at this point is the selection of a final solution from the solution set. The choice is based on acceptable levels of k (or ℓ) for the worst case privacy measurements. Similarly, the solution set in this case can be filtered so that only those solutions satisfying a minimum requirement are further analyzed. The solutions left thereafter show that further decisions on the account of privacy can be made even after a minimum level constraint is met. The difference of individual privacy levels as considered in the spread function forms the basis for these decisions.

6.4.4 Using multiple loss metrics

PBGs can also be used to find generalizations that are acceptable in terms of more than one loss metric. Finding generalizations to serve multiple purposes has not been addressed in the research community. Fig. 6.6 shows the minimal PBGs obtained when the *k*-anonimity property is evaluated against two different loss metrics, namely GLM

Table 6.2: CE and RR values in PBG-EA anonymization with different discretization vectors.

ϵ_k	ϵ_ℓ	ϵ_{GLM}	*CE*	*RR*
5	-	100	$\frac{4.3\times10^{-4}}{2.6\times10^{-7}}$	$\frac{0.95}{1.6\times10^{-3}}$
10	-	1000	$\frac{1.6\times10^{-4}}{8.0\times10^{-8}}$	$\frac{0.98}{8.2\times10^{-4}}$
50	-	10000	$\frac{1.7\times10^{-4}}{1.1\times10^{-8}}$	$\frac{1.0}{0.0}$
5	2	100	$\frac{4.9\times10^{-3}}{2.5\times10^{-7}}$	$\frac{0.92}{1.2\times10^{-3}}$
10	4	1000	$\frac{7.4\times10^{-3}}{9.3\times10^{-7}}$	$\frac{0.92}{7.0\times10^{-4}}$
50	6	10000	$\frac{1.8\times10^{-2}}{8.2\times10^{-6}}$	$\frac{0.88}{1.7\times10^{-3}}$

and CM. The heavily scattered points across the entire space signify the existence of reciprocal relationships between GLM and CM. Ideally, for a given k value, the convention is to choose the generalization with the lowest GLM. However, the multi-loss metric experiment indicates that choosing a generalization with a comparatively higher GLM can serve the dual purpose of making the anonymized data set also suitable for classification tasks.

6.4.5 Efficiency in finding representative minimal PBGs

Although the set of minimal PBGs for the data set used in the study is not unbounded in size, we experimented with several discretization vectors to demonstrate the efficiency of PBG-EA in finding a representative subset. Table 6.2 shows the performance measures for some vectors. The high representation ratio is indicative of the fact that PBG-EA solutions cover most of the non-dominated boxes generated by the use of the discretization vectors. For two property (k-GLM) anonymization, as higher ϵ values are used for the objectives, the efficiency of PBG-EA improves in terms of RR. However, using more number of properties tend to slightly affect the performance owing mostly to the limited number of iterations.

Fig. 6.7 shows a part of the boxed quality index space for $\epsilon_k = 5$ and $\epsilon_{GLM} = 5000$ in the range of $k = 1$ to 100. The archive in this case contains only one solution from each non-dominated box. The representative solution set is characterized by the property that, for any other solution, there is a solution in the archive whose k or GLM is within a difference less than the corresponding discretized level. Choosing larger discretizations

Figure 6.7: Partial ($k \in [1,100]$) representative set of solutions for the two property (k-GLM) problem with $\epsilon_k = 5$ and $\epsilon_{GLM} = 5000$.

therefore mean a smaller representative subset. Choice of the vector to use is at the discretion of the data publisher, determined by the number of solutions to analyze in the first place.

6.4.6 Node evaluation efficiency

Efficiency of PBG-EA in converging quickly to a minimal PBG is evaluated by counting the number of unique nodes that are evaluated by it during the search process. Although the evolutionary algorithm can potentially explore 2500 (25×100) distinct nodes in the lattice, a much smaller number is actually evaluated. Table 6.3 lists the average (out of the 20 runs) number of unique node evaluations performed for different problem instances. We consider this low percentage of node evaluations to be a positive indication of the convergence efficiency of PBG-EA. This is particularly promising since the entire set of minimal PBGs (all one discretization vector) is found by exploring a small 5% of nodes in the lattice. The nodes evaluated is slightly higher for three property problems. This is not surprising since the number of minimal PBGs is also comparatively higher in such cases.

6.5 Conclusions

In this chapter, we have argued that the use of existing notions of a minimal generalization burdens a data publisher with the task of model parameter selection, quite often without knowing its impact on the utility of the data. The task becomes further

Table 6.3: Average number of nodes evaluated in PBG-EA for different sets of properties. Total number of nodes in the first four sets is 17920 and that in the last set is 8960.

Objectives	Avg. node evaluations
k, GLM	916 (5.1%)
k,ℓ, GLM	946 (5.3%)
S_k, GLM	1136 (6.3%)
S_k,S_ℓ, GLM	1197 (6.7%)
k, GLM, CM	1073 (11.9%)

complicated when the data publisher has to include multiple privacy models or account for more than one possible data usage scenario.

In our approach, we propose identifying the basic properties that provide the requisite protection from a privacy breach, and then measuring them for each underlying individual. This generates a property vector for every generalization. Comparison of generalizations with respect to a single property is performed using quality index functions. Apart from the conventional worst case measures of privacy, a binary index function is presented which measures privacy using the variations in individual privacy levels. In the presence of multiple properties, a dominance based partial order relationship is used to define a property based generalization. Optimality in such generalizations is signified by non-dominated generalizations (minimal PBGs) under the dominance relation. The objective of our approach is to gather a representative subset of generalizations that demonstrate trade-offs in quality while trying to attain the different properties. The representative subset is maintained by using a box-dominance operator which guarantees that the solutions generated are optimal and the number of such solutions do not grow unbounded in size. An evolutionary algorithm is suggested to explore the generalization lattice and identify the optimal nodes. Application on a benchmark data set shows that the algorithm can quickly discover a diverse set of minimal PBGs with a small number of node evaluations.

Observations from our multi-loss experiment suggest that the problem of microdata anonymization to serve multiple usages needs to be explored in more details. No study has yet tried to understand the correlation between the multitude of loss metrics that exist in the literature.

CHAPTER 7

Disclosures in a Continuous Location-Based Service

Technological advances in location tracking and its growing embedment in mobile devices have opened up a new spectrum of on-demand services. These services deliver customized information based on the location of a mobile object. A location-based service (LBS) may simply provide information on the nearest gas station, perform targeted marketing by location-based advertising, or enhance emergency response services, among others. A mobile object can receive up-to-date and precise information by revealing its location accurately to the LBS provider. The services are classified into two types based on how frequently a mobile object communicates its location to the provider. The first one is a *snapshot* LBS where the current location of the mobile object is sufficient to deliver the service. For example, an emergency response service such as E-911 in North America acquires the location of the caller to dispatch emergency services at the precise location. The second one is a *continuous* LBS where the mobile object must periodically communicate its location as part of the service agreement. For example, a Pay-As-You-Drive insurance service must receive location updates from the mobile object to bill the consumer accurately. A real time friend finder service such as Loopt® is another example in this category.

Application domains are potentially endless with location-tracking technology. However, a serious concern surrounding their acceptance is the potential usage of the location

data to infer sensitive personal information about the mobile users. Privacy in location-based services has been studied from two different perspectives – *location anonymity* and *query privacy*. Location anonymity is related to the disclosure of exact locations that a user has visited. This knowledge can in turn reveal personal lifestyles, places of frequent visits, or even the medical problems of the involved user. With access to exact location data, sender anonymity can be violated without the capability to track a mobile user. We refer to this class of adversaries as *location-unaware adversaries*. Such adversaries use external information to perform attacks resulting in restricted space identification, observation identification and location tracking [77].

Query privacy is related to the disclosure of sensitive information in the query itself and its association to a user. Consider a marketing agency such as CellFire® that delivers mobile coupons to users based on their location and category of interest. The retail partners sponsoring such an agency are spread out across multiple categories, ranging from apparels, groceries, automotives, entertainment, electronics to insurance, telecommunication, marketing and fitness. Each category in itself can be sub-divided; apparels, for example, can be divided into men's, women's or children's. Users therefore use *service attribute* identifiers that specify their interest category. More often than not, a user's service attribute value is considered sensitive since it directly reveals personal preferences (or requirements) of the user. Addressing the sensitivity is more important in a service like Google^{TM} Adwords that can perform location-based marketing on virtually any area of interest. Query privacy is therefore an essential requirement. It has a more direct impact on user privacy than location anonymity.

7.1 Protecting Privacy in a Continuous LBS

Location obfuscation is one of the widely researched approaches to safeguard location anonymity. This technique guarantees that the location data received at the LBS provider can be associated back to more than one object – to at least k objects under the *location k-anonymity* model [77]. For this, a *cloaking region* is communicated to the service provider instead of the actual location. A k-anonymous cloaking region contains at least $k-1$ other mobile objects besides the service user. However, this approach is not sufficient to pre-

serve privacy in a continuous LBS. In the continuous case, an object maintains an ongoing session with the LBS, and successive cloaking regions may be correlated to associate the session back to the object. Such *session associations* reveal the trajectory of the involved object, and any sensitive information thereof. Cloaking regions may be susceptible to correlation attacks due to their close proximity in time or movement restriction of objects in those regions [188]. Assuring that every cloaking region contains k objects is not sufficient since the absence of an object in one of the regions eliminates the possibility that it is the session owner. Performing such elimination is much easier for a *location-aware adversary* who has the capability to monitor users. This class of adversaries has exact location information on one or more objects and uses it to eliminate possibilities and probabilistically associate the session to consistently existing objects. It may seem that these attacks can be avoided by using a different identifier for every cloaking region. However, location data can still be correlated using techniques such as multi-target tracking [147]. Besides, the provider needs to be able to distinguish updates from the same object in order to maintain service quality [18].

Session association attacks can be avoided if it can be assured that every cloaking region in a session contains k common objects. This is referred to as *historical k-anonymity* [20]. However, as a result of the movement of objects, a historically k-anonymous cloaking region is very likely to grow in size over time, thereby deteriorating service quality. Without the proper strategies to control the size of the cloaking region, historical k-anonymity is only a theoretical extension of k-anonymity for continuous LBS.

Preservation of query privacy in a LBS is similar to protection against attribute disclosures in data privacy. A typical principle used in this context is *query ℓ-diversity* [114]. A cloaking region conforming to query ℓ-diversity contains users with at least ℓ "well-represented" service attribute values. One way of enforcing the principle is to ascertain that there are users with at least ℓ distinct interest categories. Henceforth, any reference to query ℓ-diversity implies this particular enforcement. Query ℓ-diversity ensures that a user cannot be linked to less than ℓ distinct service attribute values, thereby preventing homogeneity attacks [120]. However, this approach is not sufficient to prevent query disclosures in a continuous location-based service.

Table 7.1: Successive query 3-diverse cloaking regions.

time	user set	service attribute values
t_1	$\{U_1, U_2, U_3\}$	$\{a, b, c\}$
t_2	$\{U_1, U_2, U_4\}$	$\{a, b, d\}$
t_3	$\{U_1, U_3, U_4\}$	$\{a, c, d\}$

Users issue recurrent queries in a continuous LBS over a period of time. Each query is accompanied by current location information in order to obtain updated results. An example is a mobile user cruising through an urban locality and repeatedly using a marketing service to find the real estates on sale in the neighborhood. In an attempt to maintain privacy, the continuous LBS in this case receives a sequence of cloaking regions corresponding to the recurrent queries. Let us assume that the set of users inside the cloaking regions and their service attribute values are as shown in Table 7.1. All cloaking regions generated for the involved user are query 3-diverse and location 3-anonymous. However, given the information that the three cloaking regions are generated for the same user repeatedly inquiring about a particular category of interest, it is evident that the attribute value of interest is 'a'. Further, U_1 being the only user common in all the cloaking regions, an adversary infers that U_1 has an interest in the category 'a'. This form of disclosure occurs because existing models providing query privacy do not consider the possibility of correlating consecutive sets of service attribute values generated during the recurrent use of a LBS. New techniques are therefore required to ensure that users of continuous location-based services are well protected from threats originating from query disclosures.

7.2   Related Work

While significant research has gone into algorithms that enforce location anonymity [13, 73, 77, 98], very few of them address the problem in the context of a continuous LBS. Gruteser and Liu specifically investigate privacy issues in continuous LBS [78]. They introduce the location inference problem where an adversary can infer supposedly hidden locations from prior or future location updates. They argue that privacy in continuous LBS applications can be situation dependent, hence pressing the requirement for sensitive

and insensitive areas. Their base algorithm releases location updates only if an object is in an insensitive area. Other variants try to preserve privacy by controlling the frequency of updates. Hoh and Gruteser propose a perturbation algorithm to cross paths of objects (by exchanging their pseudonyms) when they are close to each other [83]. However, the approach does not safeguard against location-unaware adversaries as actual location data is communicated to the LBS. Kido et al. use false dummies to simulate the movement of mobile nodes in order to hide the trace of an actual object [99]. An object submits $k-1$ false locations along with the true one during an update. Such a technique fails to guarantee privacy in the presence of location-aware adversaries. Xu and Cai propose using historical traces of objects to derive a spatio-temporal cloaking region that provides trajectory protection [189]. However, the approach requires that an object specifies its entire path before using the service.

Bettini et al. introduce historical k-anonymity and propose a spatio-temporal generalization algorithm to enforce it [20]. The generalization algorithm enlarges the area and time interval of the request to increase the uncertainty about the real location, and at the same time include k common objects. The generalization is accepted only if the area and time interval satisfy specified tolerance constraints. The method fails to account for mobility of the objects, without which the generalized area can easily become larger than the tolerance if it were to satisfy historical k-anonymity. Uninterrupted sessions will therefore become very difficult.

Chow and Mokbel argue that spatial cloaking algorithms should satisfy the *k-sharing* and *memorization* properties to be robust against session associations [39]. They focus on a cloaking mechanism that allows users to specify different privacy levels for location anonymity and query privacy. The two properties together imply historical k-anonymity. Their algorithm maintains groups of objects based on the two properties, along with query types involved with the objects. Unlike previous approaches, the method does not have any tolerance to satisfy in terms of the cloaking region's area. In fact, the cloaking region in this case can be much larger since a group can include objects involved in a number of different sessions. A query may also not be equally significant at all locations occupied by a group's members. Query privacy is preserved by ensuring that more than

one user in the cloaked region is interested in the same service attribute value as the issuer. We later refer to this as *many-to-one* queries. Given the restriction that the cloaking region must memorize and maintain a fixed set of users, the size of the induced cloaking region becomes an issue with this technique.

Xu and Cai propose an information theoretic measure of anonymity for continuous LBS [188]. They define a *k-anonymity area* as the cloaking region whose entropy is at least k. Although not all k-anonymity areas are historically k-anonymous, their particular implementation called *plainKAA* satisfies the property. However, the algorithm is prone to *inversion attacks* where an adversary uses knowledge of the anonymizing algorithm to breach privacy. The algorithm also fails when some objects stop using the service. The authors propose modifications to plainKAA to resolve the issue of increasing cloaking area, although that may no longer preserve historical k-anonymity.

The most recent of algorithms in this domain is *ProvidentHider* [124]. It uses a maximum perimeter constraint to ensure that cloaking regions are not too large, and the starting set of objects is as big as possible (to take care of leaving objects). This algorithm is later used in our comparative study.

Riboni et al. argue that an adversary may derive an association between a user and a service attribute value based on the distribution of service attribute values in the cloaking regions generated for the user [41]. Therefore, they propose generalizing service attribute values so that the distance between the distribution of service attribute values in cloaking regions for the user and that in regions generated for other users is below a threshold. Besides the fact that generalizing service attributes adversely affects service quality, it is also not clear if performing such generalizations can prevent disclosures emerging from correlations in consecutive cloaking regions. The *t*-closeness model [112] on which their algorithm is based upon is itself known to be sensitive to the distance metric.

The drawbacks present in the above algorithms point out three issues that must be addressed before historical k-anonymity can be efficiently enforced, namely *defunct peers*, *diverging trajectories* and *locality of requests.* Defunct peers are objects that become unavailable in time. As a result, any object dependent on a defunct peer to preserve historical k-anonymity can no longer do so. Also, if objects in a historically k-anonymous cloaking

region diverge from each other, the cloaking region would finally become too large for any reasonable service guarantee. Such divergence also allows an adversary infer crucial associations between the mobile objects and the request. Our primary contribution is therefore an anonymization algorithm, called Continuous ANONymizer (CANON), that implements explicit strategies to resolve each of the three identified issues. In particular, we argue that a cloaking region should be determined using direction information of the objects and show how this restricts the inferences that can be made about the issuer of the request. Large cloaking regions are also avoided by this process. Further, we propose using multiple cloaking regions while issuing a query in order to maintain better service quality.

This chapter also presents a formal analysis of privacy attacks leading to query disclosures in a continuous LBS. We explicitly characterize the privacy threats under consideration and state the background knowledge required to execute the underlying attacks. We model the attacks that can lead to query disclosures and formally show how a technique such as query ℓ-diversity fails to provide query privacy in a continuous LBS. While the threats analyzed here are new in the context of location-based services, similar problems have been explored for privacy protection during the re-publication of dynamic microdata. The principal of *m-invariance* [186] is of particular interest here because of the similarity in privacy issues it helps resolve and those present in a continuous LBS. *m*-Invariance infuses counterfeit entries into a data set so that a particular individual is always associated with a fixed set of sensitive parameter values irrespective of the addition or deletion of entries. Drawing upon the privacy guarantees of *m*-invariance, we formulate the principle of *query m-invariance* and show how it can be used to control the amount of risk present in the use of a continuous LBS. We further propose a cloaking algorithm to efficiently enforce the principle.

The remainder of the chapter is organized as follows. Section 7.3 presents the system architecture and highlights the issues related to historical *k*-anonymity. The CANON algorithm is presented next in Section 7.4. Section 7.5 presents results from an experimental analysis using the algorithm. The issue of query privacy in a continuous LBS is discussed next in Section 7.6. It presents the system architecture and highlights the requirement

for query *m*-invariance. A cloaking algorithm that preserves this privacy property is presented in Section 7.7. Section 7.8 details a comparison of this privacy model to location *k*-anonymity and query ℓ-diversity. Finally, Section 7.9 concludes the chapter.

7.3 Location Anonymity in Continuous LBS

Research in database privacy has shown that formal evaluation of privacy attacks and preservation techniques is difficult without explicitly stating the extent of an adversary's knowledge. Similar arguments have also been made in the context of privacy in snapshot LBS [123]. A continuous LBS susceptible to location-unaware adversaries only can usually preserve privacy by implementing the conventional *k*-anonymity model. A mobile object's position can be concealed by using an arbitrary cloaking region instead of the actual location. In the presence of location-aware adversaries only, exact object locations can be sent to the LBS as long as there is a many-to-one mapping from mobile objects to request identifiers. However, the snapshot *k*-anonymity model is not sufficient in this case. New privacy preserving algorithms are needed that will protect against both types of adversaries. We shall discuss our system architecture and then show how historical k-anonymity is preserved in the presence of both types of adversaries.

7.3.1 System architecture

Our system consists of interactions between three layers – (i) *mobile objects*, (ii) a *trusted anonymity server*, and (iii) a *continuous LBS provider*. Fig. 7.1 depicts a schematic of the architecture. The trusted anonymity server acts as a channel for any communication between mobile objects and continuous LBS providers. All privacy guarantees are therefore enforced at the trusted anonymity server. A mobile object $\mathcal{O}$ initiates a service session by registering itself with the anonymity server. The registration process includes the exchange of current location information ($\mathcal{O}.loc$) and service parameters. The service parameters collectively signify the request to forward to the LBS provider, as well as the anonymity level ($\mathcal{O}.k$) to enforce while doing so. The anonymity server issues a pseudo-identifier and uses it both as a *session identifier* ($\mathcal{O}.sid$) with the mobile object and as an *object identifier* when communicating with the LBS provider. A set of cloaking regions

Figure 7.1: Schematic of the system architecture.

is then generated for the requesting object and multiple range queries are issued to the LBS provider for these regions. Communication between the anonymity server and the LBS provider is always referenced using the object identifier so that the LBS can maintain service continuity. The candidate results retrieved from the LBS provider are filtered at the anonymity server and then communicated to the mobile object. Subsequent location updates from the mobile object is handled in a similar fashion (with the pre-assigned session identifier) until the anonymity level cannot be satisfied or the service session is terminated. A request is *suppressed* (dropped) when the anonymity requirements can no longer be met within the same service session. A new identifier is then used if the mobile object re-issues the same request. Therefore, the number of consecutive requests served with a single identifier serves as a metric to evaluate the efficiency of the anonymizing algorithm. Higher values signify longer continuity, implying better quality of service. We further assume that an object does not change its service parameters during a session. A separate session is started if a request with different service parameters is to be made. Therefore, an object can have multiple sessions running at the same time, each with a different session identifier. Without any loss of generality, we assume that a mobile object $\mathcal{O}$ has a single running session at most; identified by $\mathcal{O}.sid$.

7.3.2   Historical k-anonymity

The primary purpose of a cloaking region is to make a given mobile object $\mathcal{O}$ indistinguishable from a set of other objects. This set of objects, including $\mathcal{O}$, forms the *anonymity set* of $\mathcal{O}$. Objects in the anonymity set shall be referred to as *peers* of $\mathcal{O}$ and denoted by $\mathcal{O}.peers$. A cloaking region for $\mathcal{O}$ is usually characterized by the minimum bounding

rectangle (MBR) of the objects in $\mathcal{O}.peers$. A range query is defined over such an MBR. Therefore, the cloaking region is typically required to satisfy two properties in order to achieve an acceptable balance between privacy and service quality.

- *Sufficient anonymity:* The degree of anonymity of a cloaking region is characterized by the ambiguity introduced in differentiating one object from another with respect to the actual issuer of the request. Under the conventional *k*-anonymity model, this is achieved by ensuring that the anonymity set of a mobile object contains at least $k-1$ other objects. The value of k determines the level of privacy, and can either be a user-specific or a system parameter.

- *Sufficient quality:* Since the result set of a range query caters to every mobile object possibly present inside the cloaking region's MBR, a bigger candidate set is more likely for a larger MBR. Bigger candidate sets mean longer durations of time communicating the result with the anonymity server, longer computational overhead filtering the results, as well as, possibly higher service costs owing to I/O overheads. Therefore, a smaller cloaking region is preferred over larger ones. This is typically enforced by specifying a maximum spatial resolution as a constraint for MBR sizes. In addition, a maximum temporal resolution can also be specified to delay service requests so that smaller cloaking regions can be generated by watching the location of moving objects over a longer duration of time.

As demonstrated in a number of prior works, achieving reasonable levels of anonymity and service quality is not difficult in the case of a snapshot LBS. Since snapshot services consider every request to be mutually independent, service quality does not depend on the ability to correlate consecutive requests from the same mobile object. However, in our assumed architecture for a continuous LBS, maintaining the two properties is significantly difficult.

Consider the movement pattern of the objects depicted in Fig. 7.2. A 3-anonymous MBR is computed for O_1 during three consecutive location updates. If O_1's requests at the three time instances are mutually independent from each other (as in a snapshot LBS), then privacy level of O_1 is maintained at 3-anonymity across the different MBRs.

Figure 7.2: Conventional *k*-anonymity and historical *k*-anonymity. Size of historically *k*-anonymous MBR is influenced by movement of objects in O_1's anonymity set at time t_1.

However, when the same identifier is associated with all the MBRs (as in a continuous LBS), it only requires an adversary the knowledge of O_1, O_2 and O_3's positions at time t_1, t_2 and t_3 (a location-aware adversary) to infer that the requests are being issued by object O_1. This is because O_1 is the only common object in the intersection of the three cloaking regions. We refer to this as a case of *full disclosure.* Further, any subsequent MBR associated with the same identifier can be associated back to O_1, thereby revealing the trajectory of O_1 after time t_3. Assuming that each object is equally likely to be included in another object's cloaking region, the probability of full disclosure is unacceptably high.

Remark 1 Let $A_1, \ldots, A_n$ be a sequence of anonymity sets corresponding to $n > 1$ consecutive *k*-anonymous cloaking regions for a mobile object $\mathcal{O}$, generated from a collection of N mobile objects. Then, the probability that the intersection of the anonymity sets $\mathcal{S}_n = \underset{i}{\cap} A_i$ has at least p objects, $p > 1$, is

$$Pr(|\mathcal{S}_n| \geq p) = \left(\prod_{i=1}^{p-1} \frac{k-i}{N-i} \right)^n. \tag{7.1}$$

Since $\mathcal{S}_n$ is the intersection of anonymity sets corresponding to the same object $\mathcal{O}$, we have $\mathcal{O} \in \mathcal{S}_n$. Let $\mathcal{N}$ be the collection of N mobile objects. Consider the class of sets $\mathcal{H} = \{\mathcal{O}\} \cup Q$ where $Q \subseteq \mathcal{N} - \{\mathcal{O}\}$ and $|Q| = p - 1; p > 1$. The probability that we seek is $Pr(H \subseteq \mathcal{S}_n)$, for all $H \in \mathcal{H}$.

Consider a set A_j. We assume that each cloaking region minimally satisfies k-anonymity, i.e. $\forall j, |A_j| = k$. Hence, A_j has $k - 1$ objects other than $\mathcal{O}$. The number of ways such sets can be formed is $\binom{N-1}{k-1}$. Further, the number of such sets that has H as a subset is $\binom{N-p}{k-p}$. Therefore, $Pr(H \subseteq A_j) = \binom{N-p}{k-p} / \binom{N-1}{k-1} = \prod_{i=1}^{p-1} \frac{k-i}{N-i}$. Also, for $H \subseteq \mathcal{S}_n$, we have $\forall j, H \subseteq A_j$ and which implies $\mathcal{S}_n$ has at least as many objects as in H, i.e. p. The probability $Pr(|\mathcal{S}_n| \geq p)$ is therefore

$$Pr(\forall H \in \mathcal{H}, \forall j, H \subseteq A_j) = \left(\prod_{i=1}^{p-1} \frac{k-i}{N-i} \right)^n. \tag{7.2}$$

Remark 2 If $k \leq \frac{N+1}{2}$ then the probability of full disclosure is at least $\frac{3}{4}$. The full disclosure risk is given as $\mathcal{D}_{full} = Pr(|\mathcal{S}_n| = 1) = Pr(|\mathcal{S}_n| \geq 1) - Pr(|\mathcal{S}_n| \geq 2)$. Since intersection of the anonymity sets contain at least one object, we have $Pr(|\mathcal{S}_n| \geq 1) = 1$. Hence, $\mathcal{D}_{full} = 1 - (\frac{k-1}{N-1})^n$. With $k \leq \frac{N+1}{2}$, or $\frac{k-1}{N-1} \leq \frac{1}{2}$, we have

$$\mathcal{D}_{full} \geq 1 - \frac{1}{2^n} \geq 1 - \frac{1}{2^2} = \frac{3}{4}. \tag{7.3}$$

We also observe in Fig. 7.2 that it does not require knowledge on the objects' locations at all three time instances in order to breach O_1's privacy. In fact, location knowledge at time instances t_1 and t_2 is sufficient to lower O_1's privacy to 2-anonymity. This is referred to as a *partial disclosure.* Such disclosures occur when the intersection of anonymity sets (corresponding to the same object) contains less than the desired number of peers (the anonymity level k).

Remark 3 Although the likelihood of full disclosure reduces as k approaches N, partial disclosures can still render an anonymization algorithm ineffective. Fig. 7.3 shows the probability of partial disclosure for $N = 100$ and various anonymity levels. The partial disclosure typically increases as k increases from $(N + 1)/2$. For the case when the adversary's location knowledge extends beyond two cloaking regions, the situation is much worse as full disclosures are more likely for a larger range of anonymity levels.

A straightforward extension of the conventional k-anonymity model that can counter risks of full and partial disclosures in a continuous LBS is to ensure that all anonymity sets within a service session contain at least k common objects. In other words, k-anonymity

Figure 7.3: Risk of partial disclosure for varying anonymity levels $k > (N+1)/2$. $N = 100$ and n is the number of cloaking regions for which adversary has full location information.

should be satisfied on the intersection of anonymity sets, referred to as historical k-anonymity.

Definition 7.1 HISTORICAL k–ANONYMITY. Let $A_1, \ldots, A_n$ be a sequence of anonymity sets corresponding to the cloaking regions with the same identifier and at time instants $t_1, \ldots, t_n$, $t_i > t_j$ for $i > j$, respectively. The anonymity set A_i is then said to satisfy historical k-anonymity if $|A_1 \cap \ldots \cap A_i| \geq k$.

In other words, the sequence of anonymity sets preserve historical k-anonymity if all subsequent sets after A_1 contain at least k same objects from A_1. Therefore, even an adversary with complete location knowledge of all the mobile objects in all the cloaking regions can not associate the object identifier to less than k objects. This prohibits both cases of full and partial disclosures. Fig. 7.2 depicts how the cloaking regions should change over time in order to ensure that object O_1 always has historical 3-anonymity.

7.3.3 Implications

Since anonymity sets under historical k-anonymity also satisfy the conventional k-anonymity model, privacy breaches involving restricted space identification, observation identification and location tracking by location-unaware adversaries are implicitly impeded. In addition, historical k-anonymity also impedes session association attacks by

location-aware adversaries. Hence, the privacy guarantees of historical k-anonymity in a continuous LBS is similar to that of k-anonymity in a snapshot LBS. However, maintaining acceptable levels of service can become increasingly difficult in case of historical k-anonymity. We have identified three issues for consideration that impact the practical usage of historical k-anonymity.

(1) *Defunct peers:* A defunct peer in an anonymity set is an object that is no longer registered with the anonymity server. As a result, it can no longer be ascertained that a cloaking region includes the peer. If the first cloaking region generated during a particular session contains exactly k objects, then every other anonymity set in that session must contain the same k objects for it be historically k-anonymous. A defunct peer in this case does not allow subsequent cloaking regions to satisfy historical k-anonymity and introduces possibilities of partial disclosure. Such possibilities can be reduced either by suppressing the requests that fail to meet historical k-anonymity, or by reducing the impact of a defunct peer on the achievement of a desired anonymity level. The latter is achieved in *ProvidentHider* by generating the first cloaking region to include as many peers as possible within a MBR of fixed perimeter. It is expected that more than k objects will get included in the first anonymity set, thereby providing some flexibility for peers to defunct before influencing the minimum required anonymity set size. However, this approach cannot account for the varying density of objects across time and space. A highway, for example, can be very sparsely populated with cars during off-peak hours, but can be heavily occupied during rush hour. Using a fixed perimeter can also lead to a high percentage of suppressed requests where object distribution is relatively sparse.

(2) *Diverging peer trajectories:* The trajectories of peers influence the size of a cloaking region (satisfying historical k-anonymity) over time. Refer to Fig. 7.2. The MBR for object O_1 becomes increasingly larger owing to the trajectory of object O_3. Bigger cloaking regions have a negative impact on service quality. In general, the more divergent the trajectories are, the worse is the effect. Algorithms that use a maximum spatial resolution will not be able to facilitate service continuity as spatial constraints will not be met. In the worst case, trying to satisfy historical k-anonymity will unwantedly result in requests being serviced on a snapshot basis.

(3) *Locality of requests:* The significance of a particular service request can often be correlated with the locality where it originated. For instance, let us assume that the region shown in Fig. 7.2 corresponds to a urban locality. Further, object O_1 issues a request to periodically update itself with information (availability, price, etc.) on the nearest parking garage. At time instance t_1, an adversary cannot infer which object (out of O_1, O_2 and O_3) is the actual issuer of the request. All objects in this case are in a locality where the request holds equal significance. However, as O_3 moves away from the urban locality (suspiciously ignoring the high concentration of garages if it were the issuer), an adversary can infer that the issuer of the request is more likely to be O_1 or O_2. We say that these two objects are still in the *locality of the request*. If historical *k*-anonymity is continued to be enforced, O_3 (and most likely O_2 as well) will be positioned in different localities, thereby allowing an adversary infer with high confidence that O_1 is the issuer of the request. The significance of requests and the locality of peers can also help an adversary infer a non-uniform distribution of request issuer likelihoods. Peers ought to have some level of directional similarity in order to avoid such inferences.

Note that these three issues are primarily applicable in the context of a continuous LBS, more specifically when working in conjunction with an anonymity server enforcing historical *k*-anonymity. Defunct peers is not an issue in snapshot LBS since the set of peers can be decided on a per request basis. Further, since the same peers need not be present in subsequent anonymity sets, their trajectories do not influence the size of the next privacy preserving cloaking region. Differences in locality also do not provide additional inferential power to an adversary. However, in a continuous LBS, these three issues are direct residues of providing privacy by historical *k*-anonymity. Our attempt in this work is to find an effective technique to enforce historical *k*-anonymity in a way that produces minimal impact on the service quality. The CANON (Continuous ANONymizer) algorithm presented in the next section highlight the strategies adopted to do so.

7.4   The CANON Algorithm

CANON is an anonymization algorithm that enforces historical *k*-anonymity for use with a continuous LBS. The algorithm defines explicit procedures to handle each of the

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Procedure 7.1 CANON(Object $\mathcal{O}$)

Input: Mobile object $\mathcal{O}$ (includes all associated data).

Output: A set of peer groups (one of them includes $\mathcal{O}$); **null** if request is suppressed (cannot satisfy anonymity).

1: **if** ($\mathcal{O}.sid =$ **null**) **then**
2:     $\mathcal{O}.peers = CreatePeerSet(\mathcal{O})$
3:     $\mathcal{O}.sid =$ new session identifier
4: **else**
5:     Remove defunct objects in $\mathcal{O}.peers$
6: **end if**
7: **if** ($|\mathcal{O}.peers| < \mathcal{O}.k$) **then**
8:     $\mathcal{O}.sid =$ **null**
9:     **return null**
10: **end if**
11: $peerGroups = PartitionPeerSet(\mathcal{O})$
12: **if** ($\exists g \in peerGroups$ such that $|g| < 2$) **then**
13:     $\mathcal{O}.sid =$ **null**
14:     **return null**
15: **end if**
16: **return** $peerGroups$

---

three potential issues identified in the previous section. An overview of the algorithm is shown in Procedure 7.1.

CANON is initiated by the anonymity server whenever it receives a request from a mobile object $\mathcal{O}$. The algorithm starts by first checking if $\mathcal{O}$ has an open session with respect to the current request. If it finds one then the set of peers is updated by removing all defunct peers from the set. Otherwise, a peer set is generated for $\mathcal{O}$ through the procedure *CreatePeerSet* and a session identifier is assigned. The newly generated (or updated) peer set must have at least $\mathcal{O}.k$ objects in order to continue to the next step; otherwise the request is suppressed and the session is terminated. Historical *k*-anonymity is ensured at the end of Line 10 since at least *k* objects inserted into $\mathcal{O}.peers$ by *CreatePeerSet* is still registered with the anonymity server. The next step is to divide the peer set into groups over which the range queries will be issued. A *peer group* is defined as a subset of $\mathcal{O}.peers$. *PartitionPeerSet* divides $\mathcal{O}.peers$ into disjoint peer groups. We shall often use the term "object's peer group" to signify the group that contains $\mathcal{O}$. Each peer group defines a smaller cloaking region than that defined by the entire peer set and reduces the impact of diverging trajectories on service quality. The peer groups returned by CANON

Figure 7.4: Peer set partitioning into groups over which multiple range queries are issued with the same identifier.

are used to issue multiple range queries (one for each) with the same object identifier. Fig. 7.4 depicts this process. Line 12 checks that each peer group contains at least two objects in order to avoid the disclosure of exact location information (of any object) to location-unaware adversaries.

All object agglomerations, namely into peer sets and then into peer groups, are performed so that the *reciprocity* property is satisfied. This property states that the inclusion of any two objects in a peer set (group) is independent of the location of the object for which the peer set (groups) is being formed. Reciprocity prevents inversion attacks where knowledge of the underlying anonymizing algorithm can be used to identify the actual object. The *Hilbert Cloak* algorithm [98] was first proposed in this context for the conventional k-anonymity model. Hilbert Cloak orders the objects according to their Hilbert indices (index on a space filling curve) and then groups them into buckets of size k. The peer set of an object is the bucket that contains the object. The peer set is the same for any object in the same bucket. *CreatePeerSet* and *PartitionPeerSet* thus use Hilbert-sorted lists to incorporate these properties.

7.4.1 Handling defunct peers

As mentioned earlier, defunct peers can influence the lifetime of a service session by reducing the peer set size to below the limit that satisfies historical k-anonymity. The resolution is to include more than k objects in the first peer set. An indirect way to achieve

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this is to specify a maximum spatial boundary around the requesting object's location and then include all objects within that boundary into the peer set. However, this method suffers from object distribution problems as in *ProvidentHider*. It may be possible to adapt the spatial boundary according to available traffic information. Even so, determining the adjustments could be a difficult task. Using spatial boundaries also cannot account for the relative differences in MBR sizes corresponding to varying anonymity requirements. For example, an area of $1\,km^2$ may be sufficient to have enough peers to satisfy a historical 2-anonymity requirement, but may not be so to satisfy a stronger requirement (say historical 50-anonymity).

A more direct method to resolve the issue is to specify the desired peer set size explicitly. This removes any dependency on how the objects are distributed and the area required to cover a reasonable number of them. We can specify the size as a sufficiently big constant. However, this strategy favors objects with weaker anonymity requirements as their peer sets are allowed a comparatively higher number of peers to defunct. For instance, a constant peer set size of 20 would allow the anonymizer to tolerate up to 18 defunct peers to preserve historical 2-anonymity, but only 5 defuncts to preserve historical 15-anonymity. Therefore, the strategy adopted in CANON uses an *oversize factor τ* that relatively specifies the number of extra peers that must be included in the peer set. The minimum initial size of the peer set of an object $\mathcal{O}$ is equal to $(1 + \tau) \times \mathcal{O}.k$ with this strategy. We say "minimum" because other parameters introduced later can allow more peers to be included. Use of an oversize factor prevents the problem associated with constant peer set sizes. Note that since CANON partitions the peer set into further groups before issuing a query, the area of the cloaking region defined by the enlarged peer set has little or no influence on service quality. However, we would still not want the area to expand extensively in order to curb the issue of request locality.

7.4.2 Deciding a peer set

The *CreatePeerSet* procedure determines the initial peer set for an object. This is a crucial step since anonymity for subsequent requests in the session will be evaluated on the basis of the initial peer set. We have introduced the oversize factor to reduce the

chances of a peer set failing to meet anonymity requirements due to defunct peers. In addition, we also need to ensure that majority of the objects in the peer set are in the locality of the request. There are two requirements to address in this regard.

1. Objects in the peer set should define an area where the request is equally significant to all the peers.

2. Objects in the peer set should move so that the defined area does not expand "too much".

The first requirement will prohibit the inclusion of peers that are positioned in a locality where the issued request is unlikely to be made. The second requirement addresses locality of requests in the dynamic scenario where the trajectories of the peers could be such that they are positioned in very different localities over time. Preventing the MBR of the peer set from expanding prohibits peers from being too far away from each other. The first requirement can be fulfilled by choosing peers according to the Hilbert Cloak algorithm. Peers chosen according to Hilbert indices will induce a small MBR, thereby ensuring that they are more likely to be in the same locality. However, a peer set generated by this process cannot guarantee that the second requirement will be fulfilled for long. This is because the neighbors of an object (according to Hilbert index) may be moving in very different directions.

It is clear from the above observation that the direction of travel of the objects should be accounted for when selecting peers. The direction of travel is calculated as a vector from the last known location of the object to its current location, i.e. if $\mathcal{O}.loc_1 = (x_1, y_1)$ and $\mathcal{O}.loc_2 = (x_2, y_2)$ are the previously and currently known positions of $\mathcal{O}$ respectively, then the direction of travel is given as $\mathcal{O}.dir = \mathcal{O}.loc_2 - \mathcal{O}.loc_1 = (x_2 - x_1, y_2 - y_1)$. $\mathcal{O}.dir$ is set to $(0,1)$ (north) for newly registered objects. A *θ-neighborhood* for $\mathcal{O}$ is then defined as the set of all objects whose direction of travel is within an angular distance θ (say in degrees) from $\mathcal{O}.dir$. Therefore, a $0°$-neighborhood means objects traveling in the same direction, while a $180°$-neighborhood contains all objects. If all peers are chosen within a $0°$-neighborhood then it is possible that the area defined by the initial peer set will more or less remain constant over time. However, the initial area itself could be very large due

Procedure 7.2 CreatePeerSet(Object $\mathcal{O}$)

Input: Mobile object $\mathcal{O}$ (includes all associated data), and system globals τ, θ and α_{full}.
Output: A set of peer objects (including $\mathcal{O}$).

1: $\mathcal{L}$ = set of available mobile objects sorted by their Hilbert index
2: $k_{of} = (1 + \tau) \times \mathcal{O}.k$; $\mathcal{P} = \phi$
3: **repeat**
4:     $\mathcal{L}_c = \phi$
5:     **for all** ($l \in \mathcal{L}$ in order) **do**
6:         **if** ($|\mathcal{L}_c| \geq k_{of}$ **and** $AreaMBR(\mathcal{L}_c \cup \{l\}) > \alpha_{full}$) **then**
7:             **break**
8:         **end if**
9:         $\mathcal{L}_c = \mathcal{L}_c \cup \{l\}$
10:     **end for**
11:     $\mathcal{P}_{prev} = \mathcal{P}$; $f = 1$; $\mathcal{O}_{pivot}$ = first object in $\mathcal{L}_c$
12:     **repeat**
13:         $\mathcal{P} = (f\theta)$-neighbors of $\mathcal{O}_{pivot}$ in $\mathcal{L}_c$
14:         $f = f + 1$
15:     **until** ($|\mathcal{P}| \geq \min(k_{of}, |\mathcal{L}_c|)$)
16:     $\mathcal{L} = \mathcal{L} - \mathcal{P}$
17: **until** ($\mathcal{O} \in \mathcal{P}$)
18: **if** ($|\mathcal{P}| < k_{of}$) **then**
19:     $\mathcal{P} = \mathcal{P} \cup \mathcal{P}_{prev}$
20: **else if** ($|\mathcal{L}| < k_{of}$) **then**
21:     $\mathcal{P} = \mathcal{P} \cup \mathcal{L}$
22: **end if**
23: **return** $\mathcal{P}$

---

to the non-availability of such peers within a close distance. On the other hand, using a 180°-neighborhood essentially allows all objects to be considered and hence the area can be kept small by including close objects. Of course, the area may increase unwantedly over time. Peer set generation is therefore guided by two system parameters in CANON - the *neighborhood step size* θ and the *full-MBR resolution* α_{full}. The neighborhood step size specifies the resolution at which the θ-neighborhood is incremented to include dissimilar (in terms of travel direction) peers. The full-MBR resolution specifies some area within which the issued request is equally likely to have originated from any of the included objects, thereby making it difficult for an adversary to eliminate peers based on position and request significance. For small values of θ and some α_{full}, all objects in a peer set would ideally move in a group, in and out of a locality. Procedure 7.2 outlines the pseudo-code of *CreatePeerSet*. We assume the existence of a function *AreaMBR* that returns the area of the minimum bounding rectangle of a set of objects.

CreatePeerSet first creates a sorted list $\mathcal{L}$ of all registered objects according to their Hilbert indices. It then continues to divide them into buckets (starting from the first one in the sorted list) until the one with $\mathcal{O}$ is found (Lines 3-17). Every time a bucket is formed, $\mathcal{L}$ is updated by removing all objects in the bucket from the list (Line 16). Lines 5-10 determine a set $\mathcal{L}_c$ of candidate objects that can potentially form a bucket. Starting from the first available object in $\mathcal{L}$, we continue to include objects in $\mathcal{L}_c$ as long as the minimum peer set size (denoted by k_{of} and decided by the oversize factor) is not met, or the area of the MBR of included objects is within the full-MBR resolution. Note that, as a result of this condition (Line 6), the minimum required size of the peer set receives more prominence than the resulting area. Hence, the full-MBR resolution is only a guiding parameter and not a constraint. Next, Lines 12-15 select k_{of} objects from the candidate set to form a bucket. The first object in $\mathcal{L}_c$ is chosen as a pivot and all objects in the θ-neighborhood of the pivot are included in the bucket. If the bucket is not full up to its capacity (k_{of}) and more objects are present in $\mathcal{L}_c$, then the neighborhood is increased by the step size θ. By the end of this process, the bucket would either contain k_{of} objects or there are less than k_{of} objects in $\mathcal{L}_c$. The latter is only possible when list $\mathcal{L}$ contains less than k_{of} objects, i.e. the last bucket is being created. Note that object $\mathcal{O}$ is not explicitly used anywhere to decide the buckets, thereby guaranteeing reciprocity. Once the bucket with $\mathcal{O}$ is found, two more checks are required (Lines 18-22). First, if $\mathcal{O}$'s bucket has less than k_{of} objects (possible if it is the last one), then it is merged with the previous bucket. Second, if the number of objects remaining in $\mathcal{L}$ is less than k_{of} (implying $\mathcal{O}$'s bucket is second to last), then the remaining objects are included into $\mathcal{O}$'s bucket to maintain reciprocity.

CreatePeerSet uses θ-neighborhoods and the full-MBR resolution to balance between dissimilar peers and the resulting MBR area. While the step size θ allows incremental selection of dissimilar peers, α_{full} guides the extent of increment admissible to generate a localized peer set. Note that the creation of a peer set is a one time procedure every service session. Hence, a good estimation of the direction of travel is required to avoid diverging trajectories. One possibility is to obtain destination points of objects and generate an average direction of travel. An average direction can also be calculated based on the

displacement vector of the object from its starting position. One can also estimate a direction of travel based on a set of last known locations. CANON uses an instantaneous direction vector. We believe this method performs reasonably well in road networks, although the efficacy of other techniques remains to be determined.

7.4.3 Handling a large MBR

The full-MBR resolution parameter is used to control breaches related to request localities. Typical values are in the range of 10 to 50 km^2. The parameter is therefore not intended to help generate cloaking regions with small MBRs. A continuous LBS would require a much finer resolution to deliver any reasonable service. Further, depending on variations in velocity and the underlying road network, some extent of expansion/contraction of the MBR is very likely. The MBR of a peer set is therefore not a good candidate to issue the range queries. Instead, the peer set is partitioned into multiple disjoint groups by *PartitionPeerSet*. Partitioning of the peer set eliminates empty spaces between peers (introduced in the first place if trajectories diverge) and produces smaller MBRs for the range queries [169]. This partitioning can be done either in a way such that each peer group has a minimum number of objects or each peer group has a maximum spatial resolution. The former approach cannot guarantee that the resulting MBR will have an acceptable area. The latter method is adopted in CANON where the maximum spatial resolution of a peer group is specified as the *sub-MBR resolution* α_{sub}. α_{sub} is relatively much smaller than α_{full}. Procedure 7.3 outlines the partitioning method.

The partitioning is performed in a manner similar to Hilbert Cloak, with the difference that each bucket now induces an area of at most α_{sub} instead of a fixed number of objects. Starting from the first object in the Hilbert-sorted peer set, an object is added to a bucket as long as the sub-MBR resolution is met (Line 6); otherwise the current bucket is a new peer group (Line 8) and the next bucket is created (Line 9). Reciprocity is preserved as before. Note that the pseudo-code in Procedure 7.3 does not handle the case when a peer group contains only one object. Procedure 7.1 checks that such groups do not exist (safeguard against location-unaware adversaries); otherwise the request is suppressed. However, the partitioning algorithm itself can relax the sub-MBR resolution when a peer group with

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Procedure 7.3 PartitionPeerSet(Object $\mathcal{O}$)

Input: Mobile object $\mathcal{O}$ (includes all associated data) and system global α_{sub}.
Output: A set of peer groups.

1: Sort objects in $\mathcal{O}$.*peers* by their Hilbert index
2: *peerGroups* $= \phi$
3: *bucket* $= \phi$
4: **for all** ($l \in \mathcal{O}$.*peers* in order) **do**
5:    **if** (*AreaMBR*(*bucket*$\cup\{l\}$) $\leq \alpha_{sub}$) **then**
6:       *bucket* $= bucket\cup\{l\}$
7:    **else**
8:       *peerGroups* $=$ *peerGroups* $\cup \{bucket\}$
9:       *bucket* $= \{l\}$
10:   **end if**
11: **end for**
12: *peerGroups* $=$ *peerGroups* $\cup \{bucket\}$
13: **return** *peerGroups*

---

a single object is found. One possible modification is to merge any peer group having a single object with the group generated prior to it. Another parameter-less technique is to create partitions that result in the minimum average peer group MBR with the constraint that each group must have at least two objects. We have kept these possibilities for future exploration.

7.5 Empirical Study

The experimental evaluation compares the performance of CANON with the *ProvidentHider* algorithm. For every new request, *ProvidentHider* first groups all available objects from a Hilbert-sorted list such that each bucket holds $\mathcal{O}$.*k* objects; more if adding them does not violate a maximum perimeter (P_{max}) constraint. The peer set of an object is the bucket that contains the object. A range query is issued over the area covered by the objects in the peer set only if the maximum perimeter constraint is satisfied; otherwise the request is suppressed. As the peers move, the assigned peer set is partitioned as before and the bucket containing the issuing object is assigned as the new peer set. The new peer set must satisfy the perimeter constraint, as well as have at least $\mathcal{O}$.*k* objects. *ProvidentHider* also differentiates between objects based on their location. We assume that all objects are in the visible domain with respect to such differentiation. We measure a number of statistics to evaluate the performance.

Figure 7.5: Trace data generated on Chamblee region of Georgia, USA. The mean speed, standard deviation and traffic volume on the three road types used are shown.

- *service continuity:* average number of requests served in a session.

- *service failures:* percentage of suppressed requests.

- *safeguard against location-unaware adversaries:* average size of the peer group to which the issuing object belongs.

- *request localization:* average factor of increase in MBR area of an object's first peer set.

7.5.1 Experimental setup

We have generated trace data using a simulator [73] that operates multiple mobile objects based on real-world road network information available from the National Mapping Division of the US Geological Survey. We have used an area of approximately $168\,km^2$ in the Chamblee region of Georgia, USA for this study (Fig. 7.5). Three road types are identified based on the available data – expressway, arterial and collector. Real traffic volume data is used to determine the number of objects in the different road types [77]. The total number of objects on a road type vary proportionate to the total length and traffic volume of the road type, and reciprocally to the average speed of the objects. The mean

speed, standard deviation and traffic volumes on the road types are shown in the figure. Using the number of objects on each road type, the simulator randomly places them on the network and moves them around. The objects move with a speed drawn from a normal distribution, randomly making turns and changing speed at junctions. The simulator maintains the traffic volume statistics while moving the objects.

The used traffic volume information results in 8,558 objects with 34% on expressways, 8% on arterial roads and 58% on collector roads. The trace data consists of multiple records spanning one hour of simulated time. A record is made up of a time stamp, object number, *x* and *y* co-ordinates of object's location, and a status indicator. The status indicator signifies if the object is registered at the anonymity server. An object's status starts off randomly as being active or inactive. The object remains in the status for a time period drawn from a normal distribution with mean 10 minutes and standard deviation 5 minutes. The status is randomly reset at the end of the period and a new time period is assigned. The granularity of the data is maintained such that the Euclidean distance between successive locations of the same object is approximately 100 meters. Each object has an associated *k* value drawn from the range [2,50] by using a Zipf distribution favoring higher values and with the exponent 0.6. The trace data is sorted by the time stamp of records.

During evaluation, the first minute of records is used only for initialization. Subsequently, the status of each record is used to determine if the object issues a request. Only an active object is considered for anonymization. If the object was previously inactive or its prior request was suppressed, then it is assumed that a new request has been issued. Otherwise, the object is continuing a service session. The anonymizer is then called to determine the cloaking region(s), if possible. The process continues until the object enters an inactive (defunct) state. Over 2,000,000 anonymization requests are generated during a pass of the entire trace data.

Default values of other algorithm parameters are set as follows: $\tau = 0.0$, $\alpha_{full} = 25\,km^2$, $\alpha_{sub} = 1\,km^2$, $\theta = 180°$ and $P_{max} = 5000\,m$. All parameters take their default values unless stated otherwise. A $5000\,m$ perimeter constraint for *ProvidentHider* is approximately an area of $1.6\,km^2$. Compared to that, α_{sub} has a smaller default value. The precision is

around 1000 m (assuming a square area) which serves reasonably well for a Pay-As-You-Drive insurance service. We also study how CANON performs for services that require a higher precision. The full-MBR resolution of 25 km^2 evaluates to a locality about $\frac{1}{32}^{th}$ the size of New York City. The entire map is assumed to be on a grid of $2^{14} \times 2^{14}$ cells (a cell at every meter) while calculating the Hilbert indices [116]. Objects in the same cell have the same Hilbert index.

7.5.2 Comparative performance

Fig. 7.6a shows the average number of requests served in a session for different anonymity requirements. *ProvidentHider* demonstrates poor performance for higher k values, almost to the extent of one request per session. Comparatively, CANON maintains much better service continuity. As mentioned earlier, using a fixed area for varying anonymity requirements makes it difficult for *ProvidentHider* to keep the peer set within the required size. The task is much difficult for bigger peer sets as the algorithm does not consider the issue of diverging trajectories. In fact, a high percentage of the requests is suppressed for higher values of k (Fig. 7.6b). CANON's performance also seems to fluctuate depending on the oversize factor. In general, a maximum peer set size slightly larger than the minimum required (for example $\tau = 0.25$) gives the best performance, while any further increase degrades it. While a few extra peers is useful to handle defunct peers, having a much larger peer set implies having objects over a larger area and often far away from each other (over time). Therefore, it is possible that some peer groups are formed with a single object owing to the sub-MBR constraint. Requests are then suppressed in the absence of a strategy to handle such peer groups. This is also corroborated by the similar trend in request suppression.

Fig. 7.6c shows the average size of the peer group to which the requesting object belonged. Bigger sizes imply more ambiguity for a location-unaware adversary while trying to identify the actual object. While the sizes are larger for higher k (implying that safeguard against location-unaware adversaries scale as more safeguard against location-aware adversaries is desired), there does not seem to be any observable trend due to changes in the oversize factor.

Figure 7.6: Comparative performance of CANON with ProvidentHider (PH) for different anonymity requirements (*k*) and oversize factors (τ). (a) Average number of requests served in a session. (b) Percentage of requests suppressed (anonymity requirement could not be satisfied). (c) Average size of requesting object's peer group.

7.5.3 Impact of parameters

Each parameter in CANON is intended to address a specific issue with the use of historical k-anonymity. We perform some parametric studies to demonstrate the consequences of varying these parameters. The neighborhood step size is varied between $1°$ and $180°$, and performance is observed for three different settings of the sub-MBR ($\alpha_{sub} = 0.25, 1.0$ and $4\,km^2$) and full-MBR ($\alpha_{full} = 10, 25$ and $50\,km^2$) resolutions. Note that increasing/decreasing the full-MBR resolution will have no impact on peer sets if the required number of objects is always present within a small area. Hence, studies on α_{full} with $\theta = 180°$ produces same results. We therefore use a neighborhood step size of $15°$ while observing the impact of α_{full}. Having a small θ forces CANON to consider more similar/dissimilar peers depending on whether we have a large/small α_{full}. All parameters other than the ones mentioned take their default values.

7.5.3.1 Neighborhood step size θ

Performance in terms of service continuity does not differ a lot for varying step size (Fig. 7.7a). Some differences are observed for lower ranges of k $(2 - 15)$ where larger step sizes show a better performance. Differences are more prominent in terms of peer group size where a bigger neighborhood improves the safeguard against location-unaware adversaries (Fig. 7.7b). This behavior is expected since bigger neighborhood sizes allow the inclusion of more dissimilar peers, thereby inducing bigger peer groups due to the possibly close proximity of objects. The statistic of interest is the size of the MBR area defined by the objects in the peer set. Fig. 7.7c indicates that this area remains almost constant for the smaller step sizes, specifically for the more frequently requested anonymity levels (higher k), implying that the objects in a peer set move together as a group. We are not concerned about the high increase in area for lower k values $(2 - 15)$ because the initial area defined by the peer sets in this range is not too large to begin. We believe the instantaneous calculation of direction vectors impact performance in this range. The area increases by more than two folds (across different anonymity requirements) when direction of travel is ignored ($\theta = 180°$), while it is contained within a two factor increase for the other experimented values. For the sake of perception, a four times increase in

Figure 7.7: Impact of different neighborhood step size θ on CANON. (a) Average number of requests served in a session. (b) Average size of requesting object's peer group. (c) Average number of folds increase in requesting object's MBR area over a session.

the area of a square doubles the distance between diagonally opposite points. Therefore, most peers can be maintained in the locality of the request by using θ-neighborhoods. Even a step size of $90°$ provides considerable benefits in this regard without severely affecting the other performance measures.

7.5.3.2 Sub-MBR resolution α_{sub}

Smaller sub-MBR resolutions mean higher precision in the range queries. However, they also mean higher chances of smaller peer groups, often ones with a single object. With reference to Fig. 7.8 (top row), a smaller α_{sub} results in a higher rate of failures, inducing shorter service sessions. Services requiring high location precision will therefore fail to provide longer service continuity. An object's peer group size is also comparatively smaller. Improvements in service continuity is more prominent for weaker anonymity requirements as α_{sub} is increased. However, improvements in peer group size is more noticeable in higher k values. In effect, finding a suitably balanced α_{sub} can help achieve good overall performance. α_{sub} is decided by the service requirements in most cases. Nonetheless, depending on how stringent the requirement is, both privacy (from location-unaware adversaries) and service quality may have scope for improvement.

7.5.3.3 Full-MBR resolution α_{full}

The full-MBR resolution is observed to have little or no influence on the average number of requests served in a session (Fig. 7.8 bottom row). However, larger areas tend to have higher percentage of failures. A possible explanation is as follows. A larger area with a small step size means similar objects are preferred over the proximity of objects. As a result, a peer set includes objects distributed far apart. This leads to the suppression of requests when the sub-MBR constraint is imposed on the formation of peer groups. Objects far apart cannot be grouped without violating the constraint. This also results in a comparatively smaller peer group size. On the other hand, a smaller area allows inclusion of close proximity objects at the expense of similarity. The sub-MBR constraint is therefore easier to meet and suppression rate is lower.

Figure 7.8: Impact of spatial resolution parameters on CANON – top: sub-MBR area α_{sub} and bottom: full-MBR area α_{full} with $\theta = 15°$. (a) Average number of requests served in a session. (b) Average size of requesting object's peer group. (c) Percentage of suppressed requests.

7.5.4 Summary

The following points summarize the results from the experimental study.

- CANON has a superior performance compared to *ProvidentHider* in maintaining longer service sessions across a wide range of anonymity requirements. More requests are also successfully anonymized by CANON.

- Including a small number of extra objects in a peer set is advantageous in handling defunct peers. However, extremely large peer sets can be detrimental.

- Use of direction information during the formation of a peer set does help avoid peers drifting away from each other over time. Choice of a too small neighborhood affects service quality, but is not necessary to balance performance across different measures.

- Performance is better with larger sub-MBR resolutions. However, performance in high precision services may be improved with a good strategy to relax the constraint.

- Service continuity is marginally different for different full-MBR resolutions. However, failure to serve new requests is much lower with smaller resolutions.

7.6 Preventing Query Disclosures

No single privacy model can yet prevent all underlying threats. Hence, the types of privacy attacks considered under an adversary model should also be explicitly mentioned. Earlier studies have identified two attack categories – *identity inferencing* (association of a user with the location(s) it has visited) and *query association* (inference of the sensitive attribute(s) involved in a user's request). The extent of success while executing attacks in these categories is decided by the adversary's background knowledge. In this study, we assume that the adversary's background knowledge is in terms of location information of one or more users. Note that the background knowledge can also be in terms of query parameters. Such forms of knowledge enable an adversary to eliminate possibilities in query association attacks.

Location-unaware adversaries do not posses knowledge of exact user locations. However, identity inferencing by such adversaries is possible when the revealed location data corresponds to a private address (restricted space identification) or can be associated to a user based on observed evidence (observation identification). If the location data is from a continuous LBS, then trajectories can also be linked to a user. Location k-anonymity prevents such inferencing by cloaking the exact location data inside a bounding box containing at least k users. Query association is the next step of identity inferencing where an adversary tries to determine what service attributes are interesting to an identified user. However, a k-anonymous cloaking region can implicitly reveal service attributes if all users in it specify the same (or similar) values. Query ℓ-diversity prevents such attacks by ensuring that a cloaking region contains users with at least ℓ distinct attribute values.

On the other hand, a location-aware adversary has exact location information on one or more users, and possibly at multiple time instances. User identities are therefore assumed to be known to the adversary. Hence, exact location data may be communicated to the LBS. However, location-aware adversaries cannot infer service attributes from the location knowledge as long as every location communicated to the LBS is query ℓ-diverse. In other words, a service request should involve a set of ℓ distinct attribute values (one of which is the real one) for every location update. Note that one cannot dismiss the absence of location-unaware adversaries in a given setting. Hence, cloaking regions are still used instead of exact locations.

A continuous LBS introduces other threats in the presence of location-aware adversaries. A continuous LBS allows an adversary to obtain the successive cloaking regions of a particular user. Given that both types of adversaries may be present, a LBS may adopt one of the following two methods to prevent query association.

- *Many-to-one queries:* In this method, a k-anonymous cloaking region communicated to the LBS is associated with a single service attribute (the one belonging to the actual user). Therefore, there are at least k potential users who may be the owner of the service attribute. However, if only one user is common across all the cloaking regions, then the attribute value must be associated with that user.

Figure 7.9: Schematic of the system architecture for query privacy.

- *Many-to-many queries:* In this method, a cloaking region is communicated to the LBS with a set of service attribute values (ones belonging to the users inside the region). Query association is prevented here by enforcing query ℓ-diversity in the set of attribute values. However, as highlighted in Section 7.1, query ℓ-diversity is not sufficient in a continuous LBS.

Our focus in this chapter is in the second strategy of prevention. We shall discuss the system architecture in accordance with this strategy and then show how query *m*-invariance eliminates the issues with query ℓ-diversity in a continuous LBS.

7.6.1 System architecture

The system architecture considered here is similar to that discussed in Section 7.3.1, consisting of three layers – (i) mobile users, (ii) a trusted anonymity server, and (iii) a continuous LBS provider (Fig. 7.9). A mobile user $\mathcal{U}$ initiates a service session by registering itself with the anonymity server. The registration process includes the exchange of current location information and service parameters. The service parameters collectively signify a service attribute value ($\mathcal{U}.\mathcal{S}$) for use with the LBS, as well as the anonymity level to enforce while generating the requests. The service attribute is considered sensitive information whose disclosure results in a privacy breach. The anonymity server generates a set of cloaking regions $A_1, \ldots, A_n$ and a set S of service attribute values for the requesting user. The user is present in one of these regions. Multiple range queries are then issued to the LBS provider for each of these regions, denoted as (A_i, S) in the figure. The LBS generates the results for each query such that a user anywhere in the cloaking region A_i with an interest in any of the values in S is served. A candidate result set is formed by

Figure 7.10: Example showing movement of users during a particular session registered to U. Values within the circles signify service attribute values of the users. Each cloaking region generated for U is 2-diverse and 3-anonymous.

merging all results from the multiple range queries. The anonymity server then filters the result set and communicates the accurate result to the mobile user. A request is suppressed (dropped) when the anonymity requirements cannot be met. The mobile user periodically updates its location with the anonymity server and receives updated results. The user unregisters and terminates the session when the service is no longer required. We assume that a user does not change its service attribute value during a session. A separate session is started if a request with different service parameters is to be made.

7.6.2 Query associations in a continuous LBS

The area of a cloaking region depends on the size of the anonymity set, as well as the time instance. Given a cloaking region R at time t, we shall use the notation $Users(R,t)$ to signify the set of users inside R at time instance t. Our system architecture uses multiple cloaking regions $A_1,\dots,A_n$ while serving a single request. The requirement for this is discussed later. In the following discussion, the cloaking region of a user is the MBR of the set of users that appear in at least one A_i.

Definition 7.2 SESSION PROFILE. Let $R_1,\dots,R_n$ be the cloaking regions of a user $\mathcal{U}$ at time instances $t_1,\dots t_n$ respectively during a particular session, where $t_i > t_j$ for $i > j$. Let $S_1,\dots,S_n$ be the set of service attribute values of users in the successive anonymity sets of $\mathcal{U}$ at different time instances, i.e. $S_i = \{u.\mathcal{S}|u \in Users(R_i,t_i)\}$. The session profile of $\mathcal{U}$ is then the set $SP(\mathcal{U}) = \cup_{i=1}^{n}(\{t_i\} \times \{R_i\} \times S_i)$.

Table 7.2: Session profile $SP(U)$ and background knowledge $BK(U)$ used during a query association attack on U.

	t	$\mathcal{R}$	$\mathcal{S}$
1	t_1	R_1	a
2	t_1	R_1	b
3	t_1	R_1	c
4	t_2	R_2	a
5	t_2	R_2	b
6	t_3	R_3	a
7	t_3	R_3	b

(a) $SP(U)$

	t	x	y	u
1	t_1	5.1	2.3	Alice
2	t_1	6.4	1.8	Bob
3	t_2	5.8	3.6	Alice
4	t_2	6.9	3.5	Bob
5	t_3	5.9	5.8	Alice
6	t_3	9.2	5.5	Bob

(b) $BK(U)$

An entry in a session profile is therefore of the form $\langle t, \mathcal{R}, \mathcal{S} \rangle$. For an $e \in SP(\mathcal{U})$, we shall use $e.t$, $e.\mathcal{R}$ and $e.\mathcal{S}$ to denote the corresponding terms. We shall also refer to $\mathcal{U}$ as the owner of the session profile. Consider the movement of the users shown in Fig. 7.10. Let us assume that a session for U lasted for three time stamps t_1, t_2 and t_3, during which the 2-diverse cloaking regions R_1, R_2 and R_3 are generated. Note that users other than U may terminate their session while U's session is in progress. As a result, their service attribute value may change during U's session. Table 7.2a lists the session profile of U w.r.t. this session.

Ideally, no knowledge on the owner of the session is required to form a session profile. A continuous LBS can improve service quality if successive requests from the same user can be distinguished from others [18]. Hence, the anonymity server typically maintains some session identifier with the continuous LBS. All cloaking regions with the same identifier belong to the same user. This information, along with the request logs (time stamp and attribute values) accumulated at the LBS, is sufficient to build the session profile. The objective is to accurately associate a service attribute value to the owner of the profile. Note that, under a location-aware adversary model, identification of the owner of the profile implies a successful query association only when the anonymity server uses the many-to-one system of querying the LBS. In a many-to-many system, the adversary will still have to associate one of the many attribute values to the owner. Next, we formally state the background knowledge of the location-aware adversary that can be used to link the owner to its service attribute value.

Definition 7.3 BACKGROUND KNOWLEDGE. The background knowledge of an adversary is a set BK of tuples of the form $\langle t, x, y, u \rangle$ which implies that the user u is known to have been at the location (x, y) at time instance t.

We shall only consider a subset of the background knowledge possessed by an adversary. This subset corresponds to the information that is relevant to perform a query association along with the data in a session profile. Given a session profile $SP(\mathcal{U})$, the background knowledge corresponding to the session is given as $BK(\mathcal{U}) = \{b \in BK | i \in \{1, \ldots, n\}, b.u \in \bigcap_i Users(R_i, t_i), b.t = t_i\}$. We make the worst case assumption that the adversary is aware of the location of every user present in the cloaking regions of $\mathcal{U}$ at all time instances when the queries are issued. In other words, for every t_i when a query is issued, there exists $|\bigcap_i Users(R_i, t_i)|$ entries in $BK(\mathcal{U})$. These entries correspond to the users that are present in all the cloaking regions generated during a session, and can potentially be the owner of the session. Table 7.2b lists the background knowledge used for U. The cloaking regions contain two potential owners, complete location information on whom is listed in $BK(U)$. Background knowledge associates users to locations while a session profile associates locations to service attribute values. The adversary relates the location data in $BK(\mathcal{U})$ and $SP(\mathcal{U})$ to link users to attribute values. We call this a *query association attack.*

Definition 7.4 QUERY ASSOCIATION ATTACK. Given a session profile $SP(\mathcal{U})$ and the background knowledge $BK(\mathcal{U})$, a query association attack on user $\mathcal{U}$ is a mapping $f : BK(\mathcal{U}) \rightarrow SP(\mathcal{U})$ such that

1. every $b \in BK(\mathcal{U})$ is mapped to exactly one $e \in SP(\mathcal{U})$,

2. every $b \in BK(\mathcal{U})$ with $f(b) = e$ satisfies

   (a) $(b.x, b.y)$ is inside $e.\mathcal{R}$

   (b) $b.t = e.t$

   (c) for all $b' \in \{b^o \in BK(\mathcal{U}) | b^o.u = b.u\}$, $f(b').\mathcal{S} = e.\mathcal{S}$.

The first condition states that a user can be associated with only one attribute value in a given time instance. The second condition prohibits the adversary from arbitrarily

Table 7.3: Associating service attribute values using query association attacks. (a) Condition 2c in Def. 7.4 not met for Alice. (b) All possible query association attacks w.r.t. Table 7.2.

(a)

BK(U)	SP(U)
1: Alice	2: b
2: Bob	2: b
3: Alice	5: b
4: Bob	5: b
5: Alice	6: a
6: Bob	7: b

(b)

BK(U)	SP(U)	BK(U)	SP(U)	BK(U)	SP(U)
1: Alice	1: a	1: Alice	2: b	1: Alice	2: b
2: Bob	1: a	2: Bob	2: b	2: Bob	1: a
3: Alice	4: a	3: Alice	5: b	3: Alice	5: b
4: Bob	4: a	4: Bob	5: b	4: Bob	4: a
5: Alice	6: a	5: Alice	7: b	5: Alice	7: b
6: Bob	6: a	6: Bob	7: b	6: Bob	6: a

mapping tuples between $BK(\mathcal{U})$ and $SP(\mathcal{U})$. Conditions 2a and 2b state that a user must be inside the cloaking region (and at the specific time instance) corresponding to the entry to which it is mapped to. Condition 2c requires that a user be associated with a single attribute value across all time instances. This condition forms the basis for a successful attack since it is known that the owner of the session profile will always have the same service attribute value within the session. Consider the mapping between $BK(U)$ and $SP(U)$ shown in Table 7.3a. This mapping associates Alice with the value 'b' at time t_1 $(1 \rightarrow 2)$ and t_2 $(3 \rightarrow 5)$, but with 'a' at time t_3 $(5 \rightarrow 6)$. If Alice is the owner of the profile, then she must be associated with the same value at all time instances. In other words, the mapping fails to satisfy condition 2c and is not considered a possible query association attack. The mappings shown in Table 7.3b are the only possible query association attacks in this case. Privacy is then measured as the probability that a query association attack accurately associates a user with its service attribute value.

Definition 7.5 DISCLOSURE RISK. Given a session profile $SP(\mathcal{U})$, let $QAA(\mathcal{U})$ be the set of all possible query association attacks on user $\mathcal{U}$. Consider the subset $QAA_b(\mathcal{U})$ of query association attacks that accurately identifies the service attribute value of $\mathcal{U}$, i.e. given $b \in BK(\mathcal{U})$ with $b.u = \mathcal{U}$, $QAA_b(\mathcal{U}) = \{f \in QAA(\mathcal{U}) | \forall b, f(b).\mathcal{S} = \mathcal{U}.\mathcal{S}\}$. The disclosure risk for $\mathcal{U}$ is the fraction of query association attacks on $\mathcal{U}$ that accurately maps it with its service attribute value, given as $DR(\mathcal{U}) = \frac{|QAA_b(\mathcal{U})|}{|QAA(\mathcal{U}|}$.

With reference to Table 7.3b, we have $|QAA(U)| = 4$, out of which two mappings accurately associate U (i.e. Alice) with the service attribute value used by her during the session (i.e. 'a'). Therefore, $|QAA_b(U)| = 2$ and disclosure risk of Alice is 0.5.

Theorem 7.1 Let $R_1, \ldots, R_n$ be the cloaking regions of a user $\mathcal{U}$ at time instances $t_1, \ldots t_n$ respectively during a particular session, where $t_i > t_j$ for $i > j$. Let $S_1, \ldots, S_n$ be the set of service attribute values of users in the successive anonymity sets of $\mathcal{U}$ at different time instances, i.e. $S_i = \{u.\mathcal{S} | u \in Users(R_i, t_i)\}$. The disclosure risk of $\mathcal{U}$ is 1.0 if $|\bigcap_i S_i| = 1$.

Proof Let $f : BK(\mathcal{U}) \rightarrow SP(\mathcal{U})$ be any query association attack. Consider a tuple $b \in BK(\mathcal{U})$ such that $b.u = \mathcal{U}$ and let $f(b) = e$. Hence, for any tuple $b' \in BK(\mathcal{U})$ with $b'.u = \mathcal{U}$,

we have $f(b').\mathcal{S} = e.\mathcal{S}$ (from Def. 7.4, condition 2c). Note that $e.\mathcal{S}$ is the service attribute value that the adversary has associated with $\mathcal{U}$ under the attack f. We show that $e.\mathcal{S}$ is in fact $\mathcal{U}.\mathcal{S}$ for any f, and hence all possible query association attacks accurately associate $\mathcal{U}$ with its service attribute value, i.e. $QAA(\mathcal{U}) = QAA_b(\mathcal{U}) \implies DR(\mathcal{U}) = 1.0$.

By definition of $BK(\mathcal{U})$, every b' has a different time stamp ($b'.t$) and is therefore mapped to a different $f(b')$. Further, only one cloaking region is associated with a time stamp in a session profile. Hence, every $f(b')$ has a different cloaking region depending on the time stamp. Since there is a b' for $t_1,\ldots,t_n$, there is a $f(b')$ for every $t_1,\ldots,t_n$. S_i is the set of service attribute values associated to users in the cloaking region at time t_i. Therefore, any S_i includes the value $f(b').\mathcal{S} = e.\mathcal{S}$, implying $e.\mathcal{S} \in \cap_i S_i$. Also, $e.\mathcal{S}$ must be the only element in $\cap_i S_i$ as the size of this set is given to be one. Given that $\mathcal{U}$ belongs to all cloaking regions in the session profile and $\mathcal{U}.\mathcal{S}$ is the parameter with which it issues its query, $\mathcal{U}.\mathcal{S}$ must also be in $\cap_i S_i$. This gives us $e.\mathcal{S} = \mathcal{U}.\mathcal{S}$. $\square$

7.6.3 Query m-invariance

Theorem 7.1 underlines why location k-anonymity and query ℓ-diversity are not sufficient to prevent query association attacks. Location k-anonymity only guarantees that the number of users in every cloaking region is at least k. However, the same users may not be present across all the cloaking regions, thereby requiring a much smaller $BK(\mathcal{U})$. Historical k-anonymity guarantees that background knowledge must be available on at least k users. Nonetheless, query association attacks can still reveal the service attribute value if only one such value is consistently present across all queries. Query ℓ-diversity guarantees that there are at least ℓ distinct values in every query, but does not try to invariably maintain the same set of values across queries. The requirement for such an invariant property motivates us to consider the principle of query m-invariance.

Definition 7.6 QUERY m-INVARIANCE. Let $R_1,\ldots,R_n$ be the cloaking regions of a user $\mathcal{U}$ at time instances $t_1,\ldots t_n$ respectively during a particular session, where $t_i > t_j$ for $i > j$. A cloaking region R_j is query m-invariant if $|\cap_{i=1}^{j} S_i| \geq m$ where $S_i = \{u.\mathcal{S} | u \in Users(R_i, t_i)\}$.

Query *m*-invariance implicitly implies location *m*-anonymity and query *m*-diversity. The principle draws upon the observation that the number of possible query association attacks will increase if a user can be associated with more number of service attribute values. However, this would require multiple values to be present at all time stamps in the session profile. Query *m*-invariance guarantees that the number of such values is not less than *m*. With reference to Fig. 7.10, there are two values ('*a*' and '*b*') that are invariably present across all cloaking regions. The disclosure risk in this case is $\frac{1}{2}$. In general, the following theorem provides the upper bound on the disclosure risk for query *m*-invariant cloaking regions.

Theorem 7.2 Let $R_1, \ldots, R_n$ be query *m*-invariant cloaking regions of a user $\mathcal{U}$ at time instances $t_1, \ldots t_n$ respectively during a particular session, where $t_i > t_j$ for $i > j$. The disclosure risk of $\mathcal{U}$ is then at most $\frac{1}{m}$.

Proof Since $\mathcal{U}$ is present inside every $R_i; 1 \le i \le n$, $BK(\mathcal{U})$ contains a tuple for $\mathcal{U}$ at each time instance $t_1, \ldots, t_n$. Let $b_1, \ldots, b_n$ denote these tuples, i.e. $b_i.u = \mathcal{U}$ and $b_i.t = t_i$, for $1 \le i \le n$. Given a query association attack $f : BK(\mathcal{U}) \to SP(\mathcal{U})$, we have $f(b_1).\mathcal{S} = \ldots = f(b_n).\mathcal{S}$ by Def. 7.4, condition 2c. Consider an arbitrary b_k. Note that if f maps b_k such that $f(b_k).\mathcal{S} = \mathcal{U}.\mathcal{S}$, then every $b_i; i \ne k$ will also be mapped such that $f(b_i).\mathcal{S} = \mathcal{U}.\mathcal{S}$. By definition, such a query association attack then belongs to $QAA_b(\mathcal{U})$. Hence, we need a count of the number of attacks that map an arbitrarily chosen b_k in $\{b_1, \ldots, b_n\}$ to a session entry such that $f(b_k).\mathcal{S} = \mathcal{U}.\mathcal{S}$.

Since $b_k.u$ must be associated with the same service attribute value at all time stamps, it must be one that is present in all S_i, for $1 \le i \le n$. Let $p = |\bigcap_i S_i|$. Further, let q be the number of users in $\bigcap_i Users(R_i, t_i)$. The same users are present in $BK(\mathcal{U})$ at time stamp t_k. The function f associates the q users with one of the p service attribute values. This can be done in p^q ways, out of which p^{q-1} is the number of ways where a particular user is fixed to a specific value. Hence $\frac{p^{q-1}}{p^q} = \frac{1}{p}$ is the fraction of attacks that associate $b_k.u$ (or $\mathcal{U}$) with a particular value in $\bigcap_i S_i$ (which can be $\mathcal{U}.\mathcal{S}$ since it belongs to all S_i). Since all cloaking regions are query *m*-invariant, we have $p \ge m$, implying that the fraction is at most $\frac{1}{m}$. $\square$

Note that, by symmetry, any user in $\bigcap_i Users(R_i, t_i)$ has a disclosure risk of at most $\frac{1}{m}$. Further, the number of users in $\bigcap_i Users(R_i, t_i)$ does not affect the disclosure risk as far as query association attacks are concerned. There is no restriction on the size of common users set (as in historical k-anonymity) since, under the location-aware adversary model, user identities are already assumed to be known. As far as location aware-adversaries are concerned, it is sufficient to have k-anonymous cloaking regions.

7.7 A Cloaking Algorithm

A trivial implementation of query m-invariance is to randomly decide m distinct service attribute values (one of them must be the user's attribute value) and use it as the *invariant set* of values (we call it S in Fig. 7.9) across all cloaking regions in the session. However, this implementation is vulnerable to other inference attacks. Consider the user U who consistently uses the service attribute value 'a'. Hence, the set S in all sessions belonging to U will have 'a'. Given that other values in S will be generated randomly, an adversary can observe that the value 'a' is present with a high frequency in the set S across all sessions whenever user U is in the common users set. This allows the adversary make a highly confident association between U and 'a'. The method to prevent such inference attacks is to preserve reciprocity in the set S, i.e. the set S should be the same no matter which user in $\bigcap_i Users(R_i, t_i)$ is the owner of the session. Ideally, such a set is $\{u.S | u \in \bigcap_i Users(R_i, t_i)\}$. However, forming this set is not possible without clairvoyant knowledge about the users that will be present in every cloaking region generated in the session.

Our approach considers a m-diverse set of users in the first time stamp t_1 and uses their service attribute values as the set S. In successive instances, the cloaking region is adjusted so that each value in S is the service attribute value of at least one user inside the region. All cloaking regions then have users with service attribute values in S, thereby making $\bigcap_i S_i$ equal to S. Since the first cloaking region is m-diverse, S has at least m elements and every cloaking region is query m-invariant. Further, reciprocity in the user set is preserved by using Hilbert Cloak to determine the anonymity sets. Procedure 7.4 outlines this approach.

Procedure 7.4 m-InvariantCloak(User $\mathcal{U}$)

Input: Mobile user $\mathcal{U}$.

Output: A set of peer groups (one of them includes $\mathcal{U}$).

1: $\mathcal{L}$ = set of available mobile users sorted by their Hilbert index
2: $\mathcal{D}_{prev} = \phi; \mathcal{D} = \phi$
3: **if** ($\mathcal{U}.invSet = \phi$) **then**
4:    $\mathcal{D} = m\text{-}DiverseCloak(\mathcal{L},\mathcal{U})$
5:    $params = \mathcal{U}.invSet = \{u.\mathcal{S}|u \in \mathcal{D}\}$
6: **else**
7:    **repeat**
8:       $\mathcal{D}_{prev} = \mathcal{D}; \mathcal{D} = \phi; params = \phi$
9:       **for all** ($l \in \mathcal{L}$ in order) **do**
10:          $\mathcal{D} = \mathcal{D} \cup \{l\}$
11:          $params = params \cup \{l.\mathcal{S}\}$
12:          **if** ($|params \cap \mathcal{U}.invSet| = \mathcal{U}.m$) **then**
13:             **break**
14:          **end if**
15:       **end for**
16:       $\mathcal{L} = \mathcal{L} - \mathcal{D}$
17:    **until** ($\mathcal{U} \in \mathcal{D}$)
18: **end if**
19: **if** ($|params \cap \mathcal{U}.invSet| < \mathcal{U}.m$) **then**
20:    **if** ($\mathcal{D}_{prev} = \phi$) **then**
21:       **return** null
22:    **else**
23:       $\mathcal{D} = \mathcal{D}_{prev} \cup \mathcal{D}$
24:    **end if**
25: **end if**
26: $\mathcal{U}.invSet = \mathcal{U}.invSet \cap \{u.\mathcal{S}|u \in \mathcal{D}\}$
27: **return** $PartitionSet(\mathcal{D})$

---

m-InvariantCloak starts with a list $\mathcal{L}$ of all registered users sorted by their Hilbert index. For every registered user $\mathcal{U}$, it maintains: (i) a set of service attribute values ($\mathcal{U}.invSet$) that has invariably been present in every cloaking region generated for $\mathcal{U}$ in the current session, (ii) the anonymity requirement ($\mathcal{U}.m$) for $\mathcal{U}$ and (iii) the service attribute value ($\mathcal{U}.\mathcal{S}$) used by $\mathcal{U}$ in the current session. The set of users in the first cloaking region is generated by *m-DiverseCloak* (Lines 3-5). This function returns a *m*-diverse set of users using the Hilbert Cloak algorithm. Hilbert Cloak partitions the set of users into buckets such that each bucket is *m*-diverse. The algorithm returns the set of users in the bucket that contains $\mathcal{U}$. The invariant set for $\mathcal{U}$ is the set of service attribute values of the returned users (Line 5). For subsequent cloaking requests in the same session, the buckets are

formed such that each contains $\mathcal{U}.m$ service attribute values from $\mathcal{U}.invSet$ (Lines 7-17). New buckets are formed until the one with $\mathcal{U}$ is found. Note that if $\mathcal{U}$ is in the last bucket, then there may be less than $\mathcal{U}.m$ distinct attribute values in it. In such a case, $\mathcal{U}$'s bucket is merged with the previous one (Line 23) if it exists; otherwise the request must be suppressed (Line 21). Request suppression is not likely as long as $\mathcal{U}.m$ is not higher than the number of possible service attribute values. Once the bucket of $\mathcal{U}$ is decided, its invariant set is updated. The update is required because the first invariant set may contain more than $\mathcal{U}.m$ diverse values out of which only $\mathcal{U}.m$ fixed values are to be retained from the second instance onwards. The merging of buckets is the only reason why an invariant set may have more than $\mathcal{U}.m$ elements. The following theorem summarizes the effect of the pseudo-code.

Theorem 7.3 Let R be the minimum bounding rectangle of the users in $\mathcal{D}$, where $\mathcal{D}$ is generated by m-InvariantCloak($\mathcal{U}$) at the end of Line 25 in Procedure 7.4. Then, R preserves query m-invariance with $m = \mathcal{U}.m$.

Users in the set $\mathcal{D}$ at the end of Line 25 can be used to issue point queries along with the service attribute set $\mathcal{U}.invSet$. This will be a case of one-to-many queries. As mentioned earlier, we discourage such queries because of the possible presence of location-unaware adversaries. *m-InvariantCloak* therefore partitions $\mathcal{D}$ into peer groups using *PartitionSet*. Similar to the partitioning approach in the CANON algorithm, the objective of *PartitionSet* is to partition a *m*-invariant user set into peer groups. Every group then defines its own minimum bounding rectangle (called the sub-MBR) over which a range query is issued. Results to such a query are formed such that any user anywhere inside the rectangle is served. Procedure 7.5 restates the partitioning approach, with the difference that each bucket now must include at least 2 users. If the last group has less than 2 users then it is merged with the group formed prior to it (Lines 12-14). The partitioning can also be performed so that every group has a fixed number of users. We avoid this approach since user densities vary across time and space, as a result of which, the area of cloaking regions may be beyond acceptable levels. Note that the invariant set of service attribute values is not changed by the partitioning scheme. In fact, the same set is used for

Procedure 7.5 PartitionSet(Set $\mathcal{L}$)

Input: A set $\mathcal{L}$ of users and system global α_{sub}.
Output: A set of peer groups.
1: Sort objects in $\mathcal{L}$ by their Hilbert index
2: $peerGroups = \phi$
3: $bucket = \phi$
4: **for all** ($l \in \mathcal{L}$ in order) **do**
5:    **if** ($AreaMBR(bucket \cup \{l\}) \leq \alpha_{sub}$ **or** $|bucket| < 2$) **then**
6:       $bucket = bucket \cup \{l\}$
7:    **else**
8:       $peerGroups = peerGroups \cup \{bucket\}$
9:       $bucket = \{l\}$
10:   **end if**
11: **end for**
12: **if** ($|bucket| < 2$) **then**
13:    Remove last bucket entered into *peerGroups* and merge it with *bucket*
14: **end if**
15: $peerGroups = peerGroups \cup \{bucket\}$
16: **return** *peerGroups*

---

the range queries corresponding to each cloaking region. Hence the following theorem holds.

Theorem 7.4 Let $G_1, \ldots, G_n$ be the peer groups returned by m-InvariantCloak for a mobile user $\mathcal{U}$. With reference to the system architecture in Section 7.6.1, we define $S = \{g.\mathcal{S} | g \in \underset{i}{\cup} G_i\}$ and $A_i =$ the minimum bounding rectangle of users in G_i. The anonymity server then preserves query m-invariance for $\mathcal{U}$.

7.8 Empirical Study

This empirical study compares the effectiveness of location k-anonymity, query ℓ-diversity and query m-invariance in limiting the privacy risks in a continuous LBS. Hilbert Cloak is used to create the location k-anonymous and query ℓ-diverse cloaking regions, while *m-InvariantCloak* is used for query m-invariance. The cloaking region returned by Hilbert Cloak for k-anonymity and ℓ-diversity is partitioned similar to as in Procedure 7.5. The following statistics are used to evaluate the performance.

- *safeguard against query disclosures:* number of vulnerable sessions extracted using Theorem 7.1.

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- *service quality:* area of a sub-MBR.

- *safeguard against location-unaware adversaries:* number of users inside a sub-MBR.

- *anonymization time:* time required to compute a privacy preserving cloaking region.

7.8.1   Experimental setup

Trace data for this experiment is generated using the same simulator as described in Section 7.5.1. A record in this case is made up of a time stamp, user identifier, and x and y co-ordinates of the user's location. Duration of a session for a user is determined from a normal distribution with mean 10 minutes and standard deviation 5 minutes. A new duration is assigned at the end of a session. Each user has an associated $k/\ell/m$ value drawn from the range [2,50] by using a Zipf distribution favoring higher values. Service attribute values are assigned from a set of 100 values using another Zipf distribution. Both distributions have an exponent of 0.6. The trace data is sorted by the time stamp of records.

The session duration time is used to determine if a request is a new one or a continuing one. The anonymizer is then called to determine the cloaking region(s), if possible. The process continues until the session ends; a new session is started when the user issues the next request. A new service attribute value is assigned to a user at the beginning of every session. The anonymizer receives over 4,000,000 anonymization requests during a pass of the entire trace data. The default spatial resolution is set to $\alpha_{sub} = 0.0625\,km^2$. The precision is around $250\,m$ (assuming a square area) with this setting.

7.8.2   Simulation results

Fig. 7.11 compares the effectiveness of location k-anonymity, query ℓ-diversity and query m-invariance. Query m-invariance is most effective in preventing query association attacks (Fig. 7.11a). Query ℓ-diversity can prevent query disclosures in more number of sessions compared to location k-anonymity. This is anticipated since ℓ-diverse cloaking regions are required to have at least ℓ distinct service attribute values. The invariance property in query m-invariance further prevents the possibility of only one attribute value

Figure 7.11: Comparative performance of location *k*-anonymity, query ℓ-diversity and query *m*-invariance.

being common across the cloaking regions. Both k-anonymity and ℓ-diversity have almost 100% vulnerability for weaker anonymity requirements. In general, the fewer the number of service attribute values in the first cloaking region, the higher are the chances of not having any of those values in subsequently generated regions. However, by definition, such chances are reduced to zero in the query m-invariance model.

We also observe that query ℓ-diversity manages to reduce the number of vulnerable sessions to impressive lows for cases with higher anonymity requirements. This is not unlikely since the probability of a particular value being in a subset of all possible attribute values is higher when larger subsets are to be formed. However, the performance is not indicative of similar behavior. Query m-invariance guarantees that the invariant set has a size of at least m. On the other hand, the low percentage of vulnerable sessions with query ℓ-diversity only means that the invariant set is not of size one. The actual size of the set can very well be less than ℓ. Hence, the disclosure risk is not guaranteed to be less than $\frac{1}{\ell}$, resulting in partial disclosures.

Sub-MBR areas are typically smaller with k-anonymity and ℓ-diversity (Fig. 7.11b). Query m-invariance generates comparatively larger areas for weaker anonymity requirements. This is because the users that satisfy the invariant set requirement may often be far away from each other. This is specifically true if the invariant set has values that are not frequently requested. Nevertheless, the sub-MBR area is within the spatial resolution, implying that peer groups could be formed without violating the spatial constraint (the 'or' condition in Line 5 of Procedure 7.5). Further, the service quality is consistent across all anonymity requirements. Fig. 7.11c illustrates the average time required to anonymize a request. The query m-invariance requirement does not impose any significant overhead in terms of computation time.

Fig. 7.12a depicts the impact of α_{sub} on service quality. A very small spatial resolution (such as $\alpha_{sub} = 0.0025\,km^2$) is difficult to satisfy irrespective of the anonymity level required. Given that each peer group must contain at least 2 users, the spatial constraint is easily violated and the sub-MBR area is consequently larger than specified. The area is large enough to accommodate 2 users. Fig. 7.12b corroborates this observation since the average number of users inside a sub-MBR is 2 for such an α_{sub}. A cloaking region

Figure 7.12: Impact of spatial resolution α_{sub}.

of $0.0025\,km^2$ resolves to a precision of around $50\,m$, often not required in a LBS. The resolution has a direct impact on the number of users in a peer group. Larger resolutions allow more users to be included in a group, thereby providing stronger protection from location-unaware adversaries. Note that a 2-invariant cloaking region can potentially contain a large number of users. This is specifically true when most users are inclined towards specific service attribute values. As a result, a cloaking region that contains 2 distinct attribute values essentially contains multiple users having the same value. A peer group with 45 users on the average (in the case of $\alpha_{sub} = 1.0\,km^2$) is therefore not surprising for a weak requirement such as 2-invariance. A reasonable α_{sub} such as the default value sufficiently maintains a good balance between service quality and identity disclosure risks.

7.9   Conclusions

Identity and query privacy must be adequately guaranteed before location-based services can be deployed on a large scale. Models such as *k*-anonymity provide customizable privacy guarantees in a snapshot LBS by generating a cloaking region. However, a

k-anonymous cloaking region is not sufficient to guarantee privacy in a continuous LBS. An extended notion called historical *k*-anonymity has been proposed for such services. However, all known methods of enforcing historical *k*-anonymity significantly affects the quality of service that the service provider can deliver. In this chapter, we have identified the factors that contribute towards deteriorated service quality and suggested resolutions. We propose the CANON algorithm that delivers reasonably good service quality across different anonymity requirements. The algorithm uses tunable parameters to adjust the size of a peer set, trajectories of peers and cloaking regions over which range queries are issued. Immediate future work includes optimizing the performance of CANON in terms of better usage of directional information. We believe this optimization is crucial in order to have similar performance across all levels of anonymity requirements.

Association of service parameters to users cannot be prevented in a continuous LBS by simply assuring that a cloaking region contains k or more users, or ℓ distinct service attribute values. Assuming a location-aware adversary model, we have provided a formal analysis to show that service attributes risk disclosure if the privacy model does not guarantee that an invariant set of attribute values is present in all cloaking regions generated for the continuous LBS. We therefore propose using the principle of query *m*-invariance where all cloaking regions are required to contain users with a fixed set of service attribute values. We have shown that this requirement limits the involved risk, which in itself can be controlled by the parameter *m*. We further propose a cloaking algorithm to enforce the principle and have shown its effectiveness compared to location *k*-anonymity and query ℓ-diversity. Results on trace data generated on a real-world road network show that query *m*-invariance can be enforced without significantly affecting service quality or imposing computational overhead. Further, the approach can be used to safeguard against both location-aware and location-unaware adversaries. Future work can be directed towards understanding the risks from query association attacks under alternative forms of background knowledge. Specifically, we are interested in exploring the privacy guarantees required to tackle adversaries with limited knowledge on user locations.

Part II

Security Risk Management

CHAPTER 8

Security Hardening on Attack Tree Models

Network-based computer systems form an integral part of any information technology infrastructure today. The different levels of connectivity between these systems directly facilitate the circulation of information within an organization, thereby reducing invaluable wait time and increasing the overall throughput. As an organization's operational capacity becomes more and more dependent on networked computing systems, the need to maintain accessibility to the resources associated with such systems has become a necessity. Any weakness or vulnerability that could result in the breakdown of the network has direct consequences on the amount of yield manageable by the organization. This, in turn, requires the organization to not only consider the advantages of utilizing a networked system, but also consider the costs associated with managing the system.

With cost-effectiveness occurring as a major factor in deciding the extent to which an organization would secure its network, it is not sufficient to detect the presence or absence of a vulnerability and implement a security measure to rectify it. Further analysis is required to understand the contribution of the vulnerabilities towards any possible damage to the organization's assets. Often, vulnerabilities are not exploited in isolation, but rather used in groups to compromise a system. Similarly, security policies can have a coverage for multiple vulnerabilities. Thus, cost-effective security management requires researchers to evaluate the different scenarios that can lead to the damage of a secured

asset, and then come up with an optimal set of security policies to defend such assets.

Researchers have proposed building security models for networked systems using paradigms like *attack graphs* [10, 94, 141, 156, 166] and *attack trees* [43, 131, 145, 155], and then finding attack paths in these models to determine scenarios that could lead to damage. However, determining possible attack paths, although useful, does not help the system administrators much. They are more interested in determining the best possible way of defending their network in terms of an enumerated set of hardening options [136]. Moreover, the system administrator has to work within a given set of budget constraints which may preclude her from implementing all possible hardening measures or even measures that cover all the weak spots. Thus, the system administrator needs to find a trade-off between the cost of implementing a subset of security hardening measures and the damage that can potentially happen to the system if certain weak spots are left unpatched. In addition, the system administrator may also want to determine optimal robust solutions. These are sets of security hardening measures with the property that even if some of the measures within a set fail, the system is still not compromised.

We believe that the problem should be addressed in a more systematic manner, utilizing the different tools of optimization at hand. A decision maker would possibly make a better choice by successively exploring the different levels of optimization possible, rather than accepting a solution from an "off-the-shelf" optimizer. Towards this end, this chapter discusses our four major contributions. First, we refine and formalize the notion of attack trees so as to encode the contribution of different security conditions leading to system compromise. Next, we develop a model to quantify the potential damage that can occur in a system from the attacks modeled by the system attack tree. We also quantify the security control cost incurred to implement a set of security hardening measures. Third, we model the system administrator's decision problem as three successively refined optimization problems on the attack tree model of the system. We progressively transform one problem into the next to cater to more cost-benefit information as may be required by the decision maker. Last but not least, we discuss our thoughts and observations regarding the solutions, in particular the robust solutions identified by our optimization process, with a belief that such discussion will help the system administrator decide what

methodology to adopt.

The rest of the chapter is organized as follows. We discuss some of the previous works related to determining optimum security hardening measures in Section 8.1. In Section 8.2, we describe a simple network that we use to illustrate our problem formulation and solution. The attack tree model formalism and the cost model are presented in Sections 8.3 and 8.4 respectively. The three optimization problems are presented in Section 8.5 with some empirical results and discussion following in Section 8.6. Finally, we conclude in Section 8.7.

8.1   Related Work

Network vulnerability management has been previously addressed in a variety of ways. Noel et al. use a data structure called exploit dependency graphs [136] to compute minimum cost-hardening measures. Given a set of initial conditions in the graph, they compute Boolean assignments to these conditions, enforced by some hardening measure, so as to minimize the total cost of those measures. As pointed out in their work, these initial conditions are the only type of network security conditions under our strict control. Hardening measures applied to internal nodes can potentially be bypassed by an attacker by adopting a different attack path. Jha et al. [94] on the other hand do not consider any cost for the hardening measures. Rather, their approach involves finding the minimal set of atomic attacks critical for reaching the goal and then finding the minimal set of security measures that cover the minimal set of atomic attacks.

Such analysis is meant for providing solutions that guarantee complete network safety. However, the hardening measures provided may still not be feasible within the financial or other business constraints of an organization. Under such circumstances, a decision maker must perform a cost-benefit analysis to understand the trade-off between hardening costs and network safety. Furthermore, a minimum cost hardening measure set only means that the root goal is safe, and some residual damage may still remain in the network. Owing to these real-world concerns, network vulnerability management should not always be considered as a single-objective optimization problem.

A multi-objective formulation of the problem is presented by Gupta et al. [79]. They

Figure 8.1: Example network model.

consider a generic set of security policies capable of covering one or more generic vulnerabilities. A security policy can also introduce possible vulnerabilities, thereby resulting in some residual vulnerabilities even after the application of security policies. The multi-objective problem then is to minimize the cost of implementing the security policies, as well as the weighted residual vulnerabilities. However, the authors finally scalarize the two objectives into a single objective using relative weights for the objectives.

8.2   A Simple Network Model

To illustrate our methodology, we consider the hypothetical network as shown in Fig. 8.1. The setup consists of four hosts. A firewall is installed with a preset policy to ensure that only the *FTP* and *SMTP servers* are allowed to connect to the external network. In addition, FTP and SSH are the only two services an external user can use to communicate with these servers. We assume that an external user wants to compromise the *Data Server* which is located inside the firewall. The firewall has a strong set of policies setup to protect access to the internal hosts. There are six different attack scenarios possible to achieve the ultimate goal from a given set of initial vulnerabilities and network topology as listed in Table 8.1 and 8.2. To compromise the Data Server, an attacker can exploit the FTP and SMTP Servers. Both servers are running ftp server versions that are vulnerable to the *ftp/.rhost attack*. In addition, their *rhost directories* are not properly write-protected.

Table 8.1: Initial vulnerability per host in example network.

Host	Vulnerability	CVE#
FTP Server	Ftp .rhost attack	1999-0547
196.216.0.10	Ftp Buffer overflow	2001-0755
	Ssh Buffer overflow	2006-2421
SMTP Server	Ftp .rhost attack	1999-0547
196.216.0.1		
Terminal	LICQ remote-2-user	2001-0439
196.216.0.3	"at" heap corruption	2002-0004
Data Server	LICQ remote-2-user	2001-0439
196.216.0.2	suid Buffer overflow	2001-1180

Table 8.2: Connectivity in example network.

Host	Host	Port
..*.*	196.216.0.1	21,25
..*.*	196.216.0.10	21,22
196.216.0.1	196.216.0.2	ANY
196.216.0.1	196.216.0.3	ANY
196.216.0.3	196.216.0.2	ANY
196.216.0.10	196.216.0.2	ANY

As a consequence of the ftp/.rhost exploit, the attacker machine is able to establish a trust relation with the host machine, and introduces an *authentication bypassing vulnerability* in the victim. An attacker can then log in to these servers with user access privilege. From this point, the attacker can use the connection to the Data Server to compromise it. The attacker may also compromise the SMTP Server, or choose to compromise the *Terminal* machine in order to delay an attack. The Terminal machine can be compromised via the chain of *LICQ remote to user attack* and the *local buffer overflow attack on the "at" daemon*. Finally, the attacker from either the FTP server, SMTP server, or the Terminal machine can use the connectivity to the Data Server to compromise it through the chain of *LICQ exploit* and *"suid" local buffer overflow attack*. Such attack scenarios, as in our example network model, are represented using an attack tree, discussed in details in the next section.

8.3   Attack Tree Model

Given the complexity of today's network infrastructure, materializing a threat usually requires the combination of multiple attacks using different vulnerabilities. Representing

different scenarios under which an asset can be damaged thus becomes important for preventive analysis. Such representations not only provide a picture of the possible ways to compromise a system, but can also help determine a minimal set of preventive actions. Given the normal operational state of a network, including the vulnerabilities present, an attack can possibly open up avenues to launch another attack, thereby taking the attacker a step closer to its goal. A certain state of the network in terms of access privileges or machine connectivity can be a prerequisite to be able to exploit a vulnerability. Once the vulnerability is exploited, the state of the network can change enabling the attacker to launch the next attack in the sequence. Such a pre-thought sequence of attacks gives rise to an *attack scenario*.

It is worth noting that such a notion of a progressive attack induces a transitive relationship between the vulnerabilities present in the network and can be exploited while deciding on the security measures. Attack graph [10, 94, 136, 156] and attack tree [145, 155] representations have been proposed in network vulnerability management to demonstrate such cause-consequence relationships. The nodes in these data structures usually represent a certain network state of interest to an attacker, with edges connecting them to indicate the cause-consequence relationship. Although different attack scenarios are easily perceived in attack graphs, they can potentially suffer from a state space explosion problem. Ammann et al. [10] identified this problem and propose an alternative formulation, with the assumption of monotonicity. The monotonicity property states that the consequence of an attack is always preserved once achieved. Such an assumption can greatly reduce the number of nodes in the attack graph, although at the expense of further analysis required to determine the viable attack scenarios. An *exploit-dependency graph* can be extracted from their representation to indicate the various conjunctive and disjunctive relationships between different nodes. For the purpose of this study, we adopt the attack tree representation since it presents a much clearer picture of the different hierarchies present between attacker sub-goals. An attack tree uses explicit conjunctive and disjunctive branch decomposition to reduce the visualization complexity of a sequence of operations. The representation also helps us efficiently calculate the cost factors that are of interest to us.

Different properties of the network effectuate different ways for an attacker to compromise a system. We first define an *attribute-template* that lets us generically categorize these network properties for further analysis.

Definition 8.1 ATTRIBUTE-TEMPLATE. An attribute-template is a generic property of the hardware or software configuration of a network which includes, but is not limited to, the following.

- system vulnerabilities (which are often reported in vulnerability databases such as BugTraq, CERT/CC, or NetCat).

- network configuration such as open port, unsafe firewall configuration, etc.

- system configuration such as data accessibility, unsafe default configuration, or read-write permission in file structures.

- access privilege such as user account, guest account, or root account.

- connectivity.

An attribute-template lets us categorize most of the atomic properties of the network that might be of some use to an attacker. For example, *"running SSH1 v1.2.23 on FTP Server"* can be considered as an instance of the system vulnerabilities template. Similarly, *"user access on Terminal"* is an instance of the access privilege template. Such templates also let us specify the properties in propositional logic. We define an *attribute* with such a concept in mind.

Definition 8.2 ATTRIBUTE. An attribute is a propositional instance of an attribute-template. It can take either a *true* or *false* value.

The success or failure of an attacker reaching its goal depends mostly on what truth values the attributes in a network take. Its also lays the foundations for a security manager to analyze the effects of falsifying some of the attributes using some security policies. We formally define an attack tree model based on such attributes. Since we consider an attribute as an atomic property of a network, taking either a *true* or *false* value, most of the definitions are written in propositional logic involving these attributes.

Definition 8.3 ATTACK. Let S be a set of attributes. We define Att to be a mapping $Att : S \times S \rightarrow \{true, false\}$ and $Att(s_c, s_p) =$ truth value of s_p.

$a = Att(s_c, s_p)$ is an attack if $s_c \neq s_p \wedge a \equiv s_c \leftrightarrow s_p$. s_c and s_p are then respectively called a precondition and postcondition of the attack, denoted by $pre(a)$ and $post(a)$ respectively.

$Att(s_c, s_p)$ is a ϕ–attack if $\exists$non-empty $S' \subset S | [s_c \neq s_p \wedge Att(s_c, s_p) \equiv \bigwedge_i s_i \wedge s_c \leftrightarrow s_p]$ where $s_i \in S'$.

An attack relates the truth values of two different attributes so as to embed a cause-consequence relationship between the two. For example, for the attributes $s_c =$"*vulnerable to sshd BOF on machine A*" and $s_p =$"*root access privilege on machine A*", $Att(s_c, s_p)$ is an attack – the sshd buffer overflow attack. We would like to clarify here that the bi-conditional logical connective "$\leftrightarrow$" between s_c and s_p does not imply that s_p can be set to *true* only by using $Att(s_c, s_p)$; rather it means that given the sshd BOF attack, the only way to make s_p *true* is by having s_c *true*. In fact, $Att($"*vulnerable to local BOF on setuid daemon on machine A*",$s_p)$ is also a potential attack. The ϕ–attack is included to account for attributes whose truth values do not have any direct relationship. However, an indirect relationship can be established collectively. For example, the attributes $s_{c_1} = $ "*running SSH1 v1.2.25 on machine A*" and $s_{c_2} = $ "*connectivity(machine B, machine A)*" cannot individually influence the truth value of s_c, but can collectively make s_c *true*, given they are individually *true*. In such a case, $Att(s_{c_1}, s_c)$ and $Att(s_{c_2}, s_c)$ are ϕ–attacks.

Definition 8.4 ATTACK TREE. Let A be the set of attacks, including the ϕ–attacks. An attack tree is a tuple $AT = (s_{root}, S, \tau, \varepsilon)$, where

1. s_{root} is an attribute which the attacker wants to become *true*.

2. $S = N_{internal} \cup N_{external} \cup \{s_{root}\}$ is a multiset of attributes. $N_{external}$ denotes the multiset of attributes s_i for which $\nexists a \in A | s_i \in post(a)$. $N_{internal}$ denotes the multiset of attributes s_j for which $\exists a_1, a_2 \in A | [s_j \in pre(a_1) \wedge s_j \in post(a_2)]$.

3. $\tau \subseteq S \times S$. $(s_{pre}, s_{post}) \in \tau$ if $\exists a \in A | [s_{pre} \in pre(a) \wedge s_{post} \in post(a)]$. Further, if $s_i \in S$ and has multiplicity n, then $\exists s_1, s_2, \ldots, s_n \in S | (s_i, s_1), (s_i, s_2), \ldots, (s_i, s_n) \in \tau$.

Figure 8.2: Example attack tree.

4. ε is a set of decomposition tuples of the form $\langle s_j, d_j \rangle$ defined for all $s_j \in N_{internal} \cup$ $\{s_{root}\}$ and $d_j \in \{AND, OR\}$. d_j is AND when $\bigwedge_i [s_i \wedge (s_i, s_j) \in \tau] \leftrightarrow s_j$ is *true*, and OR when $\bigvee_i [s_i \wedge (s_i, s_j) \in \tau] \leftrightarrow s_j$ is *true*.

Fig. 8.2 shows an example attack tree, with the attribute "*root access on machine A*" as s_{root}. The multiset S forms the nodes of the tree. The multiset $N_{external}$ specify the leaf nodes of the tree. These nodes reflect the initial vulnerabilities present in a network and are prone to exploits. Since, an attribute can be a precondition for more than one attack, it might have to be duplicated, hence forming a multiset. The attribute "*machine B can connect to machine A*" in the example is one such attribute. The set of ordered pairs, τ, reflect the edges in the tree. The existence of an edge between two nodes imply that there is a direct or indirect relationship between their truth values, signified by the decomposition at each node. The *AND* decomposition at a node requires all child nodes to have a truth value of *true* for it to be *true.* The *OR* decomposition at a node requires only one child node to have a truth value of *true* for it to be *true.* Using these decompositions, the truth value of an attribute $s_j \in N_{internal} \cup \{s_{root}\}$ can be evaluated after assigning a set of truth values to the attributes $s_i \in N_{external}$. Fig. 8.3 shows the attack tree for our example network model. It depicts a clear picture of the different attack scenarios possible, as outlined in the previous section.

Figure 8.3: Attack tree of example network model.

8.4 Cost Model

In order to defend against the attacks possible, a security manager (decision maker) can choose to implement a variety of safeguard technologies, each of which comes with different costs and coverage. For example, to defend against the ftp/.rhost exploit, one might choose to apply a security patch, disable the FTP service, or simply tighten the write protection on the .rhost directory. Each choice of action can have a different cost. Besides, some measures have multiple coverage, but with higher costs. A security manager has to make a decision and choose to implement a subset of these policies in order to maximize the resource utilization. However, given the number of permutations possible in choosing this subset (2^n for n policies), this decision is not a trivial task.

Security planing begins with risk assessment which determines threats, loss expectancy, potential safeguards and installation costs. Many researchers have studied risk assessment schemes, including the National Institute of Standards and Technology (NIST) [160]. For simplicity, the security manager can choose to evaluate the risks by considering a relative magnitude of loss and hardening costs [19, 108, 160]. However, relative-cost approaches do not provide sufficient information to prioritize security measures especially

when the organization faces resource constraints. We adapt Butler's multi-attribute risk assessment framework [29, 30] to develop quantitative risk assessments for our security optimization. Butler's framework enables an aggregated representation of the various factors dominating the business model of an organization. First we define the notion of a security control in the context of the attack tree definition.

Definition 8.5 SECURITY CONTROL. Given an attack tree $(s_{root}, S, \tau, \varepsilon)$, the mapping $SC : N_{external} \rightarrow \{true, false\}$ is a security control if $\exists s_i \in N_{external} | SC(s_i) = false$.

In other words, a security control is a preventive measure to falsify one or more attributes in the attack tree, so as to stop an attacker from reaching its goal. Further, in the presence of multiple security controls SC_k, the truth value of an attribute $s_i \in N_{external}$ is taken as $\bigwedge_k SC_k(s_i)$. Given a security control SC, the set of all $s_i \in N_{external} | SC(s_i) = false$ is called the *coverage* of SC. Hence, for a given set of security controls we can define the *coverage matrix* specifying the coverage of each control. For a given set of m security controls, we use the Boolean vector $\vec{T} = (T_1, T_2, \ldots, T_m)$ to indicate if a security control is chosen by a security manager. Note that the choice of this vector indirectly specifies which attributes in the attack tree (leaf nodes) would be *false*.

8.4.1 Evaluating Potential Damage

The potential damage, P_j, represents a unit-less damage value that an organization may have to incur in the event that an attribute s_j becomes *true.* Based on Butler's framework, we propose four steps to calculate the potential damage for an attribute s_j.

Step1: Identify potential consequences of having a *true* value for the attribute, induced by some attack. In our case, we have identified five outcomes – lost revenue (monetary), non-productive downtime (time), damage recovery (monetary), public embarrassment (severity) and law penalty (severity) – denoted by $x_{1j}, x_{2j}, x_{3j}, x_{4j}$ and x_{5j}.

Step2: Estimate the expected number of attack occurrence, $Freq_j$, resulting in the consequences. A security manager can estimate the expected number of attack from the

organization-based historical data or public historical data[1].

Step3: Assess a single value function, $V_{ij}(x_{ij})$, for each possible consequence. The purpose of this function is to normalize different unit measures so that the values can be summed together under a single standard scale.

$$V_{ij}(x_{ij}) = \frac{x_{ij}}{\underset{j}{Max}\, x_{ij}} \times 100 \qquad ,1 \leq i \leq 5 \tag{8.1}$$

Step4: Assign a preference weight factor, W_i, to each possible consequence. A security manager can rank each outcome on a scale of 1 to 100. The outcome with the most concern would receive 100 points. The manager ranks the other attributes relative to the first. Finally, the ranks are normalized and set as W_i.

The potential damage for the attribute can then be calculated as given by the following equation.

$$P_j = Freq_j \times \sum_{i=1}^{5} W_i V_{ij}(x_{ij}) \tag{8.2}$$

When using an attack tree, a better quantitative representation of the cost is obtained by considering the residual damage once a set of security policies are implemented. Hence, we augment each attribute in the attack tree with a value signifying the amount of potential damage residing in the subtree rooted at the attribute and the attribute itself.

Definition 8.6 AUGMENTED-ATTACK TREE. Given an attack tree $AT = (s_{root}, S, \tau, \varepsilon)$, an augmented-attack tree $AT_{aug} = AT|\langle I, V \rangle$ is obtained by associating a tuple $\langle I_i, V_i \rangle$ to each $s_i \in S$, where

1. I_i is an indicator variable for the attribute s_i, where

$$I_i = \begin{cases} 0 & , if\ s_i\ is\ false \\ 1 & , if\ s_i\ is\ true \end{cases}. \tag{8.3}$$

2. V_i is a value associated with the attribute s_i.

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[1]Also known as an incident report published annually in many sites such as CERT/CC or SANS.ORG.

In this work, all attributes $s_i \in N_{external}$ are given a zero value. The value associated with $s_j \in N_{internal} \cup \{s_{root}\}$ is then computed recursively as follows.

$$V_j = \begin{cases} \sum_{k|(s_k,s_j)\in\tau} V_k + I_j P_j & , if\ d_j\ is\ AND \\ \max_{k|(s_k,s_j)\in\tau} V_k + I_j P_j & , if\ d_j\ is\ OR \end{cases} \tag{8.4}$$

Ideally, P_j is same for all identical attributes in the multiset. We took a "panic approach" in calculating the value at each node, meaning that given multiple subtrees are rooted at an attribute with an *OR* decomposition, we choose the maximum value. We do so because an attacker's capabilities and preferences cannot be known in advance. The residual damage of the augmented tree is then defined as follows.

Definition 8.7 RESIDUAL DAMAGE. Given an augmented-attack tree $(s_{root}, S, \tau, \varepsilon)|\langle I, V\rangle$ and a vector $\vec{T} = (T_i)$, $T_i \in \{0,1\}; 1 \leq i \leq m$, the residual damage is defined as the value associated with s_{root}, i.e.

$$RD(\vec{T}) = V_{root}. \tag{8.5}$$

8.4.2 Evaluating Security Cost

Similar to the potential damage, the security manager first lists possible security costs for the implementation of a security control, assigns the weight factor on them, and computes the normalized value. The only difference is that there is no expected number of occurrence needed in the evaluation of security cost. In this study, we have identified five different costs of implementing a security control – installation cost (monetary), operation cost (monetary), system downtime (time), incompatibility cost (scale), and training cost (monetary). The overall cost C_j, for the security control SC_j, is then computed in a similar manner as for potential damage, with an expected frequency of one. The total security cost for a set of security controls implemented is then defined as follows.

Definition 8.8 TOTAL SECURITY CONTROL COST. Given a set of m security controls, each having a cost $C_i; 1 \leq i \leq m$, and a vector $\vec{T} = (T_i)$, $T_i \in \{0,1\}; 1 \leq i \leq m$, the total

Table 8.3: Security controls for example network model.

Security Control	Action
SC_1/SC_2	Disable/Patch suid @ 196.216.0.2
SC_3/SC_4	Disable/Patch LICQ @ 196.216.0.2
SC_5	Disable "at" @ 196.216.0.3
SC_6/SC_7	Disable/Patch LICQ @ 196.216.0.3
SC_8	Disable Rsh @ 196.216.0.1
SC_9	Disable Ftp @ 196.216.0.1
SC_{10}	Disconnect Internet @ 196.216.0.1
SC_{11}	Chmod home directory @ 196.216.0.1
SC_{12}/SC_{13}	Disable/Patch Ftp @ 196.216.0.10
SC_{14}/SC_{15}	Disable/Patch SSH @ 196.216.0.10
SC_{16}	Disconnect Internet @ 196.216.0.10
SC_{17}	Disable Rsh @ 196.216.0.10
SC_{18}	Patch FTP/.rhost @ 196.216.0.10
SC_{19}	Chmod home directory @ 196.216.0.10

security control cost is defined as

$$SCC(\vec{T}) = \sum_{i=1}^{m}(T_iC_i). \tag{8.6}$$

8.5 Problem Formulation

The two objectives we consider in this study are the total security control cost and the residual damage in the attack tree of our example network model. For the attack tree shown in Fig. 8.3, we have identified 19 different security controls possible by patching or disabling of different services, as well as by changing file access permissions. With about half a million choices available (2^{19}), an enumerated search would not be an efficient approach to find the optima. The security controls are listed in Table 8.3. We also tried to maintain some relative order of importance between the different services, as in a real-world scenario, when assigning the potential damage and security control costs during the experimental evaluation.

Problem 8.1 The Single-objective Optimization Problem. Given an augmented-attack tree $(s_{root}, S, \tau, \varepsilon)|\langle I, V\rangle$ and m security controls, find a vector $\vec{T}^* = (T_i^*)$, $T_i^* \in \{0,1\}; 1 \le i \le m$, which minimizes the function

$$\alpha RD(\vec{T}) + \beta SCC(\vec{T}), \tag{8.7}$$

where α and β are preference weights for the residual damage and the total cost of security control respectively, $0 \leq \alpha, \beta \leq 1$ and $\alpha + \beta = 1$.

The single-objective problem is the most likely approach to be taken by a decision maker. Given only two objectives, a preference based approach might seem to provide a solution in accordance with general intuition. However, as we find in the case of our example network model, the quality of the solution obtained can be quite sensitive to the assignment of the weights. To demonstrate this affect, we run multiple instances of the problem using different combination of values for α and β. α is varied in the range of $[0,1]$ in steps of 0.05. β is always set to $1 - \alpha$.

Problem 8.2 The Multi-objective Optimization Problem. Given an augmented-attack tree $(s_{root}, S, \tau, \varepsilon) | \langle I, V \rangle$ and m security controls, find a vector $\vec{T}^* = (T_i^*)$, $T_i^* \in \{0,1\}; 1 \leq i \leq m$, which minimizes the total security control cost and the residual damage.

The next level of sophistication is added by formulating the minimization as a multi-objective optimization problem. The multi-objective approach alleviates the requirement to specify any weight parameters and hence a better global picture of the solutions can be obtained.

Problem 8.3 The Multi-objective Robust Optimization Problem. Let $\vec{T} = (T_i)$ be a Boolean vector. A perturbed assignment of radius r, $\vec{T}_r$, is obtained by inverting the value of at most r elements of the vector $\vec{T}$. The robust optimization problem can then be defined as follows.

Given an augmented-attack tree $(s_{root}, S, \tau, \varepsilon) | \langle I, V \rangle$ and m security controls, find a vector $\vec{T}^* = (T_i^*)$, $T_i^* \in \{0,1\}; 1 \leq i \leq m$, which minimizes the total security control cost and the residual damage, satisfying the constraint

$$\max_{\vec{T}_r} RD(\vec{T}_r) - RD(\vec{T}) \leq D, \tag{8.8}$$

where D is the maximum perturbation allowed in the residual damage.

The third problem is formulated to further strengthen the decision process by determining robust solutions to the problem. Robust solutions are less sensitive to failures in

security controls and hence subside any repeated requirements to re-evaluate solutions in the event of a security control failure.

We use a simple genetic algorithm (SGA) [75] to solve Problem 8.1. NSGA-II is used to solve Problems 8.2 and 8.3. The algorithm parameters are set as follows: population size $= 100$, number of generations $= 250$, crossover probability $= 0.9$, and mutation probability $= 0.1$.

8.6   Empirical Results

We first present the sensitivity results of NSGA-II and SGA to their parameters. Increasing the population size from 100 to 500 gives us a faster convergence rate, although the solutions reported still remains the same. The effect of changing the crossover probability in the range of 0.7 to 0.9 does not lead to any significant change of the solutions obtained. Similar results were observed when changing the mutation probability from 0.1 to 0.01. The solutions also do not change when the number of generations is changed from 250 to 500. Since we did not observe any significant change in the solutions by varying the algorithm parameters, the following results are presented as obtained by setting the parameters as chosen in the previous section.

It is usually suggested that the preference based approach should normalize the functions before combining them into a single function. However, we did not see any change in the solutions of the normalized version of Problem 8.1. Fig. 8.4 shows the solutions obtained from various runs of SGA in Problem 8.1 with varying α. A decision maker, in general, may want to assign equal weights to both the objective functions, i.e. set $\alpha = 0.5$. It is clear from the figure that such an assignment does not necessarily provide the desired balance between the residual damage and the total security control cost. Furthermore, such balance is also not obtainable by assigning weight values in the neighborhood of 0.5. The solutions obtained are quite sensitive to the weights, and in this case, much higher preference must be given to the total security control cost to find other possible solutions. Since the weights do not always influence the objectives in the desired manner, understanding their effect is not a trivial task for a decision maker. It is also not possible to always do an exhaustive analysis of the affect of the weights on the objectives. Given such

Figure 8.4: SGA solutions to Problem 8.1 with α varied from 0 to 1 in steps of 0.05.

situations, the decision maker should consider obtaining a global picture of the trade-offs possible. With such a requirement in mind, we next consider Problem 8.2.

The two solutions corresponding to $\alpha = 0.25$ and 0.1 in Fig. 8.4, including any other solutions in the vicinity, are likely candidates for a decision maker's choice. Unlike the single-objective approach, where determining such vicinal solutions could be difficult, the multi-objective optimization approach clearly revealed the existence of at least one such solution. Fig. 8.5 shows the solutions obtained from a single run of NSGA-II on Problem 8.2. NSGA-II reported all the solutions obtained from multiple runs of SGA, as well as three more solutions. Interestingly, there exists no solution in the intermediate range of [25,45] for residual damage. This inclination of solutions towards the extremities of the residual damage could be indicative of the non-existence of much variety in the security controls under consideration. The number of attack scenarios possible is also a deciding factor. Most of the security controls for the example network involve either the disabling or patching of a service, resulting in a sparse coverage matrix. For a more "continuous" Pareto-front, it is required to have security controls of comparative costs and capable of covering multiple services. A larger, more complex real-world problem would likely have more attack scenarios and a good mixture of both local and global security controls, in

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Figure 8.5: NSGA-II solutions to Problem 8.2 and sensitivity of a solution to optimum settings.

which case, such gaps in the Pareto-front will be unlikely.

Once the decision maker has a better perspective of the solutions possible, further analysis of the solutions may be carried out in terms of their sensitivity to security control failures. Such sensitivity analysis is helpful in not only reducing valuable decision making time, but also to guarantee some level of fault tolerance in the network. Fig. 8.5 shows the sensitivity of one of the solutions to a failure in one of the security controls corresponding to the solution. This solution, with security controls SC_4 and SC_{11}, will incur a high residual damage in the event of a failure of SC_4. Thus, a decision maker may choose to perform a sensitivity analysis on each of the solutions and incorporate the results thereof in making the final choice. However, the decision maker then has no control on how much of additional residual damage would be incurred in the event of failure. Problem 8.3 serves the requirements of this decision stage by allowing the decision maker to specify the maximum allowed perturbation in the residual damage. It is also possible to specify the scope of failure – the radius r – within which the decision maker is interested in analyzing the robustness of the solutions. For this study, we are mostly interested in obtaining solutions that are fully robust, meaning the residual damage should not

Table 8.4: Fully robust solutions obtained by NSGA-II with $r = 1$.

	Robust-optimum security controls	RD	SCC
R1	$SC_9, SC_{11}, SC_{13}, SC_{15}, SC_{16}, SC_{19}$	0.0	26.0
R2	$SC_3, SC_4, SC_9, SC_{11}, SC_{18}, SC_{19}$	10.5	21.0
R3	$SC_3, SC_4, SC_7, SC_{11}$	13.5	12.0
R4	SC_3, SC_4	22.8	8.0
R5	SC_7, SC_{11}	49.5	4.0
R6	null	58.8	0.0

Figure 8.6: NSGA-II solutions to Problem 8.3 with $D = 0$ and $r = 1$. Problem 8.2 solutions are also shown for comparison.

increase, and hence set D to zero. Also, because of the sparse nature of the coverage matrix, we set the perturbation radius r to 1. Fig. 8.6 shows the solutions obtained for this problem.

The solutions to Problem 8.3 reveal that none of the optimum solutions previously obtained, except the trivial zero SCC solution, is fully robust even for a single security control failure. Such insight could be of much value for a decision maker when making a final choice. Table 8.4 shows the security controls corresponding to the robust solutions. With the final goal of obtaining a solution with a good balance between the residual damage and the total security control cost, the decision maker's choice at this point can be justifiably biased towards the selection of solution R3.

Figure 8.7: Compressed attack tree showing residual damage computation with R3 as security control set.

We present certain interesting properties exploited by solution R3 from the attack tree. To point out the salient features, we compress the attack tree for our example network model as shown in Fig. 8.7. The compressed tree is obtained by collapsing all subtrees to a single node until a node covered by a security control from R3 contributes to the calculation of the residual damage. All such nodes, represented by rectangles in the figure, is labeled with the maximum residual damage that can propagate to it from the child subtree and $(+)$ the damage value that can occur at the node itself. A triangular node represents the security controls that can disable that node. The individual damage value is accrued to the residual damage from the child node only if the attached security control, if any, fails.

The solution R3 clearly identifies the existence of the subtrees $ST_1 = \{\{n_7, n_{10}\}, \{n_8, n_{11}\}, \{n_9, n_{12}\}\}$ and $ST_2 = \{\{n_3, n_7, n_{10}\}, \{n_6, n_9, n_{12}\}\}$. In the event of a failure of SC_{11}, n_7 would collect a value of 10.8. Since n_3 has an *AND* decomposition with SC_7, it will be disabled, thereby not contributing its individual damage value of 12 to the residual damage at that node (10.8). On the other hand, if SC_7 fails, SC_{11} will disable n_7 which in turn will disable n_3. In fact, in this case the residual damage at n_3 would be zero. Similarly, n_6 and n_8 also never propagate a residual damage of more than 10.8 to its parent node.

Consequently, n_2 never propagates a value more than 13.5. The individual cost of 36 at n_1 is never added to this residual damage value of 13.5 from n_2 since, owing to the AND decomposition, n_1 is always falsified by security controls SC_3 and SC_4, only one of which is assumed to fail at a time. The solution wisely applies security controls covering multiple attack scenarios, and at multiple points in those scenarios to keep the damage to a minimum.

8.7 Conclusions

In this chapter, we address the system administrator's dilemma, namely, how to select a subset of security hardening measures from a given set so that the total cost of implementing these measures is not only minimized but also within budget and, at the same time, the cost of residual damage is also minimized. One important contribution of our approach is the use of an attack tree model of the network to drive the solution. By using an attack tree in the problem, we are able to better guide the optimization process by providing the knowledge about the attributes that make an attack possible. Further, a systematic analysis enables us to approach the problem in a modular fashion, providing added information to a decision maker to form a concrete opinion about the quality of the different trade-off solutions possible.

The cost model that we adopt in this chapter is somewhat simplistic. We assume that, from a cost of implementation perspective, the security measures are independent of each other, where in real life they may not be so. In addition, we have assumed that the system administrator's decision is in no way influenced by an understanding of the cost to break the system. Furthermore, the possible decomposition of an attack tree to divide the problem into sub-problems is an interesting alternative to explore. Finally, there is a dynamic aspect to the system administrator's dilemma. During run time the system administrator may need to revise her decision based on emerging security conditions.

CHAPTER 9

Security Hardening on Pervasive Workflows

Pervasive computing aims at making the presence of computing machinery so transparent that their very presence becomes imperceptible to the end user. These applications involve interacting with heterogeneous devices having various capabilities under the control of different entities. Such applications make use of workflows that are mostly automated and do not require much human intervention. For critical applications, the workflows pass on sensitive information to the various devices and make crucial decisions. Failure to protect such information against security breaches may cause irreparable damages.

Pervasive computing applications impose a number of unique constraints that make choosing the appropriate security mechanisms difficult. Interoperability of the heterogeneous devices must be taken into account while selecting security mechanisms. Resource consumption is a very important consideration as this directly relates to the up-time and maintainability of the application. The cost of deployment must also be considered. Thus, an overall picture illustrating the cost-benefit trade-offs in the presence of these constraints is needed before a final decision can be made.

Unfortunately, security threats in a pervasive environment are very application dependent. Thus, it is not possible to give a solution that is satisfactory for all pervasive applications. For example, if a communication is between a mobile device and a base

station, then a particular type of authentication protocol may be appropriate, whereas if the communication is of an ad-hoc nature, the same protocol may be inappropriate. Further, the communication between two devices can be deemed sensitive or non-sensitive depending on the context under which the communication is taking place. Finally, the resource constraints varies for different scenarios and prohibit the usage of the same security measures even for the same class of threats.

Context based security provisioning has earlier been proposed to adapt the security level of a communication to the sensitivity of the information being exchanged [97, 132, 133]. Nonetheless, exploring the aforementioned trade-off options becomes difficult when contexts are introduced. Contexts are dynamic in nature and proactive analysis do not always capture all possible scenarios. However, it is important to realize that pervasive environments set up with a specific application in mind have predefined ways of handling the different scenarios that can appear during the lifetime of the environment. It is therefore possible that these scenarios be represented in a concise way and subjected to security evaluation techniques. This is a good beginning since an organization has a concrete understanding of the assets it has and the points of immediate interest usually involve the likelihood of potential damages to these known assets.

In this chapter, we formalize these issues and identify possible resolutions to some of the decision making problems related to securing a pervasive environment. We propose the use of workflow profiles to represent the business model of a pervasive environment and compute the cost associated with the maintenance of the workflow. We then perform a multi-objective analysis to maximize the security level of the workflow and minimize the cost incurred thereof. The multi-objective formulations take into account the energy constraints imposed by devices in the environment. We demonstrate our methodology using an evolutionary algorithm in a pervasive healthcare domain.

The rest of the chapter is organized as follows. Section 9.1 presents the related work in this field. An example healthcare pervasive environment along with the concept of workflow representations is presented in Section 9.2. Security provisioning in the workflow is discussed in Section 9.3. Section 9.4 presents the cost model for the workflow. Section 9.5 discusses the multi-objective problems, some empirical results on which are presented in

Section 9.6. Finally, Section 9.7 concludes the chapter.

9.1 Related Work

Security provisioning in pervasive environments is an open field for research. Campbell et al. present an overview of the specific security challenges that are present in this field [33] and describe their prototype implementation of a component-based middleware operating system. A more specific formulation for security provisioning in wireless sensor networks is presented by Chigan et al. [37]. Their framework has an offline security optimization module targeted towards maximizing a *security provision index*. Ranganathan et al. propose some meta-level metrics to gauge the ability of different security services in a pervasive environment [143].

Dependability issues related to the application of pervasive computing to the healthcare domain is discussed by Bohn et al. [26]. They argue that the healthcare domain can serve as a benchmark platform for pervasive computing research. They point out the security issues relevant to the setup of such a platform. Similar practical issues are also investigated by Black et al. [25].

An example usage of context information in pervasive applications is presented by Judd and Steenkiste [97]. Their approach allows proactive applications to obtain context information on an user's current environment and adapt their behavior accordingly. The use of context information for security provisioning is proposed by Mostéfaoui and Brézillon [132]. They propose using contextual graphs to appropriately decide on the security policies to enforce. Further reasoning on the contributions of combining context and security is provided by the same authors in [133]. Sanchez et al. propose a Monte Carlo based framework to model context data and evaluate context based security policies [153].

9.2 The Pervasive Workflow Model

Security threats in a pervasive environment are application dependent. Consequently, business models investing in any kind of a pervasive computing paradigm will highly benefit if formalisms are derived to enable a "case-by-case" study of the problem of se-

curity provisioning. We therefore discuss our approach using an example healthcare application.

9.2.1 A pervasive health care environment

The pervasive healthcare environment consists of devices that measure the vital signs of patients, location sensors that locate mobile resources, location-aware PDAs carried by health-care personnel, and back-end systems storing and processing records of patient data. The devices are connected through wired or wireless medium. The application consists of different workflows that get triggered by various events. The following example specifies the workflow that handles the situation when an unanticipated change occurs in a patient's vital signs (VS) monitor.

Case1: The VS monitor tries to detect the presence of the doctor within a wireless communicable distance. If the doctor is present, she can make suggestions which may or may not be based on the patient report stored at the back-end. She may also decide to request the assistance of a nurse, who is located with the help of the network infrastructure. In case of an emergency, the same infrastructure is used to notify the emergency service.

Case2: If a doctor cannot be located nearby, there is a search for a nurse. The nurse may have the requisite skills to take care of the situation, perhaps with information obtained from the back-end system. If not, the nurse requests the network infrastructure to locate a remote doctor. The remote doctor can then make suggestions to the nurse or directly interact with the monitoring devices using the network. Possibilities are also that the doctor feels the need to be immediately with the patient and informs the emergency service on her way.

Case3: If a nearby doctor or a nurse cannot be located, the VS monitor communicates with the network infrastructure to locate a remote doctor. The doctor, once located, can remotely interact with the monitoring equipments, or decide to attend to the situation physically, often asking for assistance from a nurse. Emergency services are notified on a need basis. Also, on the event that the network is unable to locate

the doctor, it informs the emergency service.

9.2.2 Computing and communication infrastructure

A pervasive application requires communication between different devices with varying degrees of processing power and resource constraints. We classify these devices into three categories: *adapters*, *composers*, and *back-end*. Adapters are devices with low processing capabilities driven by a battery source or a wired power supply. They are responsible for collecting raw sensor data and forwarding it to another suitable device. A limited amount of processing can also be performed on the collected data before forwarding them. Composers have medium processing capabilities and may have a fixed or battery driven power source. They interact with the adapters and interpret much of the data collected by them, most likely with aid from the back-end. The back-end has high processing capabilities driven by a wired power supply. Databases relevant to the pervasive environment reside in the back-end. Fig. 9.1 depicts the typical interactions that can happen between the three device classes.

Figure 9.1: Component interactions in a pervasive environment.

Examples of adapters in the pervasive healthcare environment are the devices that monitor a patient's vital signs and location sensors present in the facility that help discover a mobile resource. A composer can be a location-aware PDA carried by a doctor or a nurse, a laptop, a data relay point present as part of the network infrastructure, or the system monitored by the emergency personnel. The back-end in this example are data servers used to store patients' medical records or the high-end systems available to perform computationally intensive tasks. Note that the back-end may not be reachable directly by all composers. In such cases, a composer (a personnel's PDA, for example) will first communicate with another composer (a data relay point perhaps) which will then route the request (may be using other data relay points) to the back-end system. Adapters may communicate using a wired/wireless medium with the data relay points,

which in turn communicate over an infrastructure network to the back-end system. The composer class comprising mainly of handheld PDAs and similar devices communicate wirelessly with adapters and other composers.

9.2.3 Workflow

A workflow captures the various relationships between the participating nodes and provides a concise representation of the different contexts under which the different nodes communicate with each other.

Definition 9.1 WORKFLOW. A workflow is a tuple $\langle N, E, n \rangle$ representing one or more execution paths of a business model, where N is a multiset of nodes representing the devices in the application, E is a set of ordered pairs of the form $(n_s, n_d) \in N \times N$ denoting the communication links, and $n \in N$ is a source node that triggers the workflow.

A workflow can also be visualized in terms of transfer of work between devices. A workflow is a meta-level representation of the order in which different devices participate to achieve one or more business goals. A *feasible execution path* in such a representation resembles the order of participation of the nodes to achieve one of the goals. For example, to find the nearest doctor, transfer of work progresses as VS Monitor→Data Relay Point→Location Sensor→Data Relay Point→ ... →Doctor, and signifies an *execution path*. Although this transfer of work involves multiple location sensors and multiple data relay points, the workflow representation does not make a distinction among them. This is primarily because the objective behind the usage of a communication link between a data relay point and a location sensor is fixed (find a doctor) and hence all such links are assumed to exchange information with the same level of sensitivity.

9.2.4 Context

Although a workflow helps identify the different communication links used as part of different execution paths, it does not provide any information on the frequency with which a particular channel is used. This frequency estimate is required to determine the rate of power consumption of the two participating devices in the link. When security

measures are placed in these links, the rate of power consumption will increase depending on the computational requirements of the algorithms. Here we have an inherent conflict. Heavily used communication channels should ideally have security measures with low computing requirements in order to reduce the power consumption at the two participating devices. At the same time, strong security measures are perhaps required to safeguard the huge amount of information flowing through the channel. Quite often this is hard to achieve since strong security measures typically are computation intensive algorithms.

Definition 9.2 CONTEXT. A context is a prefix of some execution path through a workflow. It is associated with a probability estimate vector that gives the likelihood of the context changing into other contexts.

A context specifies the different nodes that are traversed when following a particular execution path. We use the notation $\mathcal{C}_v$ to represent a context with the current node v. In some cases, we use a more informal notation and describe contexts by just concatenating the names of the nodes. For example, for the context $ABCDC$, the current node is C and the node has been reached by following the path $A \rightarrow B \rightarrow C \rightarrow D \rightarrow C$. We use '$|$' to denote the operator to concatenate a node to a context, resulting in a new context. Note that a context may be a part of multiple execution paths. Context-based probability estimates will help capture the likelihood with which a context can change. This is described next.

Consider Fig. 9.2a; let $\mathcal{C}_v$ be the context and let t be its probability of occurrence. We assign a probability on each outgoing edge signifying the chances with which the current context changes. The context becomes $\mathcal{C}_v|x$ with probability p_1 and $\mathcal{C}_v|y$ with probability p_2.

For a node v with k outgoing edges numbered in a particular order, the context-based probability vector $(p_1, p_2, \ldots, p_k)_{\mathcal{C}_v}$ gives the probabilities on the outgoing edges in the same order when the current context is $\mathcal{C}_v$. The probability values for a given context can be obtained from audit logs collected over a period of time signifying how often did v act as an intermediate node in the communication between two neighbor nodes under a particular communication sequence.

Figure 9.2: (a) Context-based probability estimates. (b) Workflow probability calculation.

It is important to realize that a workflow by itself does not reveal the feasible execution paths. It is only by including context information that the feasible execution paths get defined. Context-based probability estimates tell us if an adjacent node can become a part of a particular execution path. If all probabilities on the outgoing edges of a particular node are zero, then the execution does not proceed and the current context at that point terminates as a feasible execution path. Hence, by defining a set of possible contexts, along with their context-based probability estimates, we have a precise way of defining which execution paths are feasible. We call this set the *context set* of the workflow.

Definition 9.3 CONTEXT SET. A context set for a workflow is a set of contexts that are all possible prefixes of all feasible execution paths.

The context set contains only those contexts for which there is non-zero probability of transitioning into another context. To infer the feasible execution paths from a given context set, we start at the source node and check to see if the current context (the source node alone at this point) is present in the context set. We can move onto an adjacent node if the probability on the edge to it is non-zero. The current context then changes to a different one for each reachable node. The process is repeated for all such nodes until a node is reached where the current context is not present in the context set. Such a context is then a feasible execution path.

Note that context-based probability estimates provide the probability with which an outgoing edge will be used in the current context. This probability does not, as yet, pro-

vide an estimate of the overall frequency with which the edge is used in the workflow. We shall show in Section 9.4 how the context set is used to compute the effective probability with which a particular communication link in the workflow gets used.

9.3 Security Provisioning

The problem of security provisioning in a workflow involves the identification of potentially damaging attacks and the security mechanisms that can be adopted to protect against such attacks. Depending on the nature of communication between two nodes, a subset of the known attacks can be more prominent than others in a communication channel. Besides the ability to defend against one or more attacks, the choice of a security mechanism is also dependent on the resources it consumes during execution.

Definition 9.4 SECURITY MECHANISM. Given a set of A attacks, denoted by $a_1, a_2, \ldots, a_A$, a security mechanism S_i is a Boolean vector $[S_{i1}, S_{i2}, \ldots, S_{iA}]$, where S_{ij} is 1 if it defends against attack a_j, 0 otherwise.

An attack in this definition refers to the consequence of a malicious activity. A security mechanism is capable of preventing one or more attacks. For a given set of N_S security mechanisms, we have a coverage matrix defining which attacks are covered by which mechanisms. Further, each mechanism has an associated power consumption rate, denoted by SMC_i, where $1 \leq i \leq N_S$.

To facilitate the enforcement of different security mechanisms along different communication links, we augment each edge on the workflow with an *attack vector* specifying the attacks which are of concern in the link.

Definition 9.5 ATTACK VECTOR. An attack vector on an edge of the workflow is a Boolean vector of size A with the jth component being either 1 or 0 based on whether attack a_j is plausible on the edge or not.

Given an attack vector, it is possible that no single security mechanism can provide the required defenses against all attacks of concern on the edge. Multiple mechanisms have to be selected such that they can collectively provide the coverage for the attacks.

Definition 9.6 SECURITY CONTROL VECTOR. A security control vector $SV_e = [SV_{e1}, SV_{e2}, \ldots, SV_{eN_s}]$ on the edge e is a Boolean vector, where SV_{ei} is 1 if the security mechanism S_i is chosen, 0 otherwise.

For a particular security control vector SV_e, the attacks collectively covered by SV_e is computed from the expression $\bigvee_i (SV_{ei} \cdot S_i)$, for $i = 1, \ldots, N_s$. The 'dot' operator here indicates scalar multiplication with a vector and '$\vee$' signifies the Boolean *OR* operation. The resultant of this expression is a Boolean vector of size A and signifies which attacks are covered by the combination of the security mechanisms. We shall call this vector the *covered attack vector* of the edge on which the security control vector operates.

If AV_e and CAV_e are the attack vector and covered attack vector on an edge e, then the Hamming distance between the zero vector and the vector $AV_e \wedge \neg CAV_e$, '$\wedge$' and '$\neg$' signifying the boolean *AND* and *NOT* operators respectively, computes the number of attacks initially specified as plausible on the edge but not covered by any security mechanism in the control vector. We shall denote this quantity by $H(AV_e, CAV_e)$.

9.4 Cost Computation

In this study, we are interested in the cost that an organization has to incur to keep a particular workflow running. To this effect, we consider the cost of maintenance. The cost of maintenance relates to the expenses that an organization has to incur to hire personnel for regular maintenance rounds, purchase supplies and hardware support equipments, or may be due to losses arising from downtime in services during maintenance. A reduction in the cost is possible if the workflow can be engineered to run for a longer time between two maintenance rounds. We realize that the contributing factors appear with different magnitudes and in different dimensions, often with different levels of impact, and are hence difficult to combine into one cost measure. We can adapt Butler's Multi-attribute Risk Assessment model [29, 30] to cater to these difficulties.

Maintenance cost estimates obtained using Butler's method is used along with frequency estimates obtained from the workflow to determine the total maintenance cost incurred to keep the workflow running. Before we can do so, the probability estimates on

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Procedure 9.1 EffectiveProbability(Context *c*, Probability *t*)

Input: Context *c* and probability *t*. {*context set $\mathcal{C}$, context based probability estimates and workflow graph are assumed global data*}

Output: Effective probability of edges. {*initialized to* 0.0 *on all edges*}

 1: **if** $(c \in \mathcal{C})$ **then**
 2:     $h =$ current node in context *c*
 3:     **for all** (edge *e* outgoing to a node *g* from *h*) **do**
 4:         $p_e =$ probability of going to *g* from *h* in the context *c*
 5:         Add $p_e t$ to effective probability of edge *e*
 6:         $EffectiveProbability(c|g, p_e t)$
 7:     **end for**
 8: **end if**

---

the communication links have to be aggregated to determine the overall frequency with which the link is used. This is done by calculating the *effective probability* of usage of a communication link on the event the workflow gets triggered.

Refer to Fig. 9.2a. Since the probabilities on the outgoing edges are decided independent of each other, the effective probability on the two outgoing edges can be calculated as $p_1 t$ and $p_2 t$ respectively. Also, since the probabilities on an outgoing edge are only dependent on the current context, effective probabilities accumulating on an edge from different contexts can be summed together to give the probability with which the edge is used. Fig. 9.2b shows the calculation of the effective probabilities for a small workflow. The workflow has node *A* as the source node. The outgoing edges from the source node capture all possible situations that can occur when the workflow is triggered. The given context probabilities are ordered according to the numbering on the outgoing edge at the current node. Procedure 9.1 when instantiated with arguments $(A, 1.0)$ recursively computes the effective probabilities on the edges of the workflow given the context probabilities shown in the figure. We denote the effective probability on an edge *e* by p_e. Once the effective probabilities on each edge of the workflow are calculated, the number of times a particular edge is used can be found by multiplying the number of times the workflow is triggered with the effective probability on the edge.

Let P_i and MC_i be the power capacity and total maintenance cost of the ith unique device respectively. Let $\{e_1, e_2, \ldots, e_k\}$ be the set of edges that connect the nodes corresponding to this device with other nodes. Let T_i be a constant power consumption by the

device and $PC_{ij} = \sum_{l=1}^{N_S}(SV_{e_jl} \times SMC_l)$ the power consumption rate of the device because of the security mechanisms in place on edge $e_j; j = 1, \ldots, k$. If the workflow is triggered F times, then edge e_j with effective probability p_{e_j} will be used $f = F \times p_{e_j}$ times. Hence, the total power consumed at device i after the security provisioning on edge e_j is expressed as $(T_i + PC_{ij}) \times f$. A maintenance will be required for the device every $\sum_{j=1}^{k} \frac{(T_i + PC_{ij}) \times f}{P_i}$ times of usage. The total maintenance cost for the workflow is

$$TMC = \sum_i \left(MC_i \times \sum_{j=1}^{k} \frac{(T_i + PC_{ij}) \times f}{P_i} \right). \tag{9.1}$$

9.5 Problem Formulation

The multi-objective formulations presented here are intended to help analyze the trade-offs resulting from the selection of a particular set of security control vectors for the different communication links in a workflow and the corresponding cost of maintenance. To begin with, we decide on a subset $E_p \subseteq E$ of edges on the workflow that are subjected to the security provisioning procedure.

The first optimization problem we consider is the determination of security control vectors for each edge in E_p in order to minimize the cost of maintenance and the total number of attacks left uncovered on the edges of the workflow, i.e. minimize $\sum_{e \in E_p} H(AV_e, CAV_e)$.

Problem 9.1 Scenario A. Find security control vectors for each edge $e \in E_p$ that minimizes TMC and minimizes $\sum_{e \in E_p} H(AV_e, CAV_e)$.

Although the total number of uncovered attacks provide a good idea about the potential exploits still remaining in the workflow, the damages that can result from the exploitation of the uncovered attacks can be more than that could have resulted from the covered attacks. The choice of security control vectors based on the number of covered attacks can thus be a misleading indicator of the assets that the employed security mechanisms helped protect. To this effect, instead of minimizing the number of uncovered attacks, the second formulation incorporates the minimization of the total potential damage that can result from the uncovered attacks. To facilitate the computation of the total

potential damage, we modify the attack vector to indicate the damages possible instead of just a Boolean indicator of whether an attack is plausible in an edge or not. The jth component of the attack vector is then a real valued quantity signifying the *potential damage cost* if attack a_j is not covered by a security mechanism on the edge. The quantity can be zero if the corresponding attack is not of concern on the particular edge. We do not focus on the cost models that can be adopted to estimate such damage levels. Butler's framework is a good starting point in this direction.

Problem 9.2 Scenario B. Find security control vectors for each edge $e \in E_p$ that minimizes TMC and minimizes $\sum_{e \in E_p} \langle AV_e, \neg CAV_e \rangle$, where $\langle , \rangle$ signifies the scalar product operation.

An assumption implicit in the above two formulations is that every security mechanism is capable of running in all the devices present in the workflow. This is not true when there exists devices with very low power capabilities and not all security mechanisms can be supported by them. The existence of such devices impose the constraint that certain security mechanisms can never be placed on certain communication links. Thus, we extend Problem 9.2 to generate solutions that are feasible within such constraints. Let $n_{s,e}$ and $n_{d,e}$ denote the two communicating devices on the edge e. For the security mechanisms to be able to execute, the total power consumed by the mechanisms in place on this edge has to be less than the minimum of the power capacities of the two participating devices. The optimization problem is then formulated as follows.

Problem 9.3 Scenario C. Find security control vectors SV_e for each edge $e \in E_p$ which minimizes TMC and minimizes $\sum_{e \in E_p} \langle AV_e, \neg CAV_e \rangle$, satisfying the constraints $\sum_{i=1}^{N_S} (SV_{ei} \times SMC_i) \leq min(P_{n_{s,e}}, P_{n_{d,e}})$, for all edges $e \in E_p$.

For the final problem, we explore the scenario when the adopted security mechanisms are not robust enough and are prone to failures. Non-robust security control vectors suffer from the drawback that a failure in one of the security mechanisms can heavily increase the number of uncovered attacks or the total potential damage in the workflow. Robust solutions, on the other hand, are able to contain such increases within a pre-specified acceptable level. We use the notion of a failure radius r which signifies the number of

security mechanisms that can fail at a time. For a given failure radius, we can specify an acceptable level D of increase in the total number of uncovered attacks, or the total potential damage, in the event of failure. The robust version of Problem 9.3 is then stated as follows.

Problem 9.4 Scenario D. Find security control vectors SV_e for each edge $e \in E_p$ which minimizes TMC and minimizes $PD = \sum_{e \in E_p} \langle AV_e, \neg CAV_e \rangle$, satisfying the constraints $\sum_{i=1}^{N_S} (SV_{ei} \times SMC_i) \leq min(PC_{n_{s,e}}, PC_{n_{d,e}})$, for all edges $e \in E_p$ and the constraint that the maximum increase in PD, resulting from at most r security mechanism failures, does not exceed D.

We employ the Non-Dominated Sorting Genetic Algorithm-II (NSGA-II) to solve the four multi-objective problems presented. A solution to a problem is represented by a boolean string generated by the concatenation of the security control vectors for each edge. We identified 23 different communication links of interest in the example healthcare workflow and 8 different security mechanisms, giving us a solution encoding length of $8 \times 23 = 184$. The parameters of the algorithm are set as follows: population size $= 300$, number of generations $= 1000$, crossover probability $= 0.9$, mutation rate $= 0.01$, and binary tournament selection.

Due to the non-availability of standard test data sets, the experiments performed involve hypothetical data. Nonetheless, the analysis do not make any reference to the absolute values obtained for the objectives from the optimization. The observations reveal what kind of cost-benefit information can such an analysis provide, irrespective of the exact numerical values of the quantities.

9.6 Empirical Results

The trade-off solutions obtained when minimizing the number of uncovered attacks and the total maintenance cost are shown in Fig. 9.3a. More number of attacks can be covered by enforcing a properly chosen subset of the security mechanisms, although resulting in heavy power utilization to support them. The cost of maintenance thus increase when lesser number of attacks are left uncovered. This observation conforms to our intuitive

Figure 9.3: (a) NSGA-II solutions in Scenario A. (b) Ratio of uncovered and covered damage cost for solutions obtained for Scenarios A and B. The line of shift shows the point beyond which the uncovered damage is more than the covered damage.

understanding. However, although no two solutions in the solution set (non-dominated front) are comparable in terms of their objective values, all solutions from the set do not fare equally well. Note that the number of attacks covered by a solution has no information in it about the total damage that it helped contain. This prompts us to identify the *line of shift* where the damage cost possible from uncovered attacks becomes more than that from covered ones.

A graphical illustration of the line of shift is shown in Fig. 9.3b. The figure shows the ratio of uncovered damage cost to covered damage cost. Any solution beyond the line of shift signifies a higher uncovered damage cost. Observe that a substantial number of solutions can exist beyond this line. If a decision maker's solution of choice lies beyond the line of shift, it is advisable that the process of security provisioning be rethought.

In terms of problem formulation, Problem 9.2 takes a damage-centric view of security and explicitly considers the total uncovered potential damage cost as an objective. Interestingly, this formulation can result in solutions with a lower uncovered to covered damage ratio for a given number of attacks left uncovered (Fig. 9.3b). A lower ratio indicates that the fraction of damages covered is much more than that uncovered. Hence, a Problem 9.2 solution is better than a Problem 9.1 solution since it gives the added benefit of having a lower uncovered to covered damage ratio. In this sense, solving Problem 9.2 can be a better approach even when the view of security is attack-centric.

Figure 9.4: NSGA-II solutions in Scenario B. (a) Solutions when maintenance cost of the devices are comparable. (b) Solutions when some devices have comparatively higher maintenance cost.

Fig. 9.4a shows the trade-off solutions when the uncovered damage cost is considered as one of the objectives. The non-dominated front is concave in structure with three identifiable regions of interest. Type I and Type III regions correspond to solutions where one of the objectives has a faster rate of decay than the other. From a cost of maintenance point of view, the trade-off nature in these regions signify that a decision maker can generate better outcome in one objective without much degradation on the other. This is quite difficult to perceive without having a global view of the interaction present between the two cost measures. The choice of a solution in the Type II region signify a good balance between the two cost factors. However, the number of solutions lying in each of these regions can vary significantly. Fig. 9.4b shows the same non-dominated front when certain devices have a much higher cost of maintenance compared to others. Observe that the Type I and Type III regions become more prominent in this front. This gives a decision maker better avenues to argue the selection of a solution biased towards a particular objective. Further, often solutions appear as part of a disconnected region of the non-dominated front (Fig. 9.4b (inset)). Such regions can be of special interest to a decision maker since disconnected solutions indicate that a change can be obtained in one objective by sacrificing a negligible value in the other.

The power capacity of a device restricts the usage of all possible subsets of the security

Figure 9.5: (a) NSGA-II solutions in Scenario C. Inset figure shows the sensitivity of a solution to security mechanism failures. (b) Solutions for failure radius = 1.

mechanisms in the device. Fig. 9.5a illustrates how the non-dominated front from Fig. 9.4a changes when the feasibility constraints are considered. The entire non-dominated front shifts to the right and clearly marks a region where no solution can be obtained. This in turn indicates the unavoidable damage cost that remains in the workflow. It is important that a business entity investing in a pervasive setup is aware of this residual cost in the system. This cost provides a preliminary risk estimate which, in the worst case, can become a matter of concern. If the unavoidable potential damage is too high, the setup will be running under a high risk of collapse.

The next step to the analysis involves the sensitivity of the solutions towards failure. The robustness analysis of a solution in Fig. 9.5a (inset) indicates that the uncovered potential damage cost can increase considerably for a failure radius of only 1. At this point, a decision maker can perform such analysis on every solution of interest and choose a feasible one. However, such an analysis is cumbersome and no control is possible on the actual amount of increase in the cost that an organization can sustain in the event of failure. Problem 9.4 alleviates this situation with a robust formulation of Problem 9.3.

Fig. 9.5b shows the robust solutions obtained for varying levels of acceptable cost increase. The increase in the potential damage cost stay within this level in the event of a failure of at most one security mechanism. Depending on the nature of the problem, obtaining solutions with small values of D may not be possible at all. Thus, obtaining a

certain level of robustness for a given level of security is not always feasible. However, there could be areas where experimenting with different values of D can be beneficial in understanding how the cost of maintenance changes with changing levels of robustness. As is seen in this example, the increase in the cost of maintenance is much higher when moving from a robustness level of $D = 30$ to 20 than moving from $D = 50$ to 30.

9.7 Conclusions

In this chapter, we address the problem of optimal security provisioning in pervasive environments under the presence of energy constraints. We adopt a workflow model to represent the different contexts under which a communication is established between two devices. We provide a formal statement of the problem of security provisioning and define a series of multi-objective optimization problems to understand the trade-offs involved between the cost of maintenance and the security of a pervasive setup.

Our analysis reveals important parameters that a business entity should be aware of before investing in the setup. First, the definition of "security" in the formulated problems plays an important role. Often, an attack-centric view of security is not enough and emphasis must be paid rather to a damage-centric view. Good solutions protecting against more attacks do not necessarily protect higher asset values. Also, the distribution of these solutions on the objective space provide invaluable clues to a decision maker on the amount of security gains possible across different levels of cost. The presence of energy constraints results in an unavoidable potential damage always residual in the system, early estimates on which can help the business entity invest better in risk mitigation strategies. Risk estimates also depend on the robustness of a chosen solution. Our robust formulation enables one to control the changes that can occur in the event of security mechanism failure and explore the costs involved.

We acknowledge that the presented work involves various other areas of research that require equal attention. The modeling of the different cost factors is a crucial aspect without which optimization formulations are difficult to transition to the real world. We can also explore the possibility of modifying the optimization framework to work on workflow models that can be broken down into sub-workflows for scalability.

CHAPTER 10

Security Hardening on Bayesian Attack Graphs

Traditionally, information security planning and management for an organization begins with risk assessment that determines threats to critical resources and the corresponding loss expectancy. A number of researchers have proposed risk assessment methods by building security models of network systems, using paradigms like attack graphs [10, 94, 141, 156, 166] and attack trees [43, 131, 145, 155], and then finding attack paths in these models to determine scenarios that could lead to damage. However, a majority of these models fail to consider the attacker's capabilities and, consequently, the likelihood of a particular attack being executed. Without these considerations, threats and their impact can be easily misjudged.

To alleviate such drawbacks, Dantu et al. [42] propose a probabilistic model to assess network risks. They model network vulnerabilities using attack graphs and applied Bayesian logic to perform risk analysis. In a similar effort, Liu and Man [115] use Bayesian networks to model potential attack paths in a system, and develop algorithms to compute an optimal subset of attack paths based on background knowledge of attackers and attack mechanisms. In both Dantu et al.'s and Liu and Man's works, nodes in the attack graph are assigned a probability value that describes the likelihood of attack on a node. They compute the likelihood of system compromise by chaining Bayesian belief rules on top of the assigned probabilities. The organizational risk is then computed as the product of

the likelihood of system compromise and the value of expected loss. The major problem with both these works is that they do not specify how the conditional probability value of an attack on each node is computed. Further, these works do not consider the problem of optimal risk mitigation.

System administrators are often interested in assessing the risk to their systems and determining the best possible way to defend their network in terms of an enumerated set of hardening options. Risk assessment methods such as those discussed earlier have been adopted by researchers to determine a set of potential safeguards, and related security control installation costs. There is however a dynamic aspect to the security planning process as well. For every attack, there is a certain probability of occurrence that can change during the life time of a system depending on what the contributing factors for the attack are and how they are changing. During run time, the system administrator may need to revise her decision based on such emerging security conditions. The attack tree model discussed in Section 8.3 does not allow such dynamic security planning.

Frigault et al. propose Bayesian network based attack graphs to incorporate temporal factors, such as availability of exploit codes or security patches, in order to facilitate a dynamic security planning process [71]. We can extend such concepts by encompassing the influence of ongoing attack incidents as well. In particular, we can use forward and backward probability inference techniques to assess the dynamic network security risk, given one or more nodes have been compromised by ongoing attacks.

Collectively, this chapter attempts to address the limitations highlighted above. First, we propose an alternative method of security risk assessment that we call *Bayesian Attack Graph*s (BAGs). In particular, we adapt the notion of Bayesian belief networks so as to encode the contribution of different security conditions during system compromise. Our model incorporates the usual cause-consequence relationships between different network states (as in attack graphs and attack trees) and, in addition, takes into account the likelihoods of exploiting such relationships. Second, we propose a method to estimate an organization's risk from different vulnerability exploitations based on the metrics defined in the Common Vulnerability Scoring System (CVSS) [154]. CVSS is designed to be an open and standardized method to rate IT vulnerabilities based on their base, temporal

and environmental properties. We also develop a model to quantify the expected return on investment based on a user specified cost model and likelihoods of system compromise. Third, we model the risk mitigation stage as a discrete reasoning problem and propose a genetic algorithm to solve it. The algorithm can identify optimal mitigation plans in the context of both single and multi-objective analysis. We also discuss how the above contributions collectively provide a platform for static and dynamic analysis of risks in networked systems.

The rest of the chapter is organized as follows. The test network used to illustrate our problem formulation and solution is described in Section 10.1. Section 10.2 presents the formalism for a Bayesian Attack Graph model. The likelihood estimation method in static and dynamic scenarios is discussed in Section 10.3. The risk mitigation process along with the expected cost computations is presented in Section 10.4. Empirical results are presented in Section 10.5. Finally, we conclude Section 10.6.

10.1 A Test Network

Fig. 10.1 depicts the test network used in this study. The network consists of eight hosts located within two sub-nets. A DMZ tri-homed firewall is installed with preset policies to ensure that the Web server, Mail server and the DNS server, located in the DMZ network, are separated from the local network so that the damage will only be limited to the DMZ zone if one of these servers is compromised. The firewall has a strong set of policies (shown in the inset table) to prevent remote access to the internal hosts. In particular, all machines in the DMZ zone passively receive service requests and only respond to the sender as needed. However, in order to accommodate Web service's transactions, the Web server is allowed to send SQL queries to the SQL server located in the trusted zone on a designated channel. Local machines are located behind a NAT firewall so that all communications to external parties are delivered through the Gateway server. In addition, all local desktops, including the administrator machine, have remote desktop enabled to facilitate remote operations for company employees working from remote sites. The remote connections are monitored by SSHD installed in the Gateway server.

From Host	To Host	Protocol (Port#)
0.0.0.0	Gateway server	SSHD (22)
	Mail server	IMAP (143)
		SMTP (25)
	Web server	HTTP (80)
Web server	SQL server	SQL (1433)
Gateway server	Local machines	Basic network protocols
	Root machine	Basic network protocols
Local machines	Gateway server	Basic network protocols
and root machine	Mail server	IMAP (143)
		SMTP (25)
	SQL server	SQL (1433)
	Web server	HTTP (80)
	DNS server	DNS (53 & 1024)

Figure 10.1: Test-bed network model.

A list of initial vulnerabilities/attack templates in this test network is listed in Table 10.1. Further scrutiny of this initial list using a vulnerability database reveals that eight malicious outcomes are possible in this network. However, the list of vulnerabilities alone cannot suggest the course of actions that lead to these outcomes, or accurately assess the casualty of each outcome, as it may have involved other damages along the way. These vulnerabilities produce more than 20 attack scenarios with different outcomes, ranging from information leakage to system compromise. Moreover, two of these scenarios use machines in the DMZ zone to compromise a local machine in the trusted zone.

10.2   Modeling Network Attacks

We use a Bayesian belief network to model network vulnerabilities. We extend the notion of Bayesian networks as presented by Liu and Man [115] to encode the contributions of different security conditions during a system compromise. We term such a Bayesian

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Table 10.1: Initial list of vulnerabilities in test network.

Host	Vulnerability	CVE#	Attack Template
Local desktops (10.0.0.1-127)	Remote login	CA 1996-83	remote-2-user
	LICQ Buffer Overflow (BOF)	CVE 2001-0439	remote-2-user
	MS Video ActiveX Stack BOF	CVE 2009-0015	remote-2-root
Admin machine (10.0.0.128)	MS SMV service Stack BOF	CVE 2008-4050	local-2-root
Gateway server (196.216.0.128)	OpenSSL uses predicable random	CVE 2008-0166	information leakage
	Heap corruption in OpenSSH	CVE 2003-0693	local-2-root
	Improper cookies handler in OpenSSH	CVE 2007-4752	authentication bypass
SQL Server (196.216.0.130)	SQL Injection	CVE 2008-5416	remote-2-root
Mail Server (196.216.0.19)	Remote code execution in SMTP	CVE 2004-0840	remote-2-root
	Error message information leakage	CVE 2008-3060	account information theft
	Squid port scan vulnerability	CVE 2001-1030	information leakage
DNS Server (196.216.0.20)	DNS Cache Poisoning	CVE 2008-1447	integrity
Web Server (196.216.0.20)	IIS vulnerability in WebDAV service	CVE 2009-1535	remote-2-local
			authentication bypass

network as a Bayesian Attack Graph (BAG).

Using the notion of an attribute-template as defined in Section 8.3, we define an *attribute* in a BAG as follows.

Definition 10.1 ATTRIBUTE. An attribute is a Bernoulli random variable representing the state of an instance of an attribute-template.

An attribute S is therefore associated with a state – *True* ($S = 1/T$) or *False* ($S = 0/F$) – and a probability $Pr(S)$. The state signifies the truth value of the proposition underlined by the instance of the attribute template. For example, the instance S :"*user access on Local machine*" is an attribute when associated with a truth value signifying whether an attacker has user access on the local machine. We shall also use the term "compromised" to indicate the *true* (or $S = 1$) state of an attribute. Further, $Pr(S)$ is the probability of the attribute being in state $S = 1$. Consequently, $Pr(\neg S) = 1 - Pr(S)$ is the probability of the state being $S = 0$. The success or failure of an attacker reaching its goal depends mostly on the states of the attributes in a network. It also lays the foundations for a security manager to analyze the effects of forcing some attributes to the *false* state using security measures. We formally define a BAG to capture the cause-consequence relationships between such attributes.

Definition 10.2 ATOMIC ATTACK. Let S be a set of attributes. We define $\mathcal{A}$, a conditional dependency between a pair of attributes, as a mapping $\mathcal{A} : S \times S \rightarrow [0,1]$. Then, given $S_{pre}, S_{post} \in S$, $a : S_{pre} \mapsto S_{post}$ is called an atomic attack if

1. $S_{pre} \neq S_{post}$,

2. given $S_{pre} = 1$, we also have $S_{post} = 1$ with probability $\mathcal{A}(S_{pre}, S_{post}) > 0$, and

3. $\nexists S_1, \ldots, S_j \in S - \{S_{pre}, S_{post}\}$ such that $\mathcal{A}(S_{pre}, S_1) > 0, \mathcal{A}(S_1, S_2) > 0, \ldots$, and $\mathcal{A}(S_j, S_{post}) > 0$.

Therefore, an atomic attack allows an attacker to compromise the attribute S_{post} from S_{pre} with a non-zero probability of success. Although, given a compromised attribute, another attribute can be compromised with positive probability using a chain of other

attributes, the third condition in the definition does not allow such instances to be considered as atomic attacks. Instead, each step in such a chain is an atomic attack. Informally, an attack is associated with a vulnerability exploitation, denoted by e_i, which takes the attacker from one network state (S_{pre}) to another (S_{post}). The probability of an exploitation, $Pr(e_i)$, states the ease with which an attacker can perform the exploitation. Hence, we say that $\mathcal{A}(S_{pre}, S_{post}) = Pr(e_i)$, and S_{pre} and S_{post} are respectively called a *precondition* and *postcondition* of the attack a, denoted by *pre(a)* and *post(a)* respectively.

An attack relates the states of two different attributes so as to embed a cause-consequence relationship between the two. For example, for the attributes S_{pre} ="sshd BOF vulnerability on machine A" and S_{post} ="root access privilege on machine A", the attack $S_{pre} \mapsto S_{post}$ is associated with the e_i ="sshd buffer overflow" exploit. Using this exploit, an attacker can achieve root privilege on a machine provided the machine has the sshd BOF vulnerability. $\mathcal{A}(S_{pre}, S_{post})$ is the probability of success of the exploit, i.e. $\mathcal{A}(S_{pre}, S_{post}) = Pr(e_i)$.

Definition 10.3 BAYESIAN ATTACK GRAPH. Let S be a set of attributes and A be the set of atomic attacks defined on S. A Bayesian Attack Graph is a tuple $BAG = (S, \tau, \varepsilon, \mathcal{P})$, where

1. $S = N_{internal} \cup N_{external} \cup N_{terminal}$. $N_{external}$ denotes the set of attributes S_i for which $\nexists a \in A | S_i = post(a)$. $N_{internal}$ denotes the set of attributes S_j for which $\exists a_1, a_2 \in A | [S_j = pre(a_1)$ and $S_j = post(a_2)]$. $N_{terminal}$ denotes the set of attributes S_k for which $\nexists a \in A | S_k = pre(a)$.

2. $\tau \subseteq S \times S$. An ordered pair $(S_{pre}, S_{post}) \in \tau$ if $S_{pre} \mapsto S_{post} \in A$. Further, for $S_i \in S$, the set $Pa[S_i] = \{S_j \in S | (S_j, S_i) \in \tau\}$ is called the parent set of S_i.

3. ε is a set of decomposition tuples of the form $\langle S_j, d_j \rangle$ defined for all $S_j \in N_{internal} \cup N_{terminal}$ and $d_j \in \{AND, OR\}$. d_j is AND if $S_j = 1 \Rightarrow \forall S_i \in Pa[S_j], S_i = 1$. d_j is OR if $S_j = 1 \Rightarrow \exists S_i \in Pa[S_j], S_i = 1$.

4. $\mathcal{P}$ is a set of discrete conditional probability distribution functions. Each attribute $S_j \in N_{internal} \cup N_{terminal}$ has a discrete local conditional probability distribution (LCPD) representing the values of $Pr(S_j \mid Pa[S_j])$.

Figure 10.2: BAG of test network with unconditional probabilities.

Fig. 10.2 shows the BAG for our test network. We use an in-house tool to generate such BAGs. The tool takes as input an initial vulnerability table, generated by a vulnerability scanner, and the network topology (currently provided manually to the tool). Using a sequence of SQL queries on a vulnerability exposure database, the tool creates consequence attributes for the graph until no further implications can be derived. The tool addresses circular dependencies by assuming that the preconditions of an attack are never invalidated by successfully executing another attack. This is also known as the monotonicity assumption [10]. Circular dependencies also do not influence the success probabilities of attacks in subsequent analysis.

The BAG in Fig. 10.2 depicts a clear picture of 20 different attack scenarios. Each node is a Bernoulli random variable (S_i) representing the state variable of the attribute. The set $N_{external}$ represents the entry points of the graph. These nodes reflect an attacker's

capability as discovered in a threat-source model. $N_{terminal}$ resemble the end points in the graph. These nodes reflect casualty at the end of each attack scenario. The set of ordered pair, τ, reflects the edges in the graph. The existence of an edge between two nodes imply that there is a causal dependency between their states, signified by the decomposition at each node. AND-decomposition signifies that the compromised state of a node implies that all nodes in its parent set have also been compromised. Similarly, OR-decomposition signifies that at least one parent node is in the *true* state. Note that these decompositions are uni-directional. For instance, under AND-decomposition, compromising all nodes in the parent set does not necessarily imply the node itself has been compromised. This is because the attacks relating the node with its parents can have varying levels of difficulty, or in other words, different probabilities of success. Hence, although the preconditions of the attacks have been met, there can still be a non-zero probability that the attacker is unable to carry out all the exploits successfully. The existence of this probability is what primarily differentiates a BAG from a classical attack graph. The probabilities are captured in the local conditional probability distribution of the node. The LCPD is a set of probability values specifying the chances of the node being compromised, given different combination of states of its parents.

Definition 10.4 LOCAL CONDITIONAL PROBABILITY DISTRIBUTION. Let $(S, \tau, \varepsilon, \mathcal{P})$ be a BAG and $S_j \in N_{internal} \cup N_{terminal}$. For $S_i \in Pa[S_j]$, let e_i be the vulnerability exploitation associated with the attack $S_i \mapsto S_j$. A local conditional probability distribution (LCPD) function of S_j, mathematically equivalent to $Pr(S_j \mid Pa[S_j])$, is defined as follows.

1. $d_j = AND$

$$Pr(S_j \mid Pa[S_j]) = \begin{cases} 0, & \exists S_i \in Pa[S_j] \mid S_i = 0 \\ Pr(\bigcap_{S_i=1} e_i), & otherwise \end{cases} \tag{10.1}$$

2. $d_j = OR$

$$Pr(S_j \mid Pa[S_j]) = \begin{cases} 0, & \forall S_i \in Pa[S_j], S_i = 0 \\ Pr(\bigcup_{S_i=1} e_i), & otherwise \end{cases} \tag{10.2}$$

To compute the local conditional probabilities when multiple exploits are involved, we proceed as follows. For AND-decomposition, each vulnerability exploitation is a distinct

event. The chance of compromising the target node depends on the success of each individual exploit. Therefore, we use the product rule in probability to derive $Pr(\bigcap_{S_i=1} e_i)$ as

$$Pr(\bigcap_{S_i=1} e_i) = \prod_{S_i=1} Pr(e_i). \tag{10.3}$$

For OR-decomposition, Liu et al. observed that the joint probability is equivalent to the noisy-OR operator [115], given as

$$Pr(\bigcup_{S_i=1} e_i) = 1 - \prod_{S_i=1} [1 - Pr(e_i)]. \tag{10.4}$$

10.3 Security Risk Assessment with BAG

Security risk management consists of threat analysis, risk assessment, loss expectancy, potential safeguards and risk mitigation analysis. A BAG positions itself between threat analysis and risk assessment. Threat sources and the list of initial vulnerabilities are used to build the BAG threat model. Once the graph is built, the administrator can expect better results in risk assessment and risk mitigation analysis as follows.

Static Risk Assessment: Risk assessment begins with the identification of system characteristics, potential threat sources and attacker capabilities. Threat sources are represented as the external nodes in a BAG, along with their impact on other network attributes. One set of attributes act as preconditions to an exploit, which when successfully executed by an attacker, can make the network state favorable for subsequent exploits. Estimating the amount of risk at each node therefore requires some judgment on attacker capabilities. Often this judgment is indirectly stated as the system administrator's subjective belief on the likelihood of a threat source becoming active and the difficulty of an exploit. The former is represented by the probabilities $Pr(S_i)$ for all $S_i \in N_{external}$, also called the *prior probabilities*, and is subjectively assigned by the administrator. The latter is incorporated into an internal node's LCPD. Thereafter, given the prior probability values and the LCPDs, we can compute the *unconditional probability $Pr(S_j)$* for any node $S_j \in N_{internal} \cup N_{terminal}$.

These risk estimates can be used to help locate weak spots in the system design and operations.

Dynamic Risk Assessment: A deployed system may experience first hand attack incidents during its life cycle. Formally, an attack incident is evidence that an attribute is in the *true* state. A security administrator may then want to investigate how these incidents impact the risk estimates initially derived solely based on subjective beliefs. Knowledge about attack incidents is therefore used to update the probabilities using the Bayesian inference techniques of forward and backward propagation. Forward propagation updates the probability on successor attributes that are directly influenced by the evidences. Backward propagation corrects/adjusts the initial hypothesis on all prior attributes. Thereafter, the *posterior probabilities* (updated unconditional probabilities) reflect the likelihoods of other potential outcomes under the light of detected events.

Risk Mitigation Analysis: Risk assessment paves the way for efficient decision making targeted at countering risks either in a proactive or reactive manner. Given a set of security measures (e.g. firewall, access control policy, cryptography, etc.), we can design the security plan which is the most resource efficient in terms of reducing risk levels in the system. This can be done before the deployment (static mitigation) or in response to attack incidents (dynamic mitigation).

10.3.1 Probability of vulnerability exploitation

In order to compute the local conditional probability distribution (LCPD) of an attribute, the administrator needs to estimate the probability of success when an attacker exploits a known vulnerability exploitation. We propose a method to estimate this attack likelihood using publicly available risk exposure data sources. In particular, we are interested in deriving attack likelihoods using the metrics defined in NIST's Common Vulnerability Scoring System (CVSS) [154].

A CVSS score is a decimal number on a scale of 0 to 10. It is composed of three groups – *base*, *temporal* and *environmental*. The base metrics quantify the intrinsic characteristics of a vulnerability with two sub-scores – (i) the exploitability sub-score, composed of the

Table 10.2: CVSS attributes used for estimation of attack likelihood.

CVSS metrics group	CVSS attributes	category	score
base metrics	access vector(B_AV)	local(L)	0.395
		adjacent network(A)	0.646
		network(N)	1.0
	attack complexity(B_AC)	high(H)	0.35
		medium(M)	0.61
		low(L)	0.71
	authentication instance(B_AU)	multiple(M)	0.45
		single(S)	0.56
		none(N)	0.704
temporal metrics	exploitability (tools & techniques) (T_E)	unproved(U)	0.85
		proof-of-concept(POC)	0.9
		functional(F)	0.95
		high(H)	1.0
	remediation level(T_RL)	official fix(OF)	0.87
		temporary fix(TF)	0.90
		workaround(W)	0.95
		unavailable(U)	1.0
	report confidence (T_RC)	unconfirmed(UC)	0.90
		uncorroborative(UR)	0.95
		confirmed(C)	1.0

access vector (B_AV), access complexity (B_AC) and authentication instances (B_AU), and (ii) the impact sub-score, expressing the potential damage on confidentiality (B_C), integrity (B_I) and availability (B_A). The temporal metrics quantify dynamic aspects of a vulnerability on the environment around the organization. These metrics take into account the availability of exploitable tools and techniques (T_E), remediation level (T_RL) and report confidence (T_RC). The environmental metrics quantify two aspects of impact that are dependent on the environment surrounding the organization. More details on CVSS metrics and their scoring computation can be found in the CVSS guide [154]. In this study, we are interested in likelihood estimation and hence the impact sub-score and environmental metrics are ignored in the analysis. A summary of the metrics used here is shown in Table 10.2. We refine Houmb's Misuse Frequency model [86] to estimate the probability of success in vulnerability exploitation.

Given the vulnerability exposure information (CVSS attributes), the probability of success $Pr(e_i)$ while executing a given vulnerability exploitation e_i is computed by the

following equations.

$$Pr(e_i) = (1 - \mu)MF_{init} + \mu MF_{uFac}, \text{ where} \tag{10.5}$$

$$0 \leq \mu \leq 0.5$$
$$MF_{init} = \frac{B\_AV \times B\_AC \times B\_AU}{0.49984}$$
$$MF_{uFac} = T\_E \times T\_RL \times T\_RC.$$

Note that the combination of various CVSS metrics to estimate the exploit probability has also been used earlier [71]. Our equation distinctly differs from this approach in the treatment of the impact sub-scores. We intentionally remove all impact metrics since the probability of a successful exploit is not usually related to its impact. In addition, the perceived impact of a successful exploit differs from organization to organization. The formulation in [71] directly uses the base score (includes exploitability and impact sub-scores) while our formulation only uses the exploitability sub-score (MF_{init}).

The constant μ represents the evaluator's preference weight on temporal knowledge of the exploitation. In the case where the vulnerability exploitation is unknown to the evaluator, the estimation should rely on the base score by setting μ to zero. In the case where the evaluator or organization has experienced the vulnerability exploitation, or there is an ongoing concern about the exploitation, the evaluator may set the value of μ to a specific value. However, we bound μ to a maximum value of 0.5 in order to restrict likelihood estimates based solely on temporal factors. Nonetheless, temporal metrics help capture the uncertainties in relatively new exploits. For instance, at the time of writing this dissertation, CVE announced a vulnerability in Acrobat Reader(VU#905281) where the only workaround is to disable JavaScript in Acrobat Reader. In such a case, temporal metrics often influence security decisions because of immediate needs. We design μ to capture such an aspect.

Our empirical estimation in Eq. (10.5) preserves the CVSS design characteristics and extends the range of possible values in Houmb's model from $[0.53, 0.83]$ to $[0.12, 1.00]$.

10.3.2 Local conditional probability distributions

Refer to the BAG in Fig. 10.3. Nodes *A:"root/FTP server"*, *B:"Matu FTP BOF"* and *C:"remote BOF on ssh"* are internal attributes, while node *D:"remote attacker"* is an external

Figure 10.3: Simple BAG illustrating probability computations.

attribute. A is the successor of B and C which in turn are successors of D. The values on the edges reflect the probability of success of the associated vulnerability exploitation, computed by following the procedure described in the previous section. We begin by assigning a prior probability of $Pr(D) = 0.7$ to the external attribute D. This probability represents the administrator's subjective belief on the chances of a remote attack. For the nodes A, B and C, we calculate LCPDs by the equations previously defined in Def. 10.4. For example, for node A, there are 2^2 marginal cases given the two parents B and C. The decomposition at the node dictates the rule to follow while computing the local conditional probability for each case.

10.3.3 Unconditional probability to assess security risk

Once the LCPDs have been assigned to all attributes in the BAG, we can merge the marginal cases at each node to obtain the unconditional probability at the node. This is commonly known as *marginalization*. Further, given a set of Bernoulli random variables $S = \{S_1, ..., S_n\}$ in a Bayesian belief network, the joint probability of all the variables is given by the *chain rule* as

$$Pr(S_1, ..., S_n) = \prod_{i=1}^{n} Pr(S_i \mid Pa[S_i]). \tag{10.6}$$

In Fig. 10.3, the unconditional probability at node A is derived as the joint probability of A along with all nodes that influence its outcome, which is essentially all ancestors of A. Hence we have,

$$
\begin{aligned}
Pr(A) \quad &= \quad Pr(A,B,C,D) \\
&= \quad Pr(A \mid B,C) \times Pr(B \mid D) \times Pr(C \mid D) \times Pr(D) \qquad \text{(chain rule)} \\
&= \quad \sum_{B,C,D \in \{T,F\}} [Pr(A \mid B,C) \times Pr(B \mid D) \times Pr(C \mid D) \times Pr(D)] \quad \text{(marginalization)} \\
&= \quad (1.0 \times 0.85 \times 0.7 \times 0.7)_{TTT} + (0.65 \times 0.85 \times 0.3 \times 0.7)_{TFT} + \\
&\quad\ \ \, (1.0 \times 0.15 \times 0.7 \times 0.7)_{FTT} \qquad\qquad\qquad\quad \text{(after simplification)} \\
&= \quad 0.6060 \approx 61\%.
\end{aligned}
$$

Similarly, unconditional probabilities at nodes B and C can be computed by considering the sub-network rooted at the corresponding nodes. The unconditional probabilities are shown under the LCPD table of each node. Fig. 10.2 shows the unconditional probabilities of the nodes in our test network. It exposes the weak spots of the system where the likelihood of attack is higher than others. The security administrator can use this threat model to prioritize risk and derive an effective security hardening plan so as to reduce the risk to a certain level (e.g. $< 50\%$) before deploying the system. The model can also be used to assess what-if scenarios, for e.g. while deploying new machines, services, or operations of interest.

10.3.4 Posterior probability with attack evidence

The BAG can also be used to address dynamic aspects of the security planning process. Every network state has a certain probability of occurrence. This probability can change during the life time of the system due to emerging security conditions, changes in contributing factors or the occurrence of attack incidents. The BAG can then be used to calculate the posterior probabilities in order to evaluate the risk from such emerging conditions.

Let $S = \{S_1, ..., S_n\}$ be the set of attributes in a BAG and $E = \{S'_1, ..., S'_m\} \subset S$ be a set of attributes where some evidence of exploit have been observed. We can say that attributes in E are in the *true* state, i.e. $S'_i = 1$ for all $S'_i \in E$. Let $S_j \in S - E$ be an attribute whose posterior probability has to be determined. The probability we are interested in is $Pr(S_j \mid E)$ and can be obtained by using the *Bayes Theorem*, given as

$$
Pr(S_j \mid E) = Pr(E \mid S_j) \times Pr(S_j)/Pr(E). \tag{10.7}
$$

$Pr(E \mid S_j)$ is the conditional probability of joint occurrence of $S'_1, ..., S'_m$ given the states

of S_j. $Pr(E)$ and $Pr(S_j)$ are the prior unconditional probability values of the corresponding attributes. Since evidence attributes in E are mutually independent, $Pr(E \mid S_j) = \prod_i Pr(S_i' \mid S_j)$ and $Pr(E) = \prod_i Pr(S_i')$. For example, in Fig. 10.3, assume that the system administrator detects an attack incident on A (attacker compromises FTP server). The posterior probability of C is then computed as follows.

$$
\begin{aligned}
Pr(C \mid A) &= Pr(A \mid C)Pr(C)/Pr(A) \\
&= 0.81, \text{ where} \\
Pr(A \mid C) &= \sum_{B \in \{T,F\}} [Pr(A \mid B, C = T)Pr(B)] \\
&= (1.0 \times 0.6)_T + (1.0 \times 0.4)_F \\
&= 1.0 \\
Pr(A) &= 0.61 \\
Pr(C) &= 0.49
\end{aligned}
$$

Similarly, the posterior probability at node B can be computed in the same manner. Note that the unconditional probability of node C was originally 0.49. After taking into account the attack incident at node A, the posterior probability becomes 0.81. Further, computation of posterior probabilities for successor nodes of A (forward propagation) remain the same as described in the previous sub-section, with the change that the LCPDs at those nodes only account for the $A = 1$ case while marginalization. In this manner, the security administrator can revise the probability of occurrence of every node of the graph in response to an emerging attack incident. Fig. 10.4 shows the posterior probabilities in response to two hypothetical evidences (denoted by the label Ⓔ) in the Mail server of our test network. Note that the parent ("*root access @ 196.216.0.19*") of the evidence node "*squid port scan*" has a posterior probability of less than 1.0. Ideally, given the evidence that the port scan has been executed, the attacker must have had root access on the machine. Hence, the parent node should also have an updated probability of 1.0. However, this inference assumes that the squid port scan is only executable after gaining root access on the machine. The system administrator may decide to relax such an assumption in order to account for uncertainties (e.g. zero-day attacks), achieved by replacing the zero values in Def. 10.4 with non-zero values. Such a relaxation will reduce the impact of the evidence nodes on their parents.

As can be seen in Fig. 10.4, most of the unconditional probabilities increase after the attack incidents, but not at the same rate. It is possible to have nodes with decreased

Figure 10.4: Posterior probabilities in test network after attack incidents (marked by Ⓔ).

probabilities as well. In this specific scenario, there is a significant increase in the chance that the administrator machine is targeted by an attacker. This observation shows that the attacker is likely to execute an attack to compromise the root machine. Hence, sufficient measures should be taken to protect it. Moreover, it is also possible that the mitigation plan designed earlier in static analysis may no longer be appropriate under the light of the emerging events. We will formally address this problem in the next section.

10.4   Security Risk Mitigation with BAG

Although many researchers have studied risk assessment schemes, including the NIST, the methodologies used to estimate loss varies from organization to organization. Loss can be measured in terms of monetary units, relative magnitudes [11, 19, 108, 160] or

Table 10.3: Expanded LCPD of node A (in Fig. 10.3) under the presence of security control M_0.

B	C	Pr(A)	Pr(¬A)
1	1	1.00	0.00
1	0	0.65	0.35
0	1	1.00	0.00
0	0	0.00	0.00

$\longrightarrow$

B	C	M_0	Pr(A)	Pr(¬A)
1	1	1	0.50	0.50
1	1	0	1.00	0.00
1	0	1	0.65	0.35
1	0	0	0.65	0.35
0	1	1	0.75	0.25
0	1	0	1.00	0.00
0	0	1	0.00	1.00
0	0	0	0.00	1.00

multi-units [29, 30, 51]. In a BAG, the security manager can choose to evaluate the risks by considering an expected loss/gain quantity. The expected loss/gain is computed from organization specific factors such as potential loss or gain associated with an attribute's states. It usually reflects the impact of attack likelihoods on the economic turnout of an organization. We will describe this scheme later in the section. We begin with the notion of a security control in the context of the BAG.

Definition 10.5 SECURITY CONTROL. Given a BAG $(S, \tau, \varepsilon, \mathcal{P})$, a Bernoulli random variable M_i is a security control if $\exists S_j \in N_{internal} \cup N_{external}$ such that $Pr(S_j \mid Pa[S_j], M_i = T) < Pr(S_j \mid Pa[S_j], M_i = F)$ for at least one assignment of states to $Pa[S_j]$. Further, $Pr(M_i) = 1.0$ if $M_i = T$; otherwise zero.

In other words, a security control is a preventive measure that minimizes or eliminates the likelihood of attack on one or more attributes so as to prevent an attacker from reaching its goal. We define the security measure as a Bernoulli random variable with the *true* state signifying that the control is enforced and *false* signifying that the control is known but not enforced. Enforcement of a control directly influences the LCPD of the associated attribute and indirectly impacts the unconditional probabilities of other attributes. For example, the probability of the node A in Fig. 10.3 is initially $Pr(A \mid B, C)$. Assume that a security measure M_0:*"local access control"* can influence the outcome at A. The probability distribution therefore becomes $Pr(A \mid B, C, M_0)$ and the LCPD of the node is hypothetically expanded as shown in Table 10.3. The probabilities when $M_0 = 0$ are directly taken from the original LCPD. However, probabilities for $M_0 = 1$ are assigned based on certain

subjective belief on the security measure's capacity to prevent the attribute's compromise. The modified LCPDs are used to compute the unconditional probability of nodes in the graph. It is not difficult to see that the unconditional probability of a node (and its successors) influenced by a control will reduce when the control is enforced. Note that, by definition, the unconditional probability of the control itself is 1.0 if its state is *true*.

Definition 10.6 SECURITY MITIGATION PLAN. Given set $M = \{M_1, \ldots, M_m\}$ of m security controls, a security mitigation plan is represented by a boolean vector $\vec{T} = (T_1, T_2, \ldots, T_m)$ where $M_i = T_i$.

Therefore, the mitigation plan is a specification of which controls have been chosen for enforcement as part of the hardening process. Further, for a given control M_i, consider the set $\mathcal{I}$ of all $S_j \in N_{internal} \cup N_{terminal}$ such that $Pr(S_j \mid Pa[S_j], M_i = T) < Pr(S_j \mid Pa[S_j], M_i = F)$ (for some assignment of states to $Pa[S_j]$). Then, the subset $\{S_k \in \mathcal{I} \mid Pa[S_k] \cap \mathcal{I} = \phi\}$ is the coverage of M_i. With reference to Fig. 10.3, a control such as M_0:*"disconnect from Internet"* directly changes the probability $Pr(D)$ (equal to zero if $M_0 = 1$). This in effect changes the LCPD tables at nodes B, C and D. Therefore, the set $\mathcal{I}$ contains all four nodes for M_0. However, only node D is in the coverage of M_0 since, for all other nodes, one or more parent nodes are also present in $\mathcal{I}$. Intuitively, the coverage nodes are those whose LCPDs are directly affected by a security control, rather than by indirect inference. For a given security mitigation plan $\vec{T}$, we can define the plan coverage by collecting the coverage of each enforced control in the plan. Each control M_i also has an associated cost C_i of implementation, giving us the total plan cost as

$$SCC(\vec{T}) = \sum_{i=1}^{m}(T_i C_i). \tag{10.8}$$

10.4.1 Assessing security outcomes

When using a BAG, a better quantitative representation of the loss/gain is obtained by considering the expected loss/gain once a set of security measures have been implemented. Hence, we augment the BAG with a value signifying the amount of potential loss/gain at each node, and account for the security decision during evaluation.

Definition 10.7 AUGMENTED-BAYESIAN ATTACK GRAPH. Let $BAG = (S, \tau, \varepsilon, \mathcal{P})$ be a Bayesian attack graph. An augmented-Bayesian attack graph (augmented-BAG) $BAG_{aug} = BAG|(M, \gamma, V)$ is obtained by adding a node set M, edge set γ and by associating a value V_j to each $S_j \in S$, where

1. M is the set of security controls.

2. $\gamma \subseteq M \times S$. An ordered pair $(M_i, S_j) \in \gamma$ if S_j is in the coverage of M_i.

3. V_j is the expected loss/gain associated with the attribute $S_j \in S$.

The set M extends the BAG with additional nodes representing hardening measures. The set γ represents the new edges between the controls and attributes of the graph. A new edge is inserted if a control directly influences the state of an attribute. In this work, all external attributes are given a zero cost, i.e. $V_j = 0$ for all $S_j \in N_{external}$. The value associated with $S_j \in N_{internal} \cup N_{terminal}$ is computed as

$$V_j = \big[1 - Pr(S_j)\big] \times G_j - Pr(S_j) \times L_j, \tag{10.9}$$

where L_j is the potential loss representing the damage value that an organization might have to pay when the attribute S_j is compromised, G_j is the potential gain if S_j is not compromised and $Pr(S_j)$ is the unconditional probability of S_j. If there exists $(M_i, S_j) \in \gamma$, $Pr(S_j)$ is computed as a conditional probability $Pr(S_j \mid M_i)$ where the state of M_i depends on the security plan $\vec{T} = (T_i)$. The expected loss/gain w.r.t. the security plan $\vec{T}$, denoted by $LG(\vec{T})$, is computed as the cumulative sum of all node values, i.e.

$$LG(\vec{T}) = \sum_{S_j \in S} V_j. \tag{10.10}$$

A positive value of LG signifies gain, while a negative value signifies loss. Note that we do not assume any particular cost model in our formulations, both for control cost and loss/gain evaluation. The cost model is usually subjective to organizational policies and hence can differ from one institution to another. The cost factors considered here (security control cost, potential loss and potential gain) are standard quantities that any organization must be able to determine in order to perform risk analysis.

10.4.2 Assessing the security mitigation plan

In order to defend against the attacks possible, a security manager (as a decision maker) can choose to implement a variety of safeguard technologies, each of which comes with different cost and coverage. For example, to defend against the "*ftp/.rhost*" exploit, one might choose to apply a security patch, firewall, or simply disable the FTP service. Each choice of action has a different cost and different outcome. A security administrator has to assess the technologies and make a decision towards maximum resource utilization. The two objectives we consider in this study are the total security control cost and the expected loss/gain. The single-objective problem is the most likely approach to be taken by a decision maker. The problem is stated as follows.

Problem 10.1 The Single-Objective Optimization Problem (SOOP). Given an augmented-BAG $(S, \tau, \varepsilon, \mathcal{P}) \mid (M, \gamma, V)$, find a vector $\vec{T}^* = (T_i^*)$, $T_i^* \in \{0,1\}; 1 \leq i \leq |M|$, which maximizes the function $\alpha LG(\vec{T}^*) - \beta SCC(\vec{T}^*)$, where α and β are preference weights for the expected loss/gain and the total cost of security control respectively, $0 \leq \alpha, \beta \leq 1$ and $\alpha + \beta = 1$.

The method for assessing a security plan is as follows. First, the security analyst chooses a subset of security controls to construct a security plan $\vec{T}^*$. She then updates the unconditional probability of all attributes using the plan coverage information. She computes the expected loss/gain associated with every attribute $S_j \in S$ using Eq. (10.9). Finally, the total expected loss/gain of the entire graph is taken as an assessment of the security plan's outcome. The best security plan is the one that maximizes the function $\alpha LG(\vec{T}^*) - \beta SCC(\vec{T}^*)$.

The next level of sophistication is added by formulating the optimization as a multi-objective problem. The multi-objective approach alleviates the requirement to specify any weight parameters and hence a better global picture of the solutions can be obtained.

Problem 10.2 The Multi-Objective Optimization Problem (MOOP). Given an augmented-BAG $(S, \tau, \varepsilon, \mathcal{P}) \mid (M, \gamma, V)$, find a vector $\vec{T}^* = (T_i^*)$, $T_i^* \epsilon \{0,1\}; 1 \leq i \leq |M|$, which minimizes SCC and maximizes LG.

Figure 10.5: Augmented-BAG of test network with 13 security controls.

Therefore, using the Pareto-dominance concept, a security plan $\vec{T_1}$ is optimal if there is no other plan $\vec{T_2}$ such that

1. $SCC(\vec{T_2}) < SCC(\vec{T_1})$ and $LG(\vec{T_2}) \geq LG(\vec{T_1})$, or

2. $SCC(\vec{T_2}) \leq SCC(\vec{T_1})$ and $LG(\vec{T_2}) > LG(\vec{T_1})$.

For the BAG shown in Fig. 10.5, we have identified that 13 different security controls are possible. These controls are represented by an 'eclipse' in the figure. These security controls produce 2^{13} candidate security plans. A genetic algorithm based approach is presented next to search through these candidate plans in an efficient manner.

10.4.3   Genetic algorithm

Fig. 10.6 depicts the structure of the genetic algorithm designed for this study. The algorithm begins with a population P_0 of N randomly generated security plans. A gener-

Figure 10.6: Schematic of the genetic algorithm used to solve SOOP and MOOP.

ation index $t = 0, 1, \ldots, Gen_{MAX}$ keeps track of the number of iterations of the algorithm. Each iteration proceeds as follows. The *SCC* and *LG* values of every plan in P_t are calculated. $N/2$ plans are then selected from P_t to form a mating pool M_t. The selection process is different for SOOP and MOOP, and discussed later. An offspring population Q_t (containing $N/2$ plans) is generated from the mating pool by using the standard single-point binary crossover and mutation operators [75]. The process is then repeated with $P_{t+1} = Q_t \cup M_t$ until $t = Gen_{MAX}$.

10.4.3.1   Solving SOOP

The selection process to solve SOOP is based on the objective function $\alpha LG(\vec{T}) - \beta SCC(\vec{T})$ and uses the process of binary tournament. In this process, two plans are randomly selected (with replacement) from P_t and the one with the higher objective function value is inserted into the mating pool. This process is repeated until the mating pool is full. The solution to SOOP is the plan with the highest objective value across all iterations of the algorithm.

10.4.3.2   Solving MOOP

Simple objective value comparison is not possible in the presence of more than one objective function. Hence, a different selection scheme is required for MOOP. The scheme used here is based on non-dominance ranking. Under this process, all non-dominated solutions in P_t (solutions not dominated by any other solution in P_t) are identified and assigned a rank 1. The rank 1 solutions are then removed from P_t and the non-dominated solutions in the remaining population form rank 2 solutions. The process is repeated until

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all solutions are assigned a rank. Selection of $N/2$ solutions for the mating pool is then performed according to increasing rank. The crowding distance metric [48] is used if the number of solutions required to fill the mating pool is lower than the available solutions in the rank being considered. The crowding distance of a solution is the perimeter of the rectangle formed by its two neighbors of the same rank; the distance is infinite for points having less than two neighbors (e.g. extreme points). Choice of solutions within a rank is done in decreasing order of crowding distance, thereby giving preference to solutions that are not at close proximity to others. The set of solutions to MOOP are the rank 1 solutions of $P_{Gen_{MAX}}$.

10.5   Empirical Results

In the preparation phase, we conduct risk assessment analysis to initially compute the static risk. Fig. 10.2 shows the unconditional probabilities at the nodes of the test network. We identify 13 security controls that can be used to reduce the risk. We assign a security control cost to each individual control and link each control to the attribute(s) in the BAG that are covered by it. The augmented-BAG resulting from this process is shown in Fig. 10.5. Next, we assign different damage costs and revenue to every attribute in the graph. Although we do not assume any particular cost model and values are assigned hypothetically for the purpose of demonstration, we maintain some relative difference in magnitude to account for the relative importance of different services.

In the first experiment, we assess the expected loss/gain on top of the static risk analysis results (Fig. 10.2) using Eq. (10.9). When using no security control, i.e. a mitigation plan signified by the zero vector, we have an overall expected gain of 622.0 units. Then we assess the cost on the dynamic environment where we assume that two attack incidents have been detected. Fig. 10.4 and Fig. 10.7 show the posterior probabilities and the expected loss/gain at the nodes under this situation. Note that these attack incidents quickly change the business scenario. The total expected loss/gain (LG) changes from 622.0 to -398.17 units. We also notice a change in the momentum of risk. In particular, the posterior probabilities indicate a significant change in risk level at the Administrative server owing to the two attack incidents. This change influences the priority of risks

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Figure 10.7: Augmented-BAG of test network under two attack incidents. The expected loss/gain (V_j) is shown at each node.

identified earlier during static analysis, and highlights the importance of dynamic risk analysis.

Next, we conduct several tests to assess the outcome of using a security control. The base case where no individual control is used yields an expected gain of 622.0 units. Table 10.4 shows the net benefit of using each control individually. At this point, the security administrator may want to rank the security outcomes and build a security mitigation plan from the top-ranked controls. Such a methodology has two drawbacks.

First, the ranking procedure itself is not straight forward because of reciprocal relationships between control cost and expected outcome. For example, "*disable portscan*" and "*filtering external traffic*" when applied alone raises the expected gain from 622.0 units to 875.44 (an increase of 253 units) and 1208.84 units (an increase of 587 units) respectively. The combined outcome when applying both is 1351.27 units (less than $622 + 253 + 587$).

Table 10.4: Security outcome assessment for each individual control in augmented-BAG of test network. Net benefit is $(\mathcal{B} - \mathcal{A} - 622.0)$.

Security Control	Cost $(\mathcal{A})$	Outcome $(\mathcal{B})$	Net benefit
apply OpenSSH security patch	63	1407.89	722.89
apply MS work around	14	1202.45	566.45
filtering external traffic	70	1208.84	516.84
limit DNS access	53	1000.65	325.65
disable portscan	11	875.44	242.44
disable WebDAV	250	1095.90	223.90
apply MS09-004 work around	31	861.10	208.10
add Network IDS	102	858.91	134.91
add Firewall	205	881.15	54.15
encryption	34	681.75	25.75
digital signature	33	673.28	18.28
query restriction	84	681.00	−25.00
use POP3 instead	153	704.67	−70.33

On the other hand, combining "*add Firewall*" (individual increase from 622.0 to 881.15 units) and "*apply MS work around*" (individual increase from 622.0 to 1202.45 units) can raise the outcome to 1735.6 units (greater than $622 + 259 + 580$). The latter two are better choices based on expected outcome, but the former two incurs a lower cost of implementation. This makes the ranking of controls, based on a specific cost factor, a difficult process. Second, even if a ranking has been established, greedy selection can lead to sub-optimal plans. Assume that controls are ranked based on the net benefit they incur individually. The security controls are ordered in this manner in Table 10.4. Given a control cost constraint of say 200.0 units, and a selection scheme based on the ranks, an administrator will choose the first four controls in the table. These controls have a combined cost of 200.0 units and results in an expected gain of 2673.96 units (a net benefit of 2473.96 units collectively). However, selecting the 5th and the 7th controls, instead of the 4th one, effectuates an expected gain of 2809.28 units at the cost of 189.0 units (a net benefit of 2620.28 units). This shows that the security administrator should not choose the security controls based on their individual outcomes or by greedy selection. Instead, a more sophisticated decision making platform is required. This motivates the next three experiments with single and multi-objective optimization.

We conduct three risk mitigation analysis experiments on the test network. The ge-

Figure 10.8: Genetic algorithm solutions to single objective problem obtained by using different weights.

netic algorithm discussed in Section 10.4.3 is used for this analysis. The algorithm parameters are set as follows: population size $N = 100$, $Gen_{MAX} = 50$, crossover probability $= 0.8$, and mutation probability $= 0.01$. We ran each instance of the algorithm five times to check for any sensitivity of the solutions obtained from different initial populations. We also check if running the algorithm for a higher number of iterations (upto 200 generations) results in any improved convergence. However, since the solutions always converged to the same optima (or set of optima), we dismiss the presence of such sensitivity.

In single-objective cost analysis, we run multiple instances of SOOP using different combination of values for α and β. α is varied in the range of $[0,1]$ in steps of 0.05. β is always set to $1 - \alpha$. Fig. 10.8 shows the solutions obtained from this process. It is clear from the figure that equal weight assignment does not necessarily provide the desired balance between the two objectives. Furthermore, the solutions are quite sensitive to the weights and they are not uniformly distributed across different ranges of α.

Fig. 10.9 shows the non-dominated solutions (in $P_{Gen_{MAX}}$) obtained in the multi-objective analysis. Further, all mitigation plans explored by the genetic algorithm during the iterations are highlighted. The algorithm reported all solutions generated for SOOP (using multiple α), as well as several others, specifically solutions in the range where the security control cost is between 200.0 and 700.0 units. These new solutions provide much better

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Figure 10.9: Genetic algorithm solutions to multi-objective problem with static risk assessment.

Figure 10.10: Genetic algorithm solutions to MOOP with dynamic risk assessment.

flexibility in the decision making process. Moreover, performing the multi-objective analysis is much faster than solving SOOP. This is because the security administrator has to solve SOOP with multiple parameter settings in order to identify the plan with the desired outcomes, whereas by solving MOOP, one can generate a good overview of multiple plans in one single run.

In the last experiment, we use the genetic algorithm to assess the choice of security hardening in a dynamic environment. Fig. 10.10 shows the choices of mitigation plans in response to two emerging attack incidents, previously shown in Fig. 10.4. In this plot, we compare the dynamic results with the static ones. Not surprisingly, the plans in this case effectuate lower gains owing to the damage already caused by the attacker when (and at which point) the incidents are detected. Despite this difference, the mitigation plans with similar costs are not so different between the static and dynamic solutions. The three plans highlighted in the figure are very similar to those shown in Fig. 10.9. Such minimal changes in plan characteristics can be considered a positive outcome since the security administrator is not required to revise the entire plan chosen during static analysis. Instead, she can exploit the commonalities for efficient usage of already invested resources. Results from the dynamic analysis also highlight the requirement for pro-active action in security management. Note that although not implementing any controls still results in a positive gain, the appearance of two attack incidents quickly transform this into a case with negative expected outcome.

10.6 Conclusions

In this chapter, we revisit the system administrators' dilemma, namely, how to assess the risk in a network system and select security hardening measures from a given set of controls so as to maximize resource utilization. Our solution methodology uses a Bayesian attack graph model of the network in order to accommodate the likelihoods of cause to consequence transitions during the decision making process. We have provided formal definitions for network characteristics, attacks and security measures under this model. We also show that by using a BAG, we are able to better understand the causal relationships between preconditions, vulnerability exploitations and post conditions. This

is facilitated by computing the likelihoods of different outcomes possible as a result of the cause-consequence relationships. We have demonstrated how the BAG can be used to revise these likelihoods in the event of attack incidents. Using empirical results on a test network, we show that such a dynamic analysis helps the system administrator identify evolving weak spots in a network. We also provide the necessary optimization formulations required to build a mitigation plan that reduces the risk levels. Towards this end, we propose a genetic algorithm capable of performing both single and multi-objective optimization of the administrator's objectives. While single objective analysis uses administrator preferences to identify the optimal plan, multi-objective analysis provides a complete trade-off information before a final plan is chosen.

As immediate future work, we can improve the efficiency of our evaluation algorithm. The evaluation algorithm is used to compute the unconditional probabilities and is currently implemented using a brute force DFS traversal. Posterior probability computation is expensive using this implementation and therefore impacts the decision making time in a dynamic scenario. In particular, we wish to revise the evaluation algorithm to include heuristic based update mechanisms in order to reduce the time required to complete the mitigation analysis, without scarifying the quality of results obtainable. Furthermore, the mitigation process in dynamic situations needs to be improved so that a security administrator can quickly identify the best security response that accounts for all former investments made as part of the static analysis stage.

It is worth mentioning that some security controls have been found to be commonly included in the optimal solutions. It is possible that security hardening is more critical in certain areas of the attack graph. Such areas could be nodes that have multiple fan outs. In other words, these critical areas are at-risk junctions that can be used by an attacker to cause multiple outcomes. Security controls that can reduce risk in such areas are likely to be parts of the optimal solutions. Therefore, it is worth investigating how such controls can be identified efficiently so as to reduce the search space for the optimization algorithm.

CHAPTER 11

Revisiting the Optimality of Security Policies

Over the past several years, researchers have shown considerable interest in the problem of optimal security hardening. Work in this area range from identifying the minimal set of attacks that can break a system [94], to finding the set of security measures with minimum cost that can protect a given system [136], to identifying the optimal set of security controls under a given cost assumption [51]. These works use attack models such as attack graphs and attack trees to statically enumerate a system's vulnerabilities and suggest optimal defenses based on these models.

While these works provide valuable insights into the problem of optimal security hardening, the approaches provide only a static perspective to the problem. They assume that the end goal is to identify a set of security controls that can prevent a particular security breach from occurring. Once a security control is in place, they are done. Since the administrator has to operate within fixed budgetary restrictions it often prevents her from implementing defenses at all the weak spots in the system [51]. Rather, she has to perform a tradeoff analysis between the cost of security control and residual damage for not covering some weak spots. An attacker, meanwhile, continues to explore alternative attack scenarios to inflict maximum damage possible to a system, despite the security controls that are in place. Many a times, the attacker's goal may be just to cause some damage, and not necessarily cause the specific security breach that the defender is trying

to protect against. The attacker may be aided by several factors in this quest. Defenses may have unknown vulnerabilities that can be exploited as a system evolves with time. Misconfiguration of defenses can render them susceptible to attacks. If the attacker has insider knowledge about system configuration, weak spots and defenses (or lack thereof), such knowledge can be used to increase the probability of a defense failing. Such a situation may not be acceptable to the higher ups in the organization. Their perspective may be to not only prevent a specific security breach but also accept minimal collateral damage. This may require a continual updating of the defense strategy based on attacker activities. Thus, one important objective of security hardening, often missed by existing works, is to make life as difficult for the attacker as possible by adjusting security controls. It seems worth investigating if such an arms race between the attacker and the defender will be perpetual or there exists a state involving security controls in which the defender is guaranteed that unexpected damages will never be inflicted no matter how the attacker changes its strategies.

Bistarelli et al. propose *defense trees* as an extension to attack trees to analyze the economic profitability of security measures and their deterrent effects on attackers [24]. A game theoretic perspective of the problem is presented by the same authors in an attempt to discover *Nash equilibrium* points between the security provider and the attacker [23]. An implicit assumption in their work is the existence of a payoff matrix that can be used by a software tool to deduce the points of equilibrium. However, as we explain later, the payoff matrix can be too large to be computed for a given problem.

In this chapter, we revisit the optimal security hardening problem keeping in view the attacker's perspective, namely, defenses can be broken. Our goal is to identify how security controls can be decided to maximize the return on investment for a defender, under the scenario that an attacker is actively engaged in maximizing its return on attacks. We make three major contributions to this effect. First, we show that keeping the attacker's perspective in deciding security controls may often appear to be counter-intuitive but nonetheless provides a better return on investment for an organization. This is because such an approach can lead to an equilibrium condition in which even if the attacker adapts to the installed controls, he cannot expect to cause damages that the de-

fender has not already foreseen and accounted for. From the attacker's point of view, he is better off not trying to breach any of these installed defenses. Our second contribution is that we explore the optimal security control placement problem as a dynamic engagement between the defender and the attacker. We model the problem as an "arms race". We allow both the attacker and defender to start with poor quality strategies and then improve them depending on perceived payoffs. Finally, we solve the optimization problem using the *competitive co-evolution* paradigm and show how the constant engagement between a defender and an attacker drives the solution towards a state of equilibrium. Optimal solution is reached when no further increase in payoff is noticed by either side.

The rest of the chapter is organized as follows. Section 11.1 introduces a motivating example to show how the attacker's perspective may help in choosing optimal security controls. Section 11.2 formalizes the notion of defense and attack strategies. Next, in Section 11.3 we define the payoff models for the attacker as well as the defender. This is followed by the formalization of the optimization problem in Section 11.4. A brief background of competitive co-evolution is provided in Section 11.5 to help understand the solution methodology which is discussed later in the same Section. Empirical results and discussions are presented in Section 11.6. Finally, we conclude in Section 11.7.

11.1 Exploring Defense Strategies

Incorporating the attacker's perspective in the optimal security hardening problem is not easy. We consider a hypothetical example to illustrate how a defender's decision to employ a particular strategy is influenced when the attacker's gains are kept in consideration. Consider the payoff matrix shown in Fig. 11.1. The example assumes that the defender has two possible defense strategies d_1 and d_2, and the attacker has two different attack strategies a_1 and a_2 to try out. The objective of the defender is to decide on one defense strategy to adopt. The first value in a cell (i, j) is a measure of some payoff that the defender derives by adopting strategy d_i under the situation when the attacker uses strategy a_j. Given that the defender is only interested in its payoff value, it uses an average case analysis and finds that strategy d_1 can maintain a higher average payoff than d_2 – 7.5 compared to 5.5. The defender will arrive at the same strategy even with a best case

	a_1	a_2
d_1	(5,6)	(10,2)
d_2	(6,5)	(5,3)

defense strategy

Figure 11.1: A hypothetical payoff matrix showing defender and attacker payoffs.

analysis. The defender therefore installs the defense d_1. However, the decision on d_1 can reveal itself to be flawed when the attacker's payoffs are introduced.

The second value in a cell (i,j) is a measure of the payoff that the attacker derives by adopting attack a_j when defense d_i is in place. With d_1 in place, the attacker sees that its payoff is more (6 compared to 2) by adopting strategy a_1. Hence, it will always employ a_1, in which case the defender will always derive a payoff of 5. This value is not only less than the average payoff of d_1, but is also less than the average payoff of d_2. The value is not even better than the individual payoffs possible with d_2, i.e. 6 when a_1 occurs and 5 when a_2 occurs. Further, if we consider the situation where the attacker does not know which defense is in place and wants to choose a strategy using an average or best case analysis, strategy a_1 is the favorable choice. This is because strategy a_1 always provides a higher payoff than a_2 no matter which defense is in place. In the light of this analysis, the defender should thus be choosing strategy d_2. Since the attacker's choice is inclined towards using a_1, the defender now derives a payoff of 6, compared to 5 when choosing d_1.

Another interesting facet of d_2 is the equilibrium it maintains with a_1. Let us assume that the defender does a best case analysis, as in the case when the attacker's payoffs are not known, and chooses d_1. The attacker then employs a_1 to maximize its payoff. The defender notices that its payoff is not optimal when a_1 occurred and so switches to d_2 to increase it. Hereafter, although the defense strategy has changed, the attacker's best strategy is to stick to a_1. In other words, the defender and the attacker enters a state of equilibrium where none benefits any further from changes in strategy. Hence,

even though d_1 appears to be the optimal strategy at first glance, over time the defender changes policies to finally settle down with d_2 – the *equilibrium strategy*.

One may ask what is the equilibrium solution's relation to the notion of optimal security hardening. Consider the scenario where a defender installs defenses based on some optimality criteria on a system. Over time, an attacker finds the best possible way to exploit the system under the defensive configuration. The defender notices the attacker's exploitation mechanism and modifies its policies keeping in consideration the optimality criteria. The attacker adapt to the changes and the process continues. When the defense policies corresponding to the equilibrium condition are instantiated and the attacker adapts to it, the defender is already running the optimal set of policies possible for the attacker's adaption and does not need to change it. Thus, in the long run, the notion of optimal security hardening converges towards security in equilibrium with attacks.

Performing an analysis of the nature shown in the example is relatively more difficult on a larger scale. First, the payoff matrix can be very large in a real scenario. For d defense controls and a attack points, this matrix can be as large as $2^d \times 2^a$. Filling the matrix can thus involve an immense number of evaluations. Second, even if the matrix can be computed, performing the analysis to decide on the best strategy can be impractical. Note that a best (or equilibrium) defense strategy as depicted in the example may not exist at all. For example, if the values at cell $(2,1)$ are replaced by $(4,10)$, then the best strategy for the attacker varies depending on the defense. Nonetheless, we can argue that d_1 is a better defense strategy in this case since the payoff is better than from d_2 – 5 with a_1 as the strategy of choice for the attacker compared to 4 with a_2 as the attacker's choice.

Ideally, it would be sufficient to decide on a defense strategy by comparing it against others under the light of attack strategies resulting in higher payoffs for the attacker. One may visualize the attack strategies as test cases to measure the competence of a defense strategy. Better test cases are those which are more difficult to solve, or in other words, results in inferior performance of the defense strategy. Similarly, an attack strategy should only be analyzed against defense strategies that result in higher payoffs for the defender. The presence of such cyclic dependencies in the evaluation process makes the analysis hard to conduct. Moreover, the optimal defense strategy will most likely have to be

changed over time to maintain maximum payoff depending on what strategy is chosen by the attacker. Hence we believe it is worth investigating if an equilibrium strategy exists for the security hardening problem.

11.2   Defense and Attack Strategy

We adopt the network and attack tree model described in Chapter 8 to demonstrate the concepts. Recall that nodes of an attack tree are defined as propositions and edges relate the truth value of a node with that of its children. Leaf nodes on the tree represent propositions related to the different network and system states, which may be *true* or *false* depending on what defenses are in place. The truth values of the leaf nodes progressively define if the propositions on the internal nodes would be *true* or *false*. If no defense (also called a security control) is installed, all leaf nodes would be *true*, which would finally lead to the root node to become *true* as well. In such a case, the attacker is assumed to have successfully met its objective. Due to the presence of the *AND-OR* decompositions, the root node may become *true* even if all leaf nodes are not *true*. Similarly, all leaf nodes need not be *false* for the root to become *false.*

A defender installs defenses on the network (makes some or all leaf nodes *false*) so as to prevent the root node from becoming *true*. The defender's choice of defenses may be determined by factors such as the installation cost and the potential damage residual after making the choice. From an attacker's perspective, the attack tree is a model showing the different ways it can compromise the root node. However, we do not restrict our focus to the root node alone. An attacker's strategy might as well be directed towards inflicting the most damage in the presence of defenses, rather than just compromising the root node. The choice of such a strategy is also influenced by the difficulty that the attacker has to overcome in order to bypass any installed defenses.

In order to defend against the attacks possible, the defender can choose to implement a variety of safeguard technologies (or defenses). Each choice of action can have a different cost involved. Besides, some measures have multiple coverages, but with higher costs. The defender has to make a decision and choose to implement a subset of these policies in order to maximize the resource utilization.

Figure 11.2: Payoff for different attack strategies in hypothetical attack tree. Circles denote nodes of the attack tree and the rectangle denote a defense. Value within a circle signify a payoff value that the attacker receives if it succeeds in reaching the node. Value within the rectangle is the cost that the attacker incurs to bypass the defense.

Definition 11.1 DEFENSE STRATEGY. For a given set of d defenses, the defense strategy $\vec{S_D} = (S_{D_1}, S_{D_2}, \ldots, S_{D_d})$ is a Boolean vector indicating which defenses are chosen by the defender. $S_{D_i} = 1$ if defense D_i is chosen, zero otherwise.

The choice of this vector indirectly specifies which leaf nodes in the attack tree would be *false*. An attacker typically exploit leaf nodes that are not covered by any defense in order to progressively climb up the tree, inflicting some amount of damage to the network at every step. However, it is not always correct to assume that an attacker can no longer exploit some parts of the attack tree because of the installed defenses. With the appropriate tools and knowledge, an attacker may have the potential to bypass a defense as well. In other words, leaf nodes which were made *false* by a defense can be reverted back to being *true*. We thus assume an attacker that has the requisite knowledge to breach a defense. However, in order to do so, the attacker will have to incur some cost, often related to the number of defenses in place and the difficulty to breach them. If an attacker's gains are less than the cost incurred, then its effort to breach the defense is not worth the time and value. This primarily motivates the defender to still install defenses despite there being a chance of breach.

Given that the attacker can bypass an installed defense (after incurring a cost), it can start its exploits from any leaf node on the attack tree. The attacker's progress towards

the root is then decided by the leaf nodes it chooses. Note that choosing all leaf nodes that can collectively make an intermediate node *true* need not always be the best approach for the attacker. For instance, given that defenses will be in place at different levels of the tree and the attacker will have to incur a cost to bypass them, it is possible that the attacker derives more payoff by inflicting damages at different parts of the attack tree rather than continuing along a single scenario all the way up to the root. An example of this situation is depicted in Fig. 11.2. With the given values and the defense in place, the strategy 101 generates a higher payoff than trying to reach the root node with strategy 111. This happens because the cost to breach the installed defense nullifies any gains derived from breaching it. An attack strategy is thus defined as follows.

Definition 11.2 ATTACK STRATEGY. Let a denote the number of unique leaf nodes in an attack tree. An attack strategy $\vec{S_A} = (S_{A_1}, S_{A_2}, \ldots, S_{A_a})$ is a Boolean vector indicating which leaf nodes in the tree are chosen by the attacker for exploit. $S_{A_i} = 1$ if node $A_i \in N_{external}$ is chosen, zero otherwise.

Thus, an attack strategy specifies the path(s) that the attacker pursues to an intermediate or the top level of the attack tree. The success of the strategy depends on the defense strategy adopted by the defender, as well as the number of levels it can move up on the tree. Another way to visualize an attack strategy is the set of leaf nodes that the attacker assumes to be *true,* or will make *true* by breaching the defenses protecting them.

11.3 Payoff Model

An augmented-attack tree (Section 8.4.1) extends an attack tree by associating a value V_i to every node. The value associated with a node signifies the sum of the total potential damage present in the child subtree(s) and the potential damage of the node itself. If no defense is installed, i.e. all leaf nodes are *true*, then V_{root} gives the maximum damage possible on the attack tree. When a defender decides on a defense strategy, it essentially sets the truth values of the covered leaf nodes to *false*. Uncovered leaf nodes are set to *true.* An attacker reverts any falsified leaf node to *true* if the node is chosen as part of the attack strategy. With this configuration, we can then find out the damage inflicted on the

attack tree as a result of an attack strategy.

Definition 11.3 DAMAGE INFLICTED. For a given defense strategy $\vec{S_D}$ and an attack strategy $\vec{S_A}$ on an augmented-attack tree AT_{aug}, the damage inflicted DI is given by the value of the root node of the tree, i.e.

$$DI(\vec{S_D},\vec{S_A}) = V_{root}. \tag{11.1}$$

The payoff for a defender and an attacker is an estimate of the gain it receives by adopting a particular strategy and after incurring the corresponding costs associated with the implementation of the strategy. For a defender, the cost of implementation relates to factors such as operations cost, training cost, system downtime, incompatibility cost and installation cost.

Definition 11.4 DEFENSE STRATEGY COST. Given a set of d defenses, each having a cost $C_i; 1 \leq i \leq d$ and a defense strategy $\vec{S_D}$, the defense strategy cost DSC is defined as

$$DSC(\vec{S_D}) = \sum_{i=1}^{d}(S_{D_i}C_i). \tag{11.2}$$

For an attacker, the cost of realizing an attack strategy is related to the effort it has to put forward in overcoming any defenses on its way. We model this cost under a simplistic assumption that stronger defenses are likely to have a higher cost of implementation. Under this assumption, we measure the relative difficulty to breach a defense – a value in $[0,1]$ – and assign the cost to breach it, $BC(\cdot)$, as a fraction (given by the difficulty value) of the cost of implementation of the defense, i.e.

$$BC(D_i) = \frac{C_i}{\underset{i}{Max\,C_i}} \times C_i. \tag{11.3}$$

Definition 11.5 ATTACK STRATEGY COST. Given a set of d defenses, a defense strategy $\vec{S_D}$ and an attack strategy $\vec{S_A}$ on an attack tree AT, the attack strategy cost ASC is defined as

$$ASC(\vec{S_D},\vec{S_A}) = \sum_{i=1}^{d} \sum_{j|D_i(A_j)=false} [BC(D_i)S_{D_i}S_{A_j}]. \tag{11.4}$$

The expression above iterates through the leaf nodes covered by a particular defense. Thereafter, the cost to breach the defense is added to the attack strategy cost if the defense is part of the defense strategy and the leaf node is part of the attack strategy. When a breach occurs, the cost paid by the defender to install it (C_i) is a loss, called the breach loss $BL(\cdot)$ and expressed in a manner similar to the above equation.

$$BL(\vec{S_D}, \vec{S_A}) = \sum_{i=1}^{d} \sum_{j|D_i(A_j)=false} [C_i S_{D_i} S_{A_j}] \tag{11.5}$$

We then define the defender and attacker payoffs as follows.

Definition 11.6 PAYOFF FOR DEFENDER AND ATTACKER. For a given defense strategy $\vec{S_D}$ and an attack strategy $\vec{S_A}$ on an augmented-attack tree AT_{aug}, the defender's payoff POD is given as

$$POD(\vec{S_D}, \vec{S_A}) = DI(\vec{0}, \vec{1}) + DSC(\vec{S_D}) - DI(\vec{S_D}, \vec{S_A}) - BL(\vec{S_D}, \vec{S_A}), \tag{11.6}$$

and the attacker's payoff POA is given as

$$POA(\vec{S_D}, \vec{S_A}) = DI(\vec{S_D}, \vec{S_A}) - ASC(\vec{S_D}, \vec{S_A}). \tag{11.7}$$

Here, $DI(\vec{0}, \vec{1})$ signify the maximum damage possible on the attack tree, which happens when there are no defenses installed and the attacker exploits all leaf nodes. $\vec{0}$ represent the all zero vector and $\vec{1}$ is the all one vector. Note that both payoff functions employ the same DI value derived from the attack tree. One can argue that the attacker's knowledge on the damages sustained by the defender when compromising a node is rather limited, and thus cannot be same as that of the defender. Further, the attacker need not have the complete knowledge about the cost of implementing a defense and hence will not know the exact value of ASC. We understand that both are rational arguments. Our justification to them is based on the fact that the POA function need not be an exact estimate of the actual payoff derived by the attacker. The optimization process only needs to compare payoff values to determine the relative effectiveness of two attack strategies, in which case it suffices to have a value proportional to the actual payoff. The POA function satisfies this requirement since the attacker's actual payoff is likely to be proportional to the damage it inflicts on the tree. Moreover, the cost paid by the attacker to overcome a defense will likely be proportional to the sustainability of the defense.

11.4 Problem Statement

We first normalize the *POD* and *POA* functions in order to account for differences arising in the magnitude of the values. The *POA* function is in the range of $[-ASC(\vec{S_D},\vec{S_A}),$ $DI(\vec{0},\vec{1})]$ which is remapped to $[0, ASC(\vec{S_D},\vec{S_A}) + DI(\vec{0},\vec{1})]$ by adding $ASC(\vec{S_D},\vec{S_A})$ to the value. *POD* function is in the non-negative range $[0, DSC(\vec{S_D}) + DI(\vec{0},\vec{1})]$. The normalized functions for *POD* and *POA* – in the range of $[0,1]$ – are then given as follows.

$$POD_{norm}(\vec{S_D},\vec{S_A}) = \frac{POD(\vec{S_D},\vec{S_A})}{DSC(\vec{S_D}) + DI(\vec{0},\vec{1})} \tag{11.8}$$

$$POA_{norm}(\vec{S_D},\vec{S_A}) = \frac{POA(\vec{S_D},\vec{S_A}) + ASC(\vec{S_D},\vec{S_A})}{ASC(\vec{S_D},\vec{S_A}) + DI(\vec{0},\vec{1})} \tag{11.9}$$

The normalized versions are more intuitive in understanding what the payoff functions model. The defender has an investment worth $DSC(\vec{S_D}) + DI(\vec{0},\vec{1})$ on the attack tree. POD_{norm} gives the fraction of this investment protected by the defender's strategy for a particular attack strategy. In other words, POD_{norm} gives the fractional return on investment for the defender. From an attacker's perspective, the best it can do is gathering the payoff from maximum damage and also retain the cost incurred while doing so to itself. $DI(\vec{S_D},\vec{S_A})$ is the amount that it actually derives. POA_{norm} is thus the fractional return on attack to the attacker.

The defender's optimization problem is to find a defense strategy $\vec{S_D}$ that gives maximum POD_{norm} under all possible attack strategies. The attacker's optimization problem is to find an attack strategy $\vec{S_A}$ that gives maximum POA_{norm} under all possible defense strategies. However, such a strategy may not exist. Besides, as argued earlier, evaluating a host strategy with all opponent strategies is often impractical. We introduce here the terms *host* and *opponent* to refer to the party whose strategy is being tested and the party against whom it is being tested respectively. In order to compare two host strategies, it is sufficient to evaluate them against their respective best opponent strategy (one generating the highest payoff for the opponent with the host strategy in place). Hence, a more suitable statement of the optimization problem is as follows.

Problem 11.1 Defender's Optimization Problem. Given an augmented attack tree AT_{aug} and d defenses, find the defense strategy $\vec{S_D}^*$ that maximizes $POD_{norm}(\vec{S_D}, \vec{S_A}^*)$, where $\vec{S_A}^*$ satisfies the relation $POA_{norm}(\vec{S_D}, \vec{S_A}^*) \geq POA_{norm}(\vec{S_D}, \vec{S_A})$ for any attack strategy $\vec{S_A}$.

Problem 11.2 Attacker's Optimization Problem. Given an augmented attack tree AT_{aug} and d defenses, find the attack strategy $\vec{S_A}^*$ that maximizes $POA_{norm}(\vec{S_D}^*, \vec{S_A})$, where $\vec{S_D}^*$ satisfies the relation $POD_{norm}(\vec{S_D}^*, \vec{S_A}) \geq POD_{norm}(\vec{S_D}, \vec{S_A})$ for any defense strategy $\vec{S_D}$.

The brute force method to solve each problem is to first generate the payoff matrix and then mark the cell, for every host strategy, with the highest opponent payoff. The solution is the host strategy which has the highest payoff in the marked cells. If, given the host strategy in the solution, the opponent's payoff is also the highest, and vice versa, then the solution admits a Nash equilibrium [135]. We want to emphasize here that solving just one problem is not sufficient. For example, assume that the defender has found the optimal solution to its problem. The POD_{norm} reported by the solution implicitly assumes that the attacker will launch the strategy $\vec{S_A}^*$ that gives the highest attacker payoff – established in the optimization problem by the relation. If the attacker also solves its own optimization problem, there is no guarantee that the best strategy found by it is the same $\vec{S_A}^*$ as found in solving the defender's optimization problem. The outcome in this case could be that both the attacker and the defender get sub-optimal payoffs. The desired solution is the defense and attack strategy pair $\vec{S_D}^*$ and $\vec{S_A}^*$ that satisfy the conditions

1. $POD_{norm}(\vec{S_D}^*, \vec{S_A}^*) > POD_{norm}(\vec{S_D}, \vec{S_A}^*)$, and

2. $POA_{norm}(\vec{S_D}^*, \vec{S_A}^*) > POA_{norm}(\vec{S_D}^*, \vec{S_A})$,

for any given defense strategy $\vec{S_D}(\neq \vec{S_D}^*)$ and attack strategy $\vec{S_A}(\neq \vec{S_A}^*)$.

11.5   Competitive Co-Evolution

Competitive co-evolution refers to the concurrent evolution of two distinct species in which the fitness of an individual in one species is based on its competitive abilities

against the individuals of the other species. Fitness evaluation with such reciprocal actions are hypothesized to occur in nature. Game theory based models of such interactions is first presented in Axelrod's *Prisoners' Dilemma* [12]. The evolution of species in such a competitive habitat usually leads to a *evolutionary stable strategy* [157] which cannot be invaded by the process of natural selection. In other words, the species reverts back to the stable strategy over time.

Competitive co-evolution has been successfully applied to the evolution of strategies for games such an Tic-Tac-Toe and Nim [148]. The range of potential opponent strategies is typically very big in such games, thereby making it difficult to determine an exogenous fitness evaluation function. Other domains such as software reliability testing faces a similar problem. The solution using competitive co-evolution involves using two populations, one representing the software solutions and the other representing the test cases, each taking turns in testing and being tested against the other [82]. A survey of other real world applications is available in [28].

Success of competitive co-evolution is attributable to the emergence of an evolutionary "arms race" [44]. Consider two populations of defense strategies and attack strategies. To begin with, both populations are likely to have strategies of poor quality. Most of the host strategies will have low payoffs brought forth by one or two good strategies existing in the opponent population. However, since defense strategies are evolving based on their competitive abilities against attack strategies, the success of the defender implies the failure of the attacker. When the attacker finds strategies to improve its payoff by overcoming the failure, it helps the defender identify gaps previously unthought of in its previous strategies. The same idea drives the attacker's strategies. New opponent strategies drive hosts towards better counter strategies, improving host performance by forcing it to respond to a wider range of more challenging test cases.

The next question that comes to mind is whether a good host strategy of the current generation can prove its competence against opponent strategies that are lost in the evolution of the opponent population. This is referred to as the *memory property* in co-evolution. To handle such situations, co-evolutionary algorithms employ a "hall of fame" [149] sub-population which keeps track of the best opponent solutions found from earlier

generations. Success of a competition for a host strategy is measured not only relative to the current opponent strategies, but is also dependent on its performance against the opponent's hall of fame. Other similar methods are elaborated in [70, 158]. Details of our implementation are presented next.

We begin with two randomly generated populations Pop_A and Pop_D of size N_A and N_D respectively. Pop_A refers to the population of attack strategies $\{\vec{S_A}^1,\ldots,\vec{S_A}^{N_A}\}$ and Pop_D refers to that of defense strategies $\{\vec{S_D}^1,\ldots,\vec{S_D}^{N_D}\}$. In every generation, every strategy in a population is evaluated with the best opponent strategy (one with highest fitness as described later) of the previous generation to find POA_{norm} and POD_{norm}. The notations $\vec{S_D}^{prevbest}$ and $\vec{S_A}^{prevbest}$ is used to denote the best defense and attack strategy from the previous generation respectively. For the first generation, the best strategies are chosen randomly from the populations.

Next, each strategy in the populations is assigned an *age count, Age*($\cdot$), signifying the number of iterations for which it has survived the evolutionary process. Each strategy begins with an age count of zero, which is incremented every time it manages to enter the next population. The age is reset to zero if the strategy no longer exists in the next population. With this, the fitness of a defense strategy $\vec{S_D}^i$ in generation (iteration) t is assigned as

$$F(\vec{S_D}^i,t) = \frac{F(\vec{S_D}^i,t-1) \times Age(\vec{S_D}^i) + POD_{norm}(\vec{S_D}^i,\vec{S_A}^{prevbest})}{[Age(\vec{S_D}^i)+1]}, \tag{11.10}$$

and that of an attack strategy $\vec{S_A}^j$ in generation t is assigned as

$$F(\vec{S_A}^j,t) = \frac{F(\vec{S_A}^j,t-1) \times Age(\vec{S_A}^j) + POA_{norm}(\vec{S_D}^{prevbest},\vec{S_A}^j)}{[Age(\vec{S_A}^j)+1]}. \tag{11.11}$$

The fitness is an average measurement of the payoff of a strategy throughout the evolutionary process. With this fitness assignment, each population then independently undergoes the usual process of evolution as in a genetic algorithm (GA) – selection, crossover and mutation [75] – and creates a new population of strategies. The best strategies of the past H generations replace H randomly selected strategies in the respective populations. The process is repeated until a set number of generations. Fig. 11.3 depicts the algorithm. The parameters of the algorithm are set as follows: $N_A = 100$, $N_D = 100$, $H = 10$, single

Figure 11.3: Schematic of competitive co-evolution of attack and defense strategies.

point crossover with probability 0.5, probability of mutation = 0.01, 2-tournament selection and 1000 generations. In the experiments, we use the 19 defenses used in Chapter 8 and the attack tree has 13 unique leaf nodes. The defender thus has 2^{19} defense strategies and the attacker has 2^{13} attack strategies to choose from.

11.6 Empirical Results

We first present the results on the sensitivity of the genetic algorithms used to their parameters. Some parameters involved in the GA affect the dynamics of the arms race undergoing between the two populations. Using a higher probability of crossover or mutation affects the age count of a solution. A high probability decreases the chances of a strategy surviving for long across iterations, thereby interrupting its chances of competing against a wider variety of opponent strategies. Increasing the population size gives a faster convergence rate, although the solution remains unaffected. We also increased the number of iterations from 1000 to 5000 to see if a dormant strategy becomes prominent over time. However, no such outcome is observed.

Fig. 11.4 shows how the fitness of the best strategy in the defender and attacker populations change across generations. The random initialization of the two populations usually starts off the competition with comparatively higher fractional payoff for the de-

Figure 11.4: Fitness of best defense and attack strategies in every iteration of the co-evolutionary process.

fender. However, the attacker immediately finds strategies to improve its payoff, and reciprocally decreases the payoff for the defender, as can be seen on the steep decline of the defender's payoff. There is even a phase between the 50th to 150th generations when the attacker continued to evolve strategies with similar payoff, but ones that continued to decrease the payoff for the defender. The arms race becomes prominent after this phase. The arms race is indicative of the period when the defender and the attacker continuously change their strategies to cease the decline in their payoffs brought forth by an improved opponent strategy. In a way, this arms race depicts the change in policies that the defender has to sporadically keep enforcing in order to subdue the affects of an evolving attacker.

Fig. 11.5 depicts the dynamics of the two populations during the 100th to the 200th generations. The average fitness of each population is plotted to show the interactions happening between them. The arms race is distinctly visible after the 130th generation – one population reacts to changes in the other. Rising up to a peak indicate the phase of steady improvement in host strategies against those of the opponent's. Falling down to a pit signify the reverse. As is depicted by the vertical lines, a rising period in one population results in a falling period in the other, and vice versa. Note that the rise in one population and the fall in the other are not correlated in terms of the payoff

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Figure 11.5: Average fitness of defender and attacker strategies showing "arms race" population dynamics between the 100th and the 200th generations.

values. An attacker's marginal improvement in payoff can result in significant drop in the defender's payoff. More interestingly, there is no fixed duration within which the two populations alternate between rise and fall. In other words, the dynamics of finding a strategy to tackle the currently dominating opponent strategy is not known. We stress this phenomena since any defense strategy not in equilibrium with the attacker's eventually results in a decline in the payoff. Ideally, the better the strategy, the slower will be the decline; emphasizing that the attacker faces more difficulty in finding a counter strategy to improve its payoff.

However, with the static attack tree in place, the process of arms race does not continue forever. Both the attack and the defense strategies stabilize at around the 500th generation. No host at this point manages to find a strategy to improve its payoff given the best strategy the opponent has at the point. However, this stability in the strategies is not sufficient to conclude that the attacker and defender are now in an equilibrium. This follows from the fact that there may exist an undiscovered opponent strategy that can reduce the payoff generated from the stable host strategy.

In order to demonstrate the effectiveness of competitive co-evolution in generating an equilibrium strategy pair, we perform the following supplementary analysis. The defender's best strategy $\vec{S_D}^{best_t}$ in every generation t ($1 \leq t \leq 1000$) of the process is noted.

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Figure 11.6: Payoffs when an independent GA is run for each best host strategy from every generation to find the best opponent strategy for it. **o/+** represent the payoffs when the defender/attacker is the host.

For each such strategy, we run a simple genetic algorithm to generate the attack strategy $\vec{S}_A^{best_t}$ with the highest attacker payoff. Fig. 11.6 shows the defender and attacker payoffs (in circles) for the pairs $\vec{S}_D^{best_t}$ and $\vec{S}_A^{best_t}$. A similar process is done taking the attacker's best strategy of every generation. The plus signs in the plot depict the payoffs for the pairs obtained from the process. We find that the only circle and plus coinciding corresponds to the stable strategy of the defender and the attacker as returned by the co-evolutionary optimization. If the defender chooses the stable defense strategy, the attacker's payoff is maximum when it uses the stable attack strategy. If the attacker uses the stable attack strategy, the defender's payoff is maximum when it uses the stable defense strategy. In other words, the stable defense and attack strategy pair is indeed an equilibrium point.

Fig. 11.7 depict the equilibrium solution on the attack tree of the example network. There exists two leaf nodes on the top half of the tree which must be *true* for the attacker to reach the root. Interestingly the defender's equilibrium strategy does not involve putting a defense to cover those nodes. Given that an attacker willing to maximize the damage will identify the bottleneck as well, the majority of its attack strategies will involve by-passing any defense put on those nodes. If the cost to do so is not higher than the gain derived from compromising the nodes in the lower half of the attack tree, then most certainly efforts to make the breach will be most here. The defender's strategy do identify

Figure 11.7: Defense and attack strategy of equilibrium solution.

the subtree resulting in user access on the FTP server (depicted by the star) as a launching pad for many other attacks. In the equilibrium condition, the attacker is better off not trying to breach any installed defenses. However, it can inflict damages to most parts of the tree by breaching the defense installed on the fore mentioned subtree and hence must make an effort to do so.

One of the forthcoming questions resulting from this analysis is whether an organization's investments be directed towards a static minimal cost security policy or proactively be channeled towards an equilibrium policy. The minimal policy resulting from a one-time evaluation may incur a lower cost w.r.t. a short time window, but under an evolving attacker model, this cost must be supplemented by further investments over time. Hence, the return on investment can potentially be lower than that could be received by enforcing the equilibrium policy. On the other hand, an equilibrium policy may not enforce sufficient safeguards to protect a network from short-term losses. We believe this necessitates investigating the security hardening problem from a perspective where an evolving defense model must also be integral to defining the notion of an optimal security policy.

11.7 Conclusions

Incorporating strong defenses against malicious attackers is challenging. Simply installing the best available defenses does not work for several reasons. The security admin-

istrator has to work within fixed budgetary constraints and has to explain the return on investment of security controls to the management. However, any convincing argument explaining the return on investment must take the attacker's benefits into consideration. Our first contribution in this chapter is the identification of this fact in order to motivate the pursuance of security hardening keeping both the defender and attacker payoffs into account.

We argue that the notion of optimal security hardening is often dictated by the constant interaction between the defender and the attacker. What is perceived as the optimal return on investment would cease to be so once the attacker's strategy to exploit the defensive configuration is understood. We justify this with the argument that the system administrator would more than likely be approached by the management to re-evaluate its defenses under the light of the attacker's strategy. Our second contribution highlights that the dynamic engagement between the attacker and the defender is a continuous process ending only when both enter a state of equilibrium. We thus emphasize that identifying such equilibrium conditions, if any, is where the true sense of optimal security hardening is buried. To this end, we formulate the requisite optimization problems and present the notion of equilibrium in terms of the formulated problems.

As a viable methodology, we propose the use of competitive co-evolution to generate the aforementioned equilibrium strategies. The method involves an algorithm that intrinsically models the arms race undergoing between the attacker and the defender, with the ability to effectively find the equilibrium solutions. Solutions from the hypothetical example demonstrate that keeping the attackers' perspective in consideration for security administration can result in placing security controls in places that may appear non intuitive but provide a better return on investment for an organization.

In this work, we assume the existence of a single attacker owning any generated payoff. However, the problem can be more interesting when payoff models can be designed to incorporate multiple attackers working in conjunction to achieve a particular objective and sharing the payoff so generated. Further work can be directed towards designing algorithms that can identify the existence of multiple equilibrium points simultaneously. We believe that Pareto analysis intended towards generation of such solutions is a promis-

ing avenue to explore. Formal analysis to determine if equilibrium solutions exist at all would be a major contribution as well.

Part III

Wireless Data Broadcasting

CHAPTER 12

Utility Driven Data Broadcast Scheduling

Recent advances in wireless communication technology are increasingly making the dream of pervasive computing a reality. Pervasive computing involves a network of portable computing devices so thoroughly embedded in our day-to-day work and personal life that their existence becomes difficult to perceive altogether. The devices interact with each other and with other computing devices by exchanging rapid and continuous streams of data. To facilitate almost imperceptible human-computer interaction, data access times in such environments must be maintained within a specified Quality of Service (QoS) level. Challenges in doing so arise from the fact that wireless bandwidth is typically a limited resource, and thus it is not always possible to meet the quality requirements of every device. This constraint not only makes pervasive data access a challenging problem, but also identifies "optimal resource allocation" as one of the major research problems in this domain.

A pervasive environment encompasses both peer-to-peer and client-server modes of data dissemination. For example, a typical pervasive health care system may involve multiple sensor nodes disseminating data on the monitored vital signs of a patient to a personal digital assistant carried by the attending health care personal. Data communication follows a peer-to-peer architecture in such a setting. On the other hand, a pervasive environment designed to serve queries on flight information in an airport is based on

a client-server mode of communication. Flight information usually reside in a database server from where data is disseminated based on the incoming queries. For an environment like an airport, it is appropriate to assume that the database will be queried more frequently for certain types of data. Similar situations can be imagined in a stock trading center, a pervasive traffic management system, or a pervasive supermarket. Such scenarios open up possibilities of adopting a broadcast based architecture to distribute data in a way that multiple queries for the same data item get served by a single broadcast. The focus of this chapter is directed towards data access issues in such pervasive environments.

Quality of service is an important facet in such pervasive data access. Consider the following example application - a traffic management system for a big city. The city government implements a traffic pricing policy for all vehicles on all roads based on factors such as the distance traveled, the type of road traveled on, the time of day, vehicle category and customer type (for example, special fee paying, traveling salesman, paramedic on-call etc.) The system gathers real time traffic data via thousands of sensors, such as traffic cameras, magneto-resistive traffic sensors and radars spread throughout the city. To help drivers optimize their travel costs, the traffic management system provides various routing services to drivers to avoid roadblocks, construction delays, congestion, and accidents.[1]

A driver requests and gets routing services using smart GPS equipped devices as follows. A device periodically uploads the vehicle's current location (based on GPS information), intended destination, time willing to spend for traveling to destination, a prioritized list of routing restrictions (for example, waypoints that the driver would like to visit if possible, rest stops, scenic byways etc.), vehicle category and customer type, and current speed. In response, the server replies with a set of route information. Each route information contains among other things information about road segments to travel on this route and traffic patterns on these road segments. The GPS equipped device uses

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[1]Such an application is not very far-fetched. The Dutch government is in the process of introducing a similar electronic traffic management system and pricing model. See [45].

this information and uses other parameters set by the driver to compute a good route to take. Note that since the response to many drivers will no doubt contain overlapping route segments and traffic information, it makes sense broadcasting this data.

The requests arrive at the server with various priority levels and soft deadlines. For example, a higher fee paying customer gets a higher priority than a lesser fee paying customer. A VIP's convoy may get a still higher priority. Emergency responders get the highest priority. A routing request that comes associated with a set of routing restrictions automatically gets associated with a set of soft deadlines based on the speed of the driver. The requests from different drivers may also get re-prioritized at the server so as to meet a broader goal of reducing congestion and enabling smoother traffic flows.

Let us assume that at some point there is a major traffic gridlock within the city and the traffic server gets thousands of re-routing requests from users. While these requests are queued up at the server, another request comes from a VIP's convoy with a high priority and deadline. At this time, the server needs to determine how to schedule this request. Pre-empting others may enable the server to meet the timeliness of this latest request. However, serving some or all of the earlier requests has the advantage of clearing up the gridlock earlier.

In this example, the different data that the server needs to serve is associated with different utility values. Owing to the dynamic nature of the utility of responses to queries, the time criticality factor cannot be ignored altogether when disseminating data. The server would like to satisfy as many queries in a timely manner as possible. However, some queries may be delayed beyond their specified deadlines (for example, the query from the VIP's convoy). The users, who hardly realize the broader goals of the traffic management system and various bottlenecks in the information infrastructure, would like to have their requests served at the earliest; however, it is reasonable to assume that delayed data still provide some utility even if received after a specified deadline. For example, a delayed route information may prevent a driver from visiting a particular waypoint en route or may require the driver to use the next gas station. Nonetheless, it may still allow the driver to choose a good route to the destination. An assumption of reduced data utility in the face of missed deadline, enables data broadcasts to be tailored

in such a way that total utility associated with a broadcast is maximized. This helps maintain a certain QoS level in the underlying infrastructure.

Note that the specific problem we are trying to address is, by no means, restricted only to the example application outlined above. Similar scenarios are witnessed in environments such as Stock Exchanges. Here stock brokers on the floor seek and receive periodic market data on wireless hand-held devices and notebooks and analyze them to determine which stocks are hot. They also buy and sell stocks using such devices. Popularity of stocks change throughout the day, and it is important to analyze such trends along multiple dimensions in order to buy and sell stocks. Thus, although queries from brokers are implicitly associated with deadlines, these can be considered soft. The overall goal of the Stock Exchange is still to satisfy as many requests as possible in a timely manner. To wrap up the scope of the problem domain, we would like to point out that in order to make useful decisions, the stock broker may make a request for a group of data from the Stock Exchange (for example, stock prices of three different oil companies). A typical constraint on such requests is that all these data items must be received (not necessarily in any particular order) before the stock broker can perform the relevant local analysis of the market trends. Such a request can be considered a *transactional request* (or simply a *transaction*). For scheduling in such cases, additional constraints must be placed for ensuring the validity of data received.

Wireless broadcast mechanisms have been extensively investigated earlier. However, very few of these research give attention to the effective utility involved in the timely servicing of a request. Time criticality has been earlier addressed in a number of contexts with the assumption of a *hard deadline* [69, 96, 100, 104, 107, 183, 187]. Broadcast scheduling in these works mostly focus on the timely servicing of a request to minimize the number of missed deadlines. When the assumption of a *soft deadline* is appropriate, a broadcast schedule should not only try to serve as many requests as possible, but also make a "best effort" in serving them with higher utility values. Often, heuristics are employed in a dynamic setting to determine these schedules. However, their designs do not involve the QoS criteria explicitly. Heuristic based methods make local decisions w.r.t. a request or a broadcast, and often fail to capture the sought global QoS requirement. Much

of this is due to the fact that real time decision making cannot span beyond an acceptable time limit, thereby restricting the usage of "full length" optimization techniques to the domain. It does become imperative to design hybrid strategies that can combine the fast real time performance of heuristics and the better solution qualities obtained from search based optimization.

In this chapter, we propose a dynamic scheduler that tries to maximize the overall utility of servicing requests and at the same time tries to serve as many requests in a timely manner as possible. The setup is a wireless broadcast environment as in pervasive computing applications. For this work, we choose to ignore the problem of power constraints of wireless devices. We acknowledge that this is serious problem because the entire time a wireless device is waiting to receive a broadcast, it has to keep its power status in *active* mode rather than reverting to a power saving *doze* mode. However, this can be addressed by packaging the transmitted data appropriately. A number of works have been done in *air indexes* [91, 194] that try to facilitate power saving by broadcasting index information along with the data. From the air index information, the receiver can predict the arrival time of its data and can accordingly switch its power level to doze or active mode. Such a strategy can very well be incorporated in our model.

We begin with an attempt to understand the nature of the underlying search space, and argue that traditional heuristics usually generate solutions in a worse part of this space w.r.t a given global utility measurement. We explore "local search" as an option to boost the performance of these solutions and provide arguments as to why the option is viable in a real time setting. The observations allow us to propose a light weight stochastic hill climber that surpasses the performance of a heuristic, and explicitly searches the space of schedules to maximize utility. We believe that the proposed method provides insights into better broadcast mechanisms which often clear our understanding for better heuristic design.

The rest of the chapter is organized as follows. Section 12.1 summarizes the related work in this domain. The broadcast model and the scheduling problem are discussed in Section 12.2. The explored solution methods and the experimental setup are described in Section 12.3 and Section 12.4 respectively. Results and discussions from the experiments

are summarized in Section 12.5. Finally, Section 12.6 concludes the chapter.

12.1   Related Work

Data broadcasting has been extensively studied in the context of wireless communi-
cation systems. Su and Tassiulas [161] study the problem in the context of access latency
and formulate a deterministic dynamic optimization problem, the solution to which pro-
vide a minimum access latency schedule. Their experimental results show that the mean
response times in push-based and on-demand broadcasts become similar as the request
generation rate increases. Acharya and Muthukrishnan propose the *stretch* metric [2] to
account for differences in service times arising in the case of variable sized data items.
Their work identifies that maintaining a balance between local and global performance
is a key factor in on-demand broadcasting environments. To this effect, they propose
the *MAX* heuristic to optimize the worst case stretch of individual requests. Another
attempt to balance individual and overall performance is seen in the work by Aksoy and
Franklin [7]. Their *RxW* heuristic is an attempt to combine the benefits of the MRF and
FCFS heuristics, each known to give preference to popular and long standing data items
respectively. Sun et al. propose the LDCF algorithm to account for various cost factors
typically observed in broadcast systems [162]. Factors such as access time, tuning time
and cost of handling failure requests are used to compute a delay cost for data items,
which then serves as a priority measure for the items. Hameed and Vaidya adapt a *packet
fair queuing* algorithm to the domain [80]. Their approach exploits the similarities in the
two problem classes and gives an efficient algorithm to solve the problem. Lee et al.
provide a survey of these techniques [106] and their applicability in the area of pervasive
data access.

The above mentioned algorithms ignore the time criticality factor while serving data
requests. Early work on time constrained data request is presented by Xuan et al. [190].
Earliest deadline based on-demand scheduling is the heuristic of choice in this seminal
work. Jiang and Vaidya address the issue by considering broadcast environments where
clients do not wait indefinitely to get their request served [96]. They model the user
impatience as an exponential distribution and propose the *SRM* algorithm to minimize

the mean waiting time, which in turn maximizes the expected service ratio. Lam et al. look at the time criticality factor from the perspective of temporal data validity [104]. Their approach assigns an *absolute validity interval* to determine the refresh frequencies of data items in order to keep the cache status of the items updated using broadcast strategies. Fernandez and Ramamritham propose a hybrid broadcasting approach to minimize the overall number of deadlines missed [69]. Their adaptive model takes into consideration the data and time-criticality requirements to determine periodic and on-demand schedules for data items. Kim and Chwa present theoretical results on the competitive ratios of scheduling algorithms working with time constrained requests [100]. Temporal constraints on data are revisited by Wu and Lee with the added complexity of request deadlines [183]. Their *RDDS* algorithm assigns a priority level to each requested data item based on the number of requests for it, the effective deadline and the size of the item, and broadcasts the item with the highest priority. Xu et al. propose the *SIN-α* algorithm to minimize the request drop rate [187]. However, their approach does not take variable data sizes into consideration. This motivates Lee et al. to propose the *PRDS* algorithm that takes into account the urgency, data size and access frequencies of various data items [107].

Several shortcomings of using a strict deadline based system are discussed by Ravindran et al. [144] in the context of real-time scheduling and resource management. A deadline is usually a linear-valued expression that fails to distinguish between urgency and importance. Although time-utility functions in data broadcast scheduling have been ignored for some time now, the idea has been extensively researched in other real-time scheduling domains. Jensen et al. point out that real-time systems are usually associated with a value based model which can be expressed as a function of time [93]. They introduce the idea of time-utility functions to capture the semantics of soft time constraints which are particularly useful in specifying utility as a function of completion time. An attempt to understand the benefit of utility values in hard deadline scheduling algorithms is done by Buttazzo et al. [31]. Wu et al. study a task scheduling problem where utility is considered a function of the start time of the task [182]. Similar studies [38, 113, 175] performed on utility accrual in task scheduling problems show that heuristics designed

to cater to the deadline requirement alone are not sufficient, and care should be taken to address any non-linear characteristics of the time-utility dependencies.

Although the different problem classes in scheduling have similarities in them, the idea of multiple requests getting served by a single broadcast make the data broadcast scheduling domain somewhat different. We believe time-utility functions are a better alternative to hard deadline specifications of data requests since they allow better generalization of the time constraints involved. Thereby, we introduce the notion of utility accrual to a data broadcast environment and explore the issues generated thereof.

12.2   Broadcast Scheduling

Wireless data broadcasting is an efficient approach to address data requests, particularly when similar requests are received from a large user community. Broadcasting in such cases alleviate the requirement for repeated communication between the server and the multiple clients interested in the same data item. Push-based architectures broadcast commonly accessed data at regular intervals, depending on a well known access pattern, and in the process removes the requirement of a client actually sending the request to the server. Contrary to this, on-demand architectures allow the clients to send their requests to the server. However, access to the data item is facilitated through a broadcast which, for more frequently requested data, serves multiple clients at a time. Broadcast scheduling in this context is the problem of determining the order in which data items should be broadcast so that more clients are served at a time within an acceptable quality requirement.

Data access in pervasive environments can be modeled with an on-demand broadcast architecture where particular emphasis has to be paid to the time criticality and utility of a served request. The time criticality factor stresses on the fact that the requested data is expected within a specified time window; failure to do so would result in an utility loss. Given the immense number of requests that may arrive at such a data broadcast server, it is often not possible to serve all requests in a timely manner. A broadcast schedule in such environments has to cater to the added requirement of maintaining a high utility value for a majority of the requests.

Figure 12.1: Typical on-demand broadcast architecture in a pervasive environment.

12.2.1 Broadcast model

Fig. 12.1 shows a typical on-demand data broadcast architecture in a pervasive environment. Various client devices use an uplink channel to a data provider to request various data items served by the provider. The data items are assumed to reside locally with the data provider. Each request Q_j takes the form of a tuple $\langle D_j, R_j, P_j \rangle$, where R_j is the response time within which the requesting client expects the data item D_j and asserts a priority level P_j on the request. Note that a single request involves only one data item. A client requiring multiple data items sends multiple requests for each data item separately. Requests from the same client can be served in any order. The data provider reads the requests from a queue and invokes a scheduler to determine the order in which the requests are to be served. It is important to note that new requests arrive frequently into the queue which makes the task of the scheduler rather dynamic in nature. The scheduler needs to re-examine the previously generated schedule to accommodate the time critical requirements of any new request. Data dissemination is carried out through a single channel data access point. Clients listen to this channel and consider a request to be served as soon as the broadcast for the corresponding item begins. The single channel assumption is not critical to our work, and changes in approach will be mentioned wherever appropriate.

The underlying scheduler is invoked every time a new request is received. At each

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Figure 12.2: Utility of serving a request.

invocation, the scheduler first determines the requests that are being currently served and removes them from the request queue. The data items required to serve the remaining requests are then determined and a schedule is generated to serve them. The scheduler tries to make a "best effort" at generating a schedule that respects the response time requirements of the requesting devices.

12.2.2 Utility metric

The utility of a data item broadcast is measured from the response time and priority level of the requests served by it. The response time r_j of a request Q_j arriving at time $t_{j,arr}$ and served at time $t_{j,ser}$ is given by $(t_{j,ser} - t_{j,arr})$. We assume that the utility of a data item received by a client decreases exponentially if not received within the expected response time (Fig. 12.2) [93]. For a given request Q_j, the utility generated by serving it within a response time r_j is given as

$$u_j = \begin{cases} P_j & ,r_j \leq R_j \\ P_j e^{-\alpha_j(r_j - R_j)} & ,r_j > R_j \end{cases}.$$ (12.1)

The utility of broadcasting a data item d is then given as

$$U_d = \sum_{j|d \text{ serves } Q_j} u_j.$$ (12.2)

For a given schedule S that broadcasts the data items $D_1, D_2, \ldots, D_k$ in order, the utility of the schedule is given as

$$U_S = \sum_{d=D_1}^{D_k} U_d.$$ (12.3)

In this work, we assume that the utility of a data item for a client decays by half for every factor of increase in response time, i.e.

$$\alpha_j = \frac{\ln 0.5}{R_j}. \tag{12.4}$$

12.2.3 Problem statement

A data source D is a set of N data items, $\{D_1, D_2, \ldots, D_N\}$, with respective sizes $d_1, d_2, \ldots, d_N$. A request queue at any instance is a dynamic queue Q with entries Q_j of the form $\langle D_j, R_j, P_j \rangle$, where $D_j \in D$, and $R_j, P_j \geq 0$. At an instance t_{curr}, let $Q_1, Q_2, \ldots, Q_M$ be the entries in Q. A schedule S at the instance is a total ordering of the elements of the set

$$\bigcup_{j=1,\ldots,M} \{D_j\}.$$

In the context of the broadcast scheduling problem, the request queue Q corresponds to the queue left after all requests currently being served are removed. Note that two different entries Q_j and Q_k in Q can have the same first component, i.e. $D_j = D_k$, for $j \neq k$; this implies that two different requests are interested in the same data item. However, it is important to realize that the current instance of the scheduler only needs to schedule a single broadcast for the data item. The arrival time of all requests in Q is at most equal to the current time, t_{curr}, and the scheduled broadcast time t for the data item in Q will be t_{curr} at the earliest. A schedule is thus a total ordering of the unique elements in all data items requested.

The time instance at which a particular data item from the schedule starts to be broadcasted is dependent on the bandwidth of the broadcast channel. A broadcast channel of bandwidth b can transmit b *data units* per *time unit*. If t_{ready} is the ready time of the channel (maximum of t_{curr} and the end time of current broadcast), then for the schedule $D_1 < D_2 < \ldots < D_k$, the data item D_i starts to be broadcasted at time instance $t_{D_i} = t_{ready} + \sum_{j=1}^{i-1}(d_j/b)$. All requests in Q for the data item D_i is then assumed to be served, i.e. $t_{j,ser}$ for such requests is set to t_{D_i}. We explicitly mention this computation to point out that the utility metric involves the access time, and not the tuning time, of a request. The access time is the time elapsed from the moment a client issues a request to the moment when it starts to receive the requested data item. The tuning time is the time the client has to actively scan the broadcast channel to receive the data item.

While Eq. (12.3) can be used to measure the instantaneous utility of a schedule, it is not suitable in determining the performance level of a solution method in a dynamic setting. We thus use the utility generated from the queries already served as the yardstick to compare performance. In other words, the performance of a solution methodology at an instance where queries $Q_1, \ldots, Q_K$ are already served is measured by $\sum_{j=1}^{K} u_j$. In effect, we are interested in the global utility generated by the method. The aforementioned QoS criteria could be a specification in terms of this global utility. The objective behind the design of a solution methodology is to then maximize this global utility measured at any instance.

12.2.4   Scheduling transaction data

The aforementioned problem assumes that requests are made at the data item level. A client interested in multiple data items would make independent requests for the individual items. As an extension of this problem, we also consider the case of transaction level requests. A transaction level request differs from a data item level request in the sense that a request involves more than one data item. The order of retrieval of the data items is not important, but processing at the client end cannot start until all data items are received.

The difference in problem formulation appears here in the definition of a query Q_j. An entry Q_j in the request queue now takes the form $\langle \mathcal{D}_j, R_j, P_j \rangle$, where $\mathcal{D}_j = \{D_{j_1}, \ldots, D_{j_n}\} \subseteq D$ and $R_j, P_j \geq 0$. The request is considered to be served at $t_{j,served}$, which is the time instance when the last remaining data item in $\mathcal{D}_j$ is broadcast. The utility generated by serving the request then follows from Eq. (12.2). Note that, with this formulation, scheduling at the transaction level is not a simple extension of the problem of scheduling an aggregation of multiple requests from the same client. The following factors highlight the differences in data item and transaction level scheduling.

1. Multiple data item level requests made from the same client can have different deadlines associated with different data items. However, the deadline specified in a transaction level request is specific to the transaction and not the individual data items required to complete it.

2. Each request made at the data item level would accrue some utility, irrespective of whether it came from the same client or not. The notion of dependency between the data items (to complete a transaction) is not embedded in the requests. A request at the transaction level specifies this dependency by collecting the required data items into a set and making a request for the entire set. Utility is accrued in this case only when all items in the set are served.

3. When requests are made independently, the scheduler is not required to maintain low levels of latency between broadcasts of data items requested by the same client. However, at the transaction level, the schedules generated must take the latency into consideration since the utility will keep reducing until all requested data items are retrieved by the client.

12.3   Solution Methodology

In the previous section, we identified that a solution methodology which can maximize the global utility is desired for the broadcast problem under study. However, it is often difficult to anticipate the incoming data access requests in a dynamic environment. Besides, an ongoing broadcast cannot be interrupted to accommodate higher utility from new requests. This restricts a solution methodology to focus on the current request queue only and make scheduling decisions that yield higher utility from the generated schedule. It can only be expected that a higher global utility measure would be obtained in the process.

Another constraint is the time factor involved in making a scheduling decision. Data requests usually arrive more frequently than they can be served, which in turn lead to the generation of a long request queue. Any scheduling method must be fast enough to generate a good schedule without adding much to the access time of requests. Latencies induced between broadcasts because of the scheduling time is also a detrimental factor to resource utilization. Heuristics are often used as fast decision makers, but may result in degraded solution quality. Hybrid approaches in this context can provide a suitable trade-off between solution quality and decision time.

Moreover, the assumption that a heuristic driven method is most appropriate in real time decision making can be flawed. This is specially true when the search space involved is well understood and specialized optimization methods can be devised to exploit the dynamics of the search space. In such situations, even a suboptimal solution generated by a carefully crafted optimization technique could be better than a heuristic based solution.

12.3.1 Heuristics

For the purpose of this study, we use two heuristics – *Earliest Deadline First* (EDF) and *Highest Utility First* (HUF) – which takes into account the time critical nature of a data request.

EDF starts off by first scheduling the data item which corresponds to a request with the minimum $t_{curr} - (t_{arr} + R_j)$. All requests in Q that get served by this broadcast are removed (all requests for the scheduled data item) and the process is repeated on the remaining requests. For multiple channels, the heuristic can be combined with *best local earliness* to map the data item to a channel that becomes ready at the earliest. EDF gives preference to data items that have long been awaited by some request, thereby having some level of impact on the utility that can be generated by the requests. However, it does not take into account the actual utility generated.

HUF alleviates this problem by first considering the data item that can generate the highest amount of utility. The strategy adopted by HUF may seem like a good approach particularly when the overall utility is the performance metric. However, HUF generated schedules may not be flexible enough in a dynamic environment. For example, if the most requested data item in the current request queue generates the highest utility, HUF would schedule the item as the next broadcast. If this data item requires a high broadcast time, not only will the subsequent broadcasts in the schedule suffer in utility, but new requests will also have to wait for a long time before getting served.

12.3.2 Heuristics with local search

As mentioned earlier, the emphasis in this work is understanding the performance of heuristics when coupled with local search techniques. We therefore introduce some

amount of local search to improve the schedules generated by the heuristics. The notion of local search in this context involves changing the generated schedules by a small amount and accept it as the new schedule if an improvement in the utility of the schedule is obtained. The process is repeated for a pre-specified number of iterations. Such a "hill climbing" approach is expected to improve the utility of the schedule generated by a heuristic. We employ the 2-exchange operator to search a neighborhood of the current schedule. The operator randomly selects two data items and swaps their positions in the schedule (Fig. 12.3). The notations EDF/LS and HUF/LS denote EDF and HUF coupled with local search respectively. Intuitively, these hybrid approaches should provide sufficient improvements over the heuristic schedules, w.r.t. the performance metric, as the local search would enable the evaluation of the overall schedule utility, often ignored when using the heuristics alone.

```
Old Schedule   : a b c d e f g h i j
Swap Pts       :   *     *
New Schedule   : a e c d b f g h i j
```

Figure 12.3: 2-exchange operator example.

12.3.3 $(2+1)$-ES

Evolution Strategies (ES) [22, 146] are a class of stochastic search methods based on computational models of adaptation and evolution. They were first suggested by Rechenberg during the late sixties. Most of the earlier work in ES did not present them as function optimizers, but rather as rules for automatic design and analysis of consecutive experiments to suitably drive a flexible system to its optimal setting.

Evolution strategies are typically expressed by the μ and λ parameters signifying the parent and the child population respectively. Whereas the algorithmic formulation of evolution strategies remains the same as that of a genetic algorithm [75], two basic forms have been defined for them. In the $(\mu + \lambda)$-ES, μ best of the combined parent and offspring generations are retained using truncation selection. In the (μ, λ)-ES variant, the μ best of the λ offspring replace the parents. These definitions are analogous to that of the steady-state and generational forms of genetic algorithms. The steady-state

---
Procedure 12.1 (2+1)-ES()

1: Generate two initial solutions x and y
2: Evaluate x and y
3: **repeat**
4:      Recombine x and y to generate an offspring z
5:      Mutate z with probability p
6:      Evaluate z
7:      Replace x and y by the two best solutions from $\{x,y,z\}$
8: **until** (termination criteria is met)
---

variant of genetic algorithms explicitly maintains the best solution found so far, while the generational variant blindly replaces the current population with the offspring population generated.

For our experimentation, we employ the $(\mu + \lambda)$-ES variant with $\mu = 2$ and $\lambda = 1$. This simple form of the ES is chosen to keep the dynamic scheduling time within an acceptable limit without sacrificing on the solution quality. Also, a $(2 + 1)$-ES can be seen as a type of greedy stochastic local search. It is stochastic because there is no fixed neighborhood and therefore the neighborhood does not define a fixed set of local optima. Otherwise, the method is very much a local search technique: sample the neighborhood and move to the best point. The pseudo code for the algorithm is given in Procedure 12.1.

12.3.3.1   Solution encoding and evaluation

For the data broadcasting problem mentioned in the previous section, a typical schedule can be represented by a permutation of the unique data item numbers in Q. Thus, for n unique data items, the search spans over a space of $n!$ points. In the presence of multiple channels (T say), a similar encoding can be obtained by using -1 as a delimiter between the schedules in different channels (Fig. 12.4) [140]. The evaluation of a solution involves finding the utility of the represented schedule as given by Eq. (12.3). The higher the utility, the better is the solution.

```
Channel: |--1--|  |-2-|   ....    |--T--|
Encoding: 3 1 4 -1 2 6 -1 .... -1 8 5 9
```

Figure 12.4: Solution encoding for multiple channels.

12.3.3.2 Syswerda recombination

Recombination operators for permutation problems differ from usual crossover operators in their ability to maintain the uniqueness of entries in the offspring produced. For a schedule encoded as a permutation, it is desired that recombination of two schedules does not result in an invalid schedule. A number of permutation based operators have been proposed in this context [159]. We employ the Syswerda recombination operator in this study. The operator is particularly useful in contexts where maintaining the relative ordering between entries is more critical than their adjacency. For a broadcast schedule, the relative ordering of data items in the schedule affect the time instance when a particular request gets served and hence influence the overall utility of the schedule.

```
x        :  a b c d e f g h i j
y        :  c f a j h d i g b e
Key Pos. :      * *     *   *
Offspring:  a j c d e f g h i b
```

Figure 12.5: Syswerda recombination.

The operator randomly selects several key positions and the order of appearance of the elements in these positions are imposed from one solution to the other. In Fig. 12.5, the entries at the four key positions from y, $\{a,j,i,b\}$, are rearranged in x to match the order of occurrence in y. The offspring is x with the rearranged entries.

12.3.3.3 Mutation using insertion

The elementary mutation operators define a certain neighborhood around a solution which in turn dictates the number of states which can be reached from the parent state in one step [22]. Insertion based mutation selects two random positions in a sequence and the element at the first chosen position is migrated to appear after the second chosen position (Fig. 12.6).

12.3.3.4 Initialization and termination

The initial solutions determine the starting points in the permutation space where the search begins. Thus, a good solution produced by EDF or HUF could be a choice

```
Parent     : a b c d e f g h i j
Mutate Pts:  *      *
Offspring : a c d e b f g h i j
```

Figure 12.6: Mutation using the insertion operator.

for the purpose. However, we did not want to add the complexity of determining an EDF or HUF schedule to the search process, and hence generated the initial solutions x and y randomly. The ES algorithm is terminated after a fixed number of iterations. If schedules generated by other heuristics are taken as initial solutions, the termination criteria could as well be specified as the point where a particular level of improvement has been obtained over the starting schedules. It is important that an alternative is also suggested since improvement based termination may never get realized.

12.3.4 Heuristics for transaction scheduling

In order to perform transaction level scheduling, the heuristics are slightly modified. The EDF heuristic for transaction level scheduling, denoted by T-EDF, sequentially schedules all data items in a request by giving preference to the request having the earliest deadline. The HUF heuristic for transaction level scheduling, denoted by T-HUF, sequentially schedules all data items in a request with preference to the one which can generate the highest amount of utility. The modifications enable the heuristics to make scheduling decisions based on the deadline, or utility, of serving a request, rather than the data items contained in it.

One might argue that T-EDF and T-HUF can be represented by EDF and HUF respectively by considering each request to be for a single data item whose size is given by the total size of the data items requested. This is a viable representation since T-EDF and T-HUF do not take the data items contained in a request into account. However, note that the schedule generated is always at the data item level. Hence, there is room to exploit any existing overlaps in the data items of two different requests. For example, consider the requests A for data items $\{D_1, D_2, D_3, D_4\}$ and B for data items $\{D_2, D_4\}$, with every data item having the same size. Representing A as a request for a data item D_A (whose size is the sum of the sizes of D_1, D_2, D_3 and D_4) and B as a request for a data item D_B

loses the fact that the data items in B are a subset of those in A. Further, exploiting the existence of such subsets is also not trivial. This is specifically true for T-EDF which makes decisions based solely on the deadline. If A has an earlier deadline than B then the schedule generated will be $D_1 < D_2 < D_3 < D_4$; otherwise $D_2 < D_4 < D_1 < D_3$. Observe that, in the former case, both requests A and B will be served at the same time instance (the instance when D_4 starts broadcasting), while in the latter case, B will get served earlier. Hence, irrespective of the deadlines, the latter schedule is always equal to or better than the former. It is possible that even T-HUF fails to exploit the relationship. If serving A first generates higher utility than serving B first, T-HUF would generate the former schedule, thereby pushing the finish time of B further in the time line. It misses the observation that by serving B first it can still maintain the same utility for A. The situation is further complicated when the data items have varying sizes. Given such observations, augmenting the heuristics with local search would allow for an interesting exploration at the data item level. Recall that the local search would attempt to improve on the heuristic generated schedule by swapping data items at different positions. The corresponding local search variants for T-EDF and T-HUF are denoted by T-EDF/LS and T-HUF/LS respectively.

The $(2+1)$-ES does not make a distinction between data item level and transaction level requests. It always operates at the data item level irrespective of how the request was made. Similar to the earlier case, the $(2+1)$-ES here first determines the set of data items to be scheduled in order to serve all requests in the queue – $\bigcup \mathcal{D}_j$. It then generates a schedule with these items to maximize the utility. Since requests are served only when all requested data items are received by a client, random initial solutions are likely to disperse dependent data items (belonging to the same transaction). Hence, we modify the initialization strategy to direct the search from a favorable region in the search space. The modification creates the initial solutions by choosing random requests from the queue and sequentially laying out the data items contained in the requests on the schedule. For example, if $\mathcal{D}_1 = \{D_1, D_2, D_3\}$ and $\mathcal{D}_2 = \{D_3, D_2\}$ are two transactions on the request queue, and transaction 1 is randomly chosen under this process, then data items D_1, D_2 and D_3 will be consecutive on the initial schedule. The process generates a favorable

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Table 12.1: Experiment parameters.

Parameter	Value	Comment
N	300	Number of data items
d_{min}	5KB	Minimum data item size
d_{max}	1000KB	Maximum data item size
b	120 KB/s	Broadcast channel bandwidth
m	60s/180s	Request response time mean
σ	20s/40s	Request response time std. dev.
r	5	Request arrival rate
r_T	5	Transaction size Poisson rate
s	0.8	Zipf's law characterizing exponent
P	Low(1), Medium(2), High(4)	Priority levels
p	0.5	Mutation probability
Gen	1000	Iterations for local search and ES

arrangement of the data items for some of the requests. The remaining functionality of the ES is maintained as described in Section 12.3.3.

12.4   Experimental Setup

The data sets used in our experiments are generated using various popular distributions that are known to capture the dynamics of a public data access system. The different parameters of the experiment are tabulated in Table 12.1 and discussed below. We generate two different data sets with these parameter settings, one involving data item level requests and another involving transaction level requests. Experiments are run independently on the two data sets using the corresponding heuristics and the $(2+1)$-ES.

Each data set contains 10,000 requests generated using a Poisson distribution with an arrival rate of r requests per second. Each request consists of an arrival time, data item number (or set of data item numbers for transaction level requests), an expected response time, and a priority level. For transaction level requests, the number of data items contained in a transaction is drawn from another Poisson distribution with rate r_T items per transaction.

We would like to point here that the number of data items (N) to be handled by a scheduler need not be very big in a pervasive setup. Each broadcast station would

typically be responsible for catering requests involving a sub-domain of the entire setup, in which case its data domain would also be comparatively smaller. The data items requested are assumed to follow the commonly used *Zipf*-like distribution [27] with the characterizing exponent of 0.8. Under this assumption, the first data item becomes the most requested item, while the last one is the least requested. Broadcast schedules can be heavily affected by the size of the most requested data item. Hence, we consider two different assignments of sizes to the data items from existing literature [80, 107]. The *INC* distribution makes the most requested data item the smallest in size, while the *DEC* distribution makes it the largest in size.

$$INC : d_i = d_{min} + \frac{(i-1)(d_{max} - d_{min} + 1)}{N}, \qquad i = 1, \ldots, N \tag{12.5}$$

$$DEC : d_i = d_{max} - \frac{(i-1)(d_{max} - d_{min} + 1)}{N}, \qquad i = 1, \ldots, N \tag{12.6}$$

Expected response times are assigned from a normal distribution with mean m and standard deviation σ. The particular settings of these parameters in our data item level experiment results in expected response times to be generated in the range of $[0, 120]$s with a probability of 0.997. For transaction level requests, the mean and standard deviation are changed so that the interval changes to $[60, 300]$s. Any value generated outside the range of the intervals is changed to the lower bound of the corresponding interval.

We use three different priority levels for the requests – *low, medium,* and *high*. Numeric values are assigned to these levels such that the significance of a level is twice that of the previous one. Since the total utility is related to the priority of the requests, we make assignments from these levels in such a way that the maximum utility obtainable by serving requests from each priority level is probabilistically equal. To do so, we use a roulette-wheel selection [75] mechanism which effectively sets the probability of selecting a priority level as: $P(Low) = \frac{4}{7}$, $P(Medium) = \frac{2}{7}$, and $P(High) = \frac{1}{7}$. Such a scheme restricts the scheduler from favoring high priority requests unless the utility generated by them is in fact higher than the cumulative utility from low priority requests.

Workloads on the scheduler can be varied by either changing the request arrival rate, or the channel bandwidth. We use the latter approach and specify the bandwidth used

wherever different from the default.

The local search for EDF/LS, HUF/LS, T-EDF/LS and T-HUF/LS are run for *Gen* number of iterations. For $(2+1)$-ES, the same number of iterations is chosen as the termination criteria. The number of iterations has been fixed such that the scheduling time is not more than 0.01s (on a 2.4 GHz Pentium 4 with 512 MB memory running Fedora Core 7) when the request queue contains requests for around 150 unique data items on the average. This is critical so that the runtime of the iterative algorithms do not impose high latencies in the decision making process. The performance of each method is measured at the time instance when all the requests get served. In other words, the performance of each method is given by the sum of the utilities generated by serving each of the requests in the data set.

12.5 Empirical Results

We first present the overall performance results obtained for the five different solution methodologies on the data item level data set. Although, the data item level scheduling problem can be viewed as a special case of the transaction level scheduling, we observed that a method's performance can be quite different in these two problem classes.

12.5.1 Data-item level scheduling

Fig. 12.7 shows the performance in terms of the percentage of maximum total utility returned by using each of the methods. The maximum total utility is obtained when every request is served within its expected response time, in which case it attains an utility equal to its priority level. Thus, summing up the priorities of all requests gives the maximum total utility that can be obtained.

For the *INC* data size distribution, HUF, HUF/LS and ES have similar performance. Although, EDF and EDF/LS have a slightly lower performance, both the methods do reasonably well. A major difference is observed in the amount of improvement gained by EDF by using local search, as compared to that of HUF. This is because EDF does not take into consideration the utility factor of requests and hence performing a local search based on utility results in a substantial level of improvement. HUF does reasonably well

Figure 12.7: Percentage of maximum total utility obtained by the solution methods. The utility obtainable suffers when the most requested data item is the largest in size.

in creating the initial schedules; hence local search does not provide significant additional improvement. Further explanation on this issue is presented later.

12.5.1.1 EDF vs. HUF

Differences arising in the performance of EDF and HUF can be explained using Fig. 12.8. The top row in the figure shows the utility obtained by serving a request. Clearly, the accumulation of points is mostly concentrated in the [0,0.5] range for EDF (left). For HUF (right), three distinct bands show up near the points 4, 2, and 1 on the y-axis. A high concentration of points in these regions indicate that a good fraction of the requests are served within their response time requirements. Moreover, even if the response time requirement could not be met, HUF manages to serve them with a good utility value. The figure confirms this point as the band near 0 utility in HUF is not as dense as in EDF.

The bottom row in Fig. 12.8 plots the utility of the requests served during a particular broadcast. We notice the presence of vertical bands in EDF (left) which shows that a good number of requests get served by a single broadcast. This is also validated by the fact that EDF does almost half the number of broadcasts as done by HUF (right). For an intuitive understanding of this observation, consider the instance when a broadcast is ongoing. Multiple new requests come in and accumulate in the queue until the current broadcast

Figure 12.8: Utility derived by using EDF and HUF with *INC* data size distribution. Top: Utility obtained from each of the request for EDF (left) and HUF (right). Bottom: Utility of requests served during a broadcast for EDF (left) and HUF (right).

ends. Most of these requests would be for data items that are more frequently requested. When EDF generates a schedule for the outstanding requests, it gives preference to the request which is closest to the deadline, or has crossed the deadline by the largest amount. Since the queue will mostly be populated with requests for frequently requested items, chances are high that EDF selects one of such requests. Thus, when a broadcast for such an item occurs, it serves all of the corresponding requests. This explanation is invalid for HUF since preference is first given to a data item that can generate the most utility. Since the data item with highest utility may not be the most requested one, more broadcasts may be needed for HUF.

Further, when the request queue gets long, EDF's preference to requests waiting for a long time to be served essentially results in almost no utility from serving that request. If we extend this observation into the scheduling decisions taken over a long time, EDF will repeatedly schedule older and older requests. If the queue size continues to grow, this essentially results in EDF failing to generate any substantial utility after a certain point of time. This is clearly visible in Fig. 12.8 (bottom) as the early drop in the utility level of broadcasts to zero. The vertical bands in EDF suggest that the request queue size did grow to a size where a single broadcast took care of multiple requests.

12.5.1.2   Impact of local search

The impact of performing the local search improves the EDF and HUF results for the *DEC* distribution – up to almost 75% to 100%. Recall that the *DEC* distribution assigns the maximum size to the most requested data item. If a schedule is not carefully built in this situation, there could be heavy losses in utility because other requests are waiting while the most requested data item is being broadcast. It is important that the scheduler does not incorporate the broadcast of heavy data items too frequently into its schedule and instead find a suitable trade-off with the induced utility loss. Unfortunately EDF and HUF generated schedules fail to maintain this sought balance. To illustrate what happens when local search is added, we refer to Fig. 12.9.

To generate Fig. 12.9, we enabled the local search mechanism when the 3000th scheduling instance with EDF is reached. At this point, the request queue contained requests for

Figure 12.9: Improvements obtained by doing 50000 iterations of the 2-exchange operator for the EDF schedule generated during the 3000th scheduling instance. The *DEC* data size distribution is used here. A newly obtained solution is considered better only if it exceeds the utility of the current solution by at least 20.

127 unique data items. The local search mechanism use the 2-exchange operator to generate a new schedule. To start with, the 2-exchange operator is applied to the EDF schedule to generate a new one. If the utility improves by more than 20, the new schedule is considered for any further 2-exchange operation. The process is repeated for 50000 iterations. The plot shows the factor of improvement obtained from the EDF schedule at each iteration of the local search. The factor of improvement is computed as *utility of solution / utility of EDF solution*. A gray point (x,y) in the plot illustrates that the factor of improvement for the solution obtained in the xth iteration of the search is y. The solid line joins the points where a generated schedule had an utility improvement of 20, or more, over the current one.

The horizontal bands in the figure illustrate the fact that the schedule space is mostly flat in structure. For a given schedule, most of its neighboring schedules (obtained by the 2-exchange operator) have, more or less, equal amounts of utility associated with them. Hence the improvement factor values accumulate around a region to generate the bands. A more interesting observation is the amount of improvement obtained across the different iterations. We see that the schedule utility improves to a factor of 2.5 within the first 1000 iterations of the local search (inset Fig. 12.9), after which the progress slows

Figure 12.10: Empirical cumulative density function of the % success rates of performing 100 independent iterations of the 2-exchange operation in each scheduling instance with EDF (left) and HUF (right). A success means the operation resulted in a better schedule.

down. This implies that as the schedules become better, the local search mechanism finds it difficult to further improve the schedule (observe the long horizontal band stretching from around the 15000 to the 50000 iteration). Hence, local search mechanisms are only useful when the initial schedules generated by the heuristics are not "good" ones.

Since the results indicate that EDF and HUF both gain substantial improvement with local search, it can be inferred that their schedules have much room for improvement when the utility measurement is the metric of choice. This is evidenced in Fig. 12.10. To generate the plots, the space near an EDF (HUF) generated schedule is sampled 100 times. Each sample is obtained by applying the 2-exchange operator to the heuristic generated schedule. A success is noted if there is an increase in utility of the schedule. With the success rate (number of success/number of samples) obtained in all the scheduling instances for the 10000 requests, an empirical cumulative density function is constructed. A point (x, y) in the plot evaluates to saying that in y fraction of the scheduling instances, a better schedule is obtained 0 to x times out of the 100 independent samples taken. For EDF, about 84% (97% − 13%) of the schedules have been improved 40 to 60 times. This high success rate for a majority of the schedules generated indicate that EDF is not a good heuristic to consider in the context of utility. HUF displays a similar increase in utility, but with a relatively lower success rate. HUF schedules generate higher utilities than EDF

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schedules and hence the success rate is low.

The observations leave us with the following conclusions. The nature of the search space tells us that significant improvements can be obtained by using local search on heuristic schedules, specially when the schedule utilities are substantially lower than what can be achieved. We observe that EDF and HUF in fact generate schedules that are not difficult to improve with a single swap of the data item positions. Thereby, combining local search with both the heuristics enable us to at least "climb up" the easily reachable points in the search space.

12.5.1.3   HUF/LS vs. $(2 + 1)$-ES

Our justification as to why local search with an operator like 2-exchange fails after a certain extent is based on the fact that these operators are limited by the number of points they can sample – the neighborhood. This can also be verified from Fig. 12.9, where the appearance of thin bands are indicative of the low sampling of the area. As schedules become better, much variation in them is required to obtain further improvement. Mutation based operators are designed to limit this variation while recombination can sample the search space more diversely [21, 84]. This is the primary motivation behind using a recombinative ES to get improved schedules.

To analyze the performance differences in HUF/LS and ES, we make the problem difficult by reducing the bandwidth to 80 KB/s. The *DEC* data size distribution is used and the broadcast frequencies for the 300 data items are noted. Fig. 12.11 (top) shows the frequency distribution. A clear distinction is observed in the frequencies for average sized data items. Recall that HUF first schedules the data item with the highest utility. However, it fails to take into account the impact of broadcasting that item on the utility that can be generated from the remaining requests. Since a majority of the requests are for the larger data items, it is highly likely that such items get scheduled more frequently. As a result most of the bandwidth is used up transmitting heavy data items. The impact of this is not felt on small data items as they are not requested often. However, for average data items which do have a substantial presence in the requests, utility can suffer. The difference between the HUF/LS schedule and ES schedule appears at this point.

Figure 12.11: Top: Broadcast frequency of the N data items for ES (left) and HUF/LS (right). Bottom: Fraction of maximum utility obtained from the different broadcasts for ES (left) and HUF/LS (right). The *DEC* distribution is used with a 80 KB/s bandwidth.

HUF schedules broadcast average sized items too infrequently, which implies that most requests for them wait for a long time before getting served. ES schedules have a comparatively higher frequency of broadcast for such data items, thereby maintaining a trade-off between the utility loss from not servicing frequently requested items faster and the utility gain from servicing average data items in an uniform manner. As can be seen from Fig. 12.11 (bottom), the pitfalls of the absence of this balance in HUF/LS is observed after a majority of the broadcasts have been done. HUF/LS schedules do perform better in maintaining a good fraction of the maximum utility during the initial broadcasts (notice that majority of the points are above the 0.2 limit on the y-axis prior to the 600th broadcast). Much of the difference in performance arises because of the utility losses resulting after that. In contrast to that, ES schedules consistently balance losses and gains to perform well almost till the end of the last broadcast.

This insight suggests that heuristics that can take into consideration the expected broadcast frequency of data items and their relative sizes should do well in the context of data utility. Probabilistic [179] and disk-based [1] broadcasts employ these notion for

Figure 12.12: Percentage of maximum total utility obtained by the solution methods for transaction level scheduling. Local search results in substantial improvement in the heuristic generated schedules.

push based architectures. It is thus worth investigating how they can be tailored for on-demand architectures. Further, balancing the utility losses and gains is a crucial aspect that any well performing heuristic needs to take into consideration.

12.5.2 Transaction level scheduling

Fig. 12.12 shows the performance of the solution methods on the transaction level scheduling problem in terms of the percentage of maximum utility attained. Recall that the maximum utility is the sum of the priorities for the requests in the data set. Similar to the data item level scheduling, the problem is not difficult to solve for the *INC* type data size distribution. Hybrid methods involving the use of local search to a heuristic generated schedule appear to be particularly effective. The problem is comparatively difficult to solve with the *DEC* type distribution. Nonetheless, local search provides substantial improvement on the quality of the heuristic generated schedules in this case as well. For both distributions, the $(2 + 1)$-ES generate comparatively lower utility schedules for this problem. An interesting observation is that, unlike for data item level scheduling, T-EDF and T-HUF both perform equally well here. Further explanation on these observations is provided later.

Figure 12.13: Whisker plots of response time for earliest deadline and highest utility heuristics on transaction level scheduling with the *INC* distribution. T-EDF fails to distinguish between requests of different priority levels. T-HUF makes the distinction on the average but response times are often high. Local search suitably modifies the schedules in both cases.

12.5.2.1 Performance on *INC*

The *INC* type distribution assigns the smallest size to the most requested data item. With this distribution, most requests have smaller data items in the requested set. Typical bandwidth values, as used in these experiments, thus allow multiple (and different) requests to be served in a relatively short span of time. Most difficulties in scheduling arise when sporadic broadcasts of bigger data items are made and the request queue grows in this duration.

Fig. 12.13 shows whisker plots of the response times for the earliest deadline and the highest utility heuristics, along with their local search variants, when applied on the data set with the *INC* distribution. The response time of a request is the difference in time between its arrival and the broadcast of the last remaining data item in the requested set. Note that, despite the high variance in response time, T-HUF manages a higher utility than T-EDF. The only visible factor better in T-HUF is the median value of these response times. We thus focus our attention to this statistic. Using the median response time as the metric, we see that T-EDF maintains similar values across requests of different

priorities. It fails to identify that high priority requests can generate more utility and hence the data items in such requests should receive some level of preference over low priority requests. The observation is not surprising since T-EDF's efforts are restricted to the deadline requirement only. In effect, there is no distinction between two requests with the same amount of time remaining to their deadlines, but with different priorities. T-HUF makes the distinction clear and further differentiates between such requests with the added advantage of being able to identify broadcasts that might serve multiple low priority requests instead of just a single high priority request.

Although T-EDF and T-HUF's differences come from the ability to distinguish between priority levels of requests, it does not seem to be the only factor affecting the utility. Local search on these methods result in substantial improvement. The contributing factor to this is the low median response time maintained along with the priority demarcations. Note that T-EDF/LS and T-HUF/LS generate very similar distributions in the response times. As described in the hypothetical example in Section 12.3.4, both T-EDF and T-HUF do not take into account the effect of intermingling data items on the schedule utility. Often it is possible to defy the logic behind a heuristic to some extent without degrading the utility of the heuristic generated schedule. In certain cases, this can in fact result in a better schedule. By swapping data items across the schedule, local search effectuates better exploitation of the fact that commonly requested data items are present in a significant fraction of the requests in the queue. What is uncertain is whether the similar distributions in the response times is indicative of a local or a global optima.

12.5.2.2 Performance on *DEC*

A major difference between data item level scheduling and transaction level scheduling is in the number of data items that has to be broadcasted to serve the requests in the data set. With an average transaction size of 5 data items in each request, transaction level scheduling has to make 5 different broadcasts before a single request is served. Thus, heuristics in this problem must be able to exploit any overlaps between requests to be effective in utility. The problem is further complicated with the *DEC* type distribution. All requests in this case will contain at least one or more big data items as a result of

the Zipf distribution. If broadcast of such items (taking a long time to broadcast) serves a very small fraction of the requests, then utility will most likely suffer in the long run. Further, average or smaller sized data items will be requested with a less frequency, often by requests separated far apart in the time line. Once again, utility will suffer if a heuristic waits for a long time to accumulate requests for such data items with the objective of serving multiple such requests with a single broadcast. Thus, a heuristic must be able to balance these two aspects of broadcasting when working on transaction level scheduling with the *DEC* distribution.

Fig. 12.14 shows the broadcast frequency of the different data items on this problem. The frequency distribution is very dissimilar than in the case of data item level scheduling. T-EDF, which performs better than T-HUF here, manages to maintain an almost uniform distribution. In fact, if the first 20% of the bigger data items is not considered, all methods show a similar trend. The uniform distribution is an extreme form of the balance we seek as indicated above. T-HUF encounters a problem when seeking this balance and results in a marginal drop in performance as compared to T-EDF. In transaction level scheduling, requests which overlap significantly with parts of other requests are the ones that can generate the highest utility. The overlap mostly occurs in the frequently requested data items. T-HUF schedules such requests first without taking into account that delaying a broadcast for a commonly requested item can actually help accumulate more requests interested in the same data item. Unless a new request has better overlap features than the ones already existing in the queue, the T-HUF schedule undergoes very small changes and virtually remains consistent, specially in the first few data items in the broadcast queue, for a certain period of time. What disrupts the consistency is the accumulation of enough new requests to change the overlap features altogether. However, multiple data items (often the bigger ones) have already been broadcast by this time and T-HUF is needed to schedule another broadcast for them. In the case of T-EDF, this effect is overpowered to some extent by the deadline requirement. Whenever a new request with a smaller deadline and a lesser overlap with the existing requests arrive, T-EDF immediately gives preference to it. As a result, broadcast times of data items in the existing schedule is delayed, which helps serve more requests now. The frequency of broadcast of commonly

Figure 12.14: Broadcast frequency of the N data items for transaction level scheduling in the *DEC* distribution. Distribution of average and below-average sized data items reaches an uniform trend for high utility solution methods.

requested data items in T-EDF and T-HUF corroborates our justification.

Performance improvements in these heuristics when aided by local search is mostly attributable to the higher frequency of broadcasts for average and smaller sized data items. Recall that a request is served only when data items of interest are received by it, irrespective of the order of retrieval. Consider a request interested in the data items D_1, D_2, and D_3, with D_1 being a commonly requested item under the *DEC* distribution. Next, consider the two schedules – $D_1 < D_2 < D_3$ and $D_3 < D_2 < D_1$. In both cases, the client making the request completes retrieval of the data items at the same time instance. However, the latter schedule has the advantage of delaying the broadcast of D_1, which in effect improves the chances of it serving newly accumulated requests as well. Exploiting such strategies is not possible by T-EDF and T-HUF unless aided by an exchange operator of the kind present in the local search variant. The side effect of this is that infrequently requested data items will be more often broadcast. This is clearly visible in T-EDF/LS and T-HUF/LS.

12.5.2.3 Dynamics of Local Search

The ability to exploit the overlap features of requests makes local search a promising aid to the heuristic generated schedules in both data item level and transaction level scheduling. Recall from Fig. 12.9 that the local search is most effective when schedules can be easily improved by adopting a hill climbing approach. The search space becomes more and more flat as better schedules are discovered. Often, reaching this flat region is not difficult in data item level scheduling, both for the *INC* and *DEC* distribution. However, the dynamics are somewhat different in transaction level scheduling.

Fig. 12.15 shows the improvement gained over a T-EDF generated schedule during the 5000th scheduling instance in T-EDF/LS. The plot depicts the progress of local search in its search for a better schedule. The T-EDF generated schedule is closer to the flat region for the *INC* distribution than for the *DEC* distribution. Hence, improvements are often slow in the *INC* distribution. However, observe that for the *DEC* distribution, at the end of the 1000th iteration of local search, the improvements are still showing an increasing trend. This indicates that, given more iterations, local search can continue to

Figure 12.15: Improvements obtained by local search in T-EDF/LS during the 5000th scheduling instance. Improvements are minor when the T-EDF schedule is already "good" (*INC* distribution). Utility shows an increasing trend at the end of the 1000th iteration of the local search for the *DEC* distribution.

easily improve the schedules before hitting the flat regions. Given the diverse number of factors determining the utility of schedules in the *DEC* distribution, it is not surprising to see that T-EDF generated schedules are far from being optimal. Besides, a hill climbing strategy like local search can effectively exploit the dynamics of the search space to adjust these schedules to bring them closer to the optimal. The only hindrance we face in doing so is the right adjustment of the number of iterations without imposing a bottleneck in the running time of the scheduler. We provide some suggestions on how this can be done in a later subsection.

12.5.2.4 Performance of $(2+1)$-ES

Our primary motivation behind using $(2+1)$-ES in data item level scheduling is to enable a diverse sampling of the search space when a schedule's utility no longer show quick improvements. However, for transaction level scheduling, other factors starts dominating the performance of this method.

Transaction level scheduling leads to an explosion in the size of search space. For data item level scheduling, a request queue of size n with requests for distinct items has a schedule search space of size $n!$, whereas in transaction level scheduling with k items

Figure 12.16: Improvements obtained by local search and ES in the schedule generated by T-EDF during the 5000th scheduling instance. For transaction level scheduling, local search performs better in avoiding local optima compared to $(2+1)$-ES.

per transaction, the search space can become as big as $(kn)!$. A bigger search space not only requires more exploration, but can also inject a higher possibility of prematurely converging to a local optima. Fig. 12.16 depicts the exploration with local search and the $(2+1)$-ES. In order to generate the plots, we ran T-EDF until the 5000th request arrives. A schedule for the requests in the queue at this point is then generated by running T-EDF/LS and $(2+1)$-ES for 5000 iterations each. This guarantees that both methods are operating on the same queue (same search space) and for sufficient number of iterations. Utility improvements by the ES starts stagnating after the 2000th iteration, suggesting convergence towards a local optima. Local search, on the other hand, has a better rate of improvement before starting to stagnate. This is visible for both *INC* and *DEC* distributions.

The premature convergence in $(2+1)$-ES is mostly due to the loss in *genetic diversity* of the solutions. As more and more recombination takes place, the two parents involved in the method starts becoming more and more similar, and finally being unable to generate a diverse offspring. This phenomenon is what typically identifies the convergence of the method. However, given the bigger search space and the small population (two) involved in exploring it, the phenomenon results in the method getting trapped in the local optima present across the space. It is suggestive from the broadcast frequency of different data items (Fig. 12.14) that, on the overall, the method failed to balance the broadcast of different sized data items for the *DEC* distribution. Changing the μ and λ

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parameters, as well as the mutation probability, of the method usually helps resolve such premature convergence. However, one should keep the real time runtime constraints into consideration while experimenting with the parameters. Local search, on the other hand, does not face such a problem. Ideally, given enough number of iterations with random restarts, a hill climbing approach is more resilient in this regard.

12.5.3 Scheduling time

The number of generations allowed to local search, or ES, can affect the quality of solutions obtained and the time required to make scheduling decisions. In our experiments, this value is set so that an average request queue can be handled in a small amount of time. However, the average queue size will greatly vary from problem to problem, often depending on the total number of data items served by the data source. In such situations, it may seem difficult to determine what a good value for the number of iterations should be. Further, in a dynamic environment, the average queue length itself may be a varying quantity. Nonetheless, one should keep in mind that scheduling decisions need not always be made instantaneously. The broadcast time of data items vary considerably from one to the other. The broadcast scheduled immediately next cannot start until the current one finishes. This latency can be used by a scheduler to continue its search for better solutions, specially with iterative methods like a local search or an ES.

12.6 Conclusions

In this chapter, we address the problem of time critical data access in pervasive environments where the time criticality can be associated with a QoS requirement. To this end, we formulate a utility metric to evaluate the performance of different scheduling methods. The earliest deadline first (EDF) and highest utility first (HUF) heuristics are used in two problem domains – data item level scheduling and transaction level scheduling. In the data item level domain, our initial observation on their performance conforms to the speculation that HUF performs better since it takes into consideration the utility of requests while making scheduling decisions. Further analysis of the nature of the scheduling problem shows that EDF and HUF generated schedules can be greatly improved by

introducing a minor amount of local search to them. The impact of the local search is found to be a direct consequence of the schedules generated by these heuristics which are found to belong to a region of the search space from where obtaining improvements is not difficult.

The observations drawn from this understanding of the behavior of local search aided heuristics enable us to propose an evolution strategy based search technique that provides more variance to a simple local search. The utility induced by such a technique surpasses that of both EDF and HUF, and their local search variants. This result also shows that search based optimization techniques are a viable option in real time broadcast scheduling problems, and often suboptimal solutions generated by such a technique can be better than those obtained from a heuristic.

The transaction level domain appear to be a relatively harder problem to solve. Balancing the broadcast frequency of data items is important here and we observe that T-EDF performs better in doing so as compared to T-HUF. Local search still remains a promising candidate to boost the performance of these heuristics. For certain data size distributions, local search improvements can continue if allowed to run outside the runtime restrictions set in our experiments. Better strategies to trigger the scheduler is thus required. Transaction level scheduling also has different search space dynamics, often with an explosive size and the presence of local optima. The evolution strategy method is affected by this, often leading to lower utility schedules than local search. Further experimentation is required in this direction to see how the parameters of the ES can be changed to avoid susceptibility towards such situations.

Future work in this context may be inspired from the insights obtained from the analysis conducted on the $(2+1)$-ES. From an utility standpoint, we intend to explore the option of designing heuristics that pay special attention to factors like broadcast frequency and loss-gain trade-off during scheduling decisions. Also, when requests involve multiple data items which may undergo regular updates, the validity of a broadcast must also be taken into account. Timely delivery of a data item then has to consider a validity deadline as well.

CHAPTER 13

Scheduling Ordered Data Broadcasts

Many pervasive application domains involve clients interested in groups of related data items that can be processed one at a time following some order. Consider the traffic management system example introduced in the previous chapter. The system gathers current traffic information using sensors and disseminates it to drivers in real time on demand. Drivers use smart GPS navigation units to request road conditions so that they can be routed and re-routed along the best possible roads. The GPS unit periodically requests traffic information for the roads that are still remaining in its route plan. The traffic service needs to provide the road conditions in the same order that the roads are in the route plan. That is, if the current route plan is $\langle r_1, r_2, r_3, r_4 \rangle$, where r_i is a road identifier, then the traffic service should provide the information as $\langle rc_1, rc_2, rc_3, rc_4 \rangle$, where rc_i is the road condition for r_i. A scheduling problem occurs in such an application when the number of data access requests is larger than the bandwidth capacity of the server.

Various soft deadlines may also be imposed on the requested data items, which if not served within a specific time window may result in the data item having near zero utility when finally received. For example, from the time that a driver makes a request for traffic information on road r_i to the time the monitoring service reports back with a gridlock on r_i, the driver may already have missed a more convenient exit for an alternate route. Given the resource limitations, it is not always possible that the time constraints

of every incoming request be satisfied. Thus, a broadcast schedule is sought which can serve clients with as much utility as possible. The scheduling problem is more difficult in a real-time setting where generation of a high utility schedule has to respect run time constraints as well.

Although ordered queries have been studied in the mobile computing paradigm [35, 90, 87], schedule evaluation is rarely based on the utility of responses. In this chapter, we first propose an utility accrual method for data requests involving constraints on the order of the requested data items. Second, we define a modified version of the Syswerda recombination operator for use in methods with recombination. Finally, the utility function is used by an evolution strategy based schedule optimizer to evaluate the effectiveness of the schedules. We pay particular attention to the run time constraint of the scheduler and argue that simple variants of an evolution strategy can be employed to satisfy this requirement.

The remainder of the chapter is organized as follows. Section 13.1 presents the utility function and a formal statement of the problem. Section 13.2 discusses the search algorithms. Some empirical results are presented in Section 13.3. Finally, Section 13.4 concludes the chapter.

13.1 Ordered Data Broadcasts

A data source $D = \{D_1, D_2, \ldots, D_N\}$ is a set of N ordered sets (or data groups), where $D_j = \{d_{1j}, d_{2j}, \ldots, d_{N_j j}\}$ with N_j being the cardinality of D_j and $j = 1, \ldots, N$. All data items d_{ij} are assumed to be unique and are of equal size d_{size}. A request is considered to be "fully served" when the last data item in the requested data group is retrieved, otherwise it is considered "partially served". A request queue at any instance is a dynamic queue Q with entries Q_j of the form $\langle D_j, R_j \rangle$, $D_j \in D$ and $R_j \geq 0$. At an instance t_{curr}, let $Q_1, Q_2, \ldots, Q_M$ be the entries in Q. We define the notation $Wait[Q_j]$ to denote the data item that the request Q_j is currently waiting for. Further, we define $Rem[Q_j]$ as the ordered subset of data items that has been requested in Q_j but not yet received, i.e. $Rem[Q_j] \subseteq D_j$. A schedule is a total ordering of the elements in the multi-set $\bigcup_{j=1,\ldots,M} Rem[Q_j]$.

The time instance at which a particular data item from the schedule starts to be broad-

cast is dependent on the bandwidth of the broadcast channel. If t_{ready} is the ready time of the channel (maximum of t_{curr} and the end time of current broadcast), then for the schedule $d_{i_1j_1} < d_{i_2j_2} < \ldots < d_{i_pj_p}$, the data item $d_{i_kj_k}$ can be retrieved by an interested client at time instance $t_{i_kj_k} = t_{ready} + [(k-1)d_{size}/b]$. All requests Q_j in Q with $Wait[Q_j] = d_{i_kj_k}$ are then partially served, i.e. t_{i_kj} for such requests is set to $t_{i_kj_k}$, and $Rem[Q_j]$ is changed to $Rem[Q_j] - \{d_{i_kj_k}\}$. The request is fully served when $Rem[Q_j] = \phi$.

In order to facilitate soft deadlines for ordered data, we make the assumption that the utility of a data item received by a client decreases exponentially if not received within the expected response time. For request Q_j arriving at time T_j and involving the data group $D_j = \{d_{1j}, d_{2j}, \ldots, d_{N_jj}\}$, let $t_{1j}, t_{2j}, \ldots, t_{N_jj}$ be the time when the respective data items are retrieved by the client. The utility generated by serving the first data item is given as

$$u_j[t_{1j}] = \begin{cases} 1 & , t_{1j} - T_j \leq R_j \\ e^{-\alpha(t_{1j} - T_j - R_j)} & , t_{1j} - T_j > R_j \end{cases}. \tag{13.1}$$

The utility from the subsequent items is then given as, for $i = 2, \ldots, N_j$ and inter-item response time R_T,

$$u_j[t_{ij}] = \begin{cases} u_j[t_{(i-1)j}] & , t_{ij} - t_{(i-1)j} \leq R_T \\ u_j[t_{(i-1)j}]e^{-\alpha(t_{ij} - t_{(i-1)j} - R_T)} & , t_{ij} - t_{(i-1)j} > R_T \end{cases}. \tag{13.2}$$

The utility of a data item for a client decays by half as a function of the response time, i.e. $\alpha = \ln 0.5/R$, where $R = R_j$ for the first data item in the requested group and $R = R_T$ for any subsequent data item.

If all data items are broadcast in a timely manner, a maximum utility of 1 will be generated by each data item. However, when a data item's broadcast time exceeds its expected response time, not only will its utility drop, it will also influence the maximum utility that can be obtained from subsequent items. We then say that the utility generated by serving the request is given by the utility generated at the last item of the data group, or $U_j = u_j[t_{N_jj}]$. The quality of service desired in an application domain can be directly specified as a fraction of the utility derived from serving the requests. For a schedule S, generated to serve the requests $Q_1, Q_2, \ldots, Q_M$ in the queue, the utility generated by the schedule is the aggregation of the utilities for the requests in the queue, given as

$$U_S = \sum_{k=1}^{M} U_k. \tag{13.3}$$

13.2 Solution Methodology

A schedule can be represented by a permutation of the data items currently in the $Rem[\cdot]$ sets of requests. Since the same data item may be present multiple times in this permutation, a request identifier is attached to every data item. For the requests $Q_1, Q_2, \ldots, Q_M$ currently in the request queue, the data items in $Rem[Q_j]$ of a request are identified by the tuples $\langle j, d_{i_k j} \rangle$, where $d_{i_k j} \in Rem[Q_j]$. The first component of a tuple identifies the request for which it exists in the permutation, and the second component identifies the data item number. The request identifier is only used in the permutation and is not part of the broadcast for the data item. Hence, if a request is waiting on a data item, say $d_{i_{k_1} j_1}$, then upon broadcast it will be retrieved by the request irrespective of the request identifier attached to the data item in the tuple, j_1 in this case.

In terms of a solution methodology, we focus on simple stochastic local search (μ, λ)-ES and $(\mu + \lambda)$-ES variants by setting $\mu = 1$. We also experiment with a $(2 + 1)$-ES with recombination and a simple genetic algorithm (GA). The simplest ES methods employ a mutation operator only, which implies a low overhead on the running time of the scheduler in a dynamic setting. "Shift" mutation selects two random positions in a permutation and the element at the first chosen position is removed and inserted at the second chosen position. The second random position is chosen in a way such that the ordering constraints for the data item in the tuple is preserved.

Recombination for the $(2 + 1)$-ES and the GA is achieved by a modified version of the Syswerda order-based crossover operator [167]. Syswerda's operator chooses random positions on a parent for exchange with the other. However, doing so can disrupt the ordering constraints of the permutation on the offspring (Fig. 13.1). The modified operator, called the *constrained-Syswerda* operator, eliminates this problem by restricting the choice of positions to a random contiguous block instead.

A synthetic dataset containing 10000 requests is used to empirically evaluate the performance of a search algorithm. The requests are generated using a Poisson distribution with an arrival rate of 3 requests per second. Each request consists of an arrival time, data group number, and an expected response time for the first data item in the group. We

```
Valid Parent 1        :  a b c d e f g h i j
Valid Parent 2        :  a d e b c h i f g j
Cross Positions       :  *   *   *
Invalid Offspring     :  a b e d c f g h i j
```

Figure 13.1: A counter example for the Syswerda order-based crossover operator. Ordering constraints are $\{a,b,c\}$, $\{d,e,f,g\}$ and $\{h,i,j\}$; constraint $\{d,e,f,g\}$ is violated in the offspring.

consider 100 different data groups, the number of data items in each drawn from an exponential distribution with rate parameter 10. The bandwidth for the broadcast channel is set at $200KB/s$.

The data groups requested are assumed to follow the commonly used Zipf distribution [27] with the characterizing exponent of 0.8. Under this assumption, the first data group becomes the most requested one, while the last one is the least requested. Broadcast schedules can be heavily affected by the size of the most requested data group. Thus, we consider the two different assignments: INC – most requested data group has the smallest number of data items, and DEC – most requested data group is the largest in size [80, 107]. Each data item is of size $50KB$.

Expected response times for the first data item are assigned from a normal distribution with mean 60s and standard deviation 20s. Any negative value is replaced by zero. Response time for intermediate data items is set at 1s.

For different variants of the $(1 + \lambda)$-ES and $(1, \lambda)$-ES, we fixed the maximum number of function evaluations to 15000. The number of function evaluations is fixed at 5000 for the $(2 + 1)$-ES. The parameters for the GA is set as follows: population size = 100, number of function evaluations = 15000, crossover probability = 0.8, mutation probability = 0.05, and 2-tournament selection.

13.3   Empirical Results

We present the results obtained from different variants of the ES on the two different data group assignments – INC and DEC. The results are averaged for 20 runs for each variant. Fig. 13.2 shows the percentage global utility obtained by running $(1 + \lambda)$-ES

Figure 13.2: Percentage global utility for the different λ values in the ES. For the *INC* assignment, a high utility level is obtainable with $\lambda = 1$. For the *DEC* assignment, increasing the sampling rate λ and the number of generations results in improvement of the global utility.

Figure 13.3: Mean difference between time of request arrival and time of first data item broadcast for data groups of different sizes in the *DEC* assignment.

with different λ values. Recall that each request can have a maximum utility of 1. The percentage global utility is thus computed from the fraction of this maximum utility generated in the requests in the data set. For the *INC* type assignment, a $(1 + 1)$-ES yields an acceptably high ($> 90\%$) utility level. Given the bandwidth limit of $200KB/s$, up to 4 data items can be transmitted in a single time unit. In the case of an *INC* type assignment, this can serve at least 4 different requests for the frequently requested data groups. Note that the *INC* assignment has the most requested data group as the smallest in size, which is one data item in our experimental data set.

The *DEC* type assignment has 20 data items in the most frequently requested data group. Further, the *Zipf* distribution makes the larger data groups more often requested than the smaller ones. The $200KB/s$ bandwidth poses a hard bottleneck in this situation. Quite often, different requests for the same data group arrive at different times prohibiting the *Wait*[·] value of those requests to be the same. The $(1 + 1)$-ES fails to provide the same level of performance as it does for the *INC* assignment. Increasing the sampling rate λ to 3,5 and 10 show improvements up to 80% utility levels. Although increasing the number of generations to 2000 improved the utility level up to 86%, the number of function evaluations ($2000 \times 10 = 20000$) exceeded the maximum set limit of 15000.

The primary difference between the schedule utilities generated by $(1 + 1)$-ES for

1000 generations and $(1+10)$-ES for 2000 generations is attributable to the latency that the scheduler puts in between the time of arrival of a request and the time when the first data item for the request is broadcast. Fig. 13.3 shows the mean values of this latency for data groups of different sizes. The $(1+10)$-ES maintains a higher mean latency for the larger data groups, whereas the $(1+1)$-ES maintains a higher value for the smaller data groups. Given that most requests are for the larger data groups in the DEC assignment, postponing the first data item broadcast for such requests provides gaps in the schedule to serve pending (or partially served) requests. Recall that the expected response time is the highest for the first data item (between 0 and 120s). Once a client has received the first data item, the expected response time drops to $R_T = 1s$. Hence, by delaying the first data item broadcast for the most requested data groups, the $(1+10)$-ES maintains a better flexibility in serving pending requests.

For the experimental data set, a simple $(1+1)$-ES for 1000 generations is sufficient when the QoS requirement is not too high $(< 75\%)$. For the case when the utility requirement is higher, a higher sampling rate is desired. However, note that results obtained from running the different variants for 2000 generations (more function evaluations) do not always yield a high difference in the utility levels as compared to those from running the same variants for 1000 generations. This is observed in both the comma and plus variants of the ES.

Fig. 13.4a shows the changes in the objective function (schedule utility given by (13.3)) value during the scheduling instance when the 5000th request arrives. The scheduler is working with DEC and $\lambda = 5$. Further, at each iteration, the plot shows the utilities of 100 randomly sampled points (using shift mutation) near the current parent. Note that the rise in the utility of the schedule is faster during the first 250 generations and then slows down considerably. In other words, as better schedules are obtained, improvements are harder to find. This in turn makes the progress slower. Moreover, the randomly sampled points around the parent of the current generation show very small differences in the utility values. This makes it difficult for the ES to maintain a steady increase in the schedule utility.

Fig. 13.4b shows the percentage global utility generated by the $(2+1)$-ES and the

Figure 13.4: (a) Schedule utility progress during the 5000th scheduling instance with *DEC*. Utilities are also plotted for 100 points sampled around the parent of every generation. (b) Percentage global utility for $(2+1)$-ES and a genetic algorithm.

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GA. Performance of both methods is on a par with the stochastic search variants for the *INC* size distribution. The GA's performance on the *DEC* distribution is easily overpowered by a $(1+3)$-ES. Further, the GA does a maximal utilization of the allowed function evaluations of 15000. The $(2+1)$-ES achieves an utility of 83%, equivalent to that of the $(1+5)$-ES with 2000 generations. An important factor to consider here is the number of function evaluations used in the two methods – 5000 in $(2+1)$-ES compared to 10000 in the $(1+5)$-ES. Although this performance is marginally better (about 3%) than the $(1+5)$-ES with 1000 generations, we stress that even obtaining such marginal improvements is difficult with the *DEC* distribution.

The enhanced performance of $(2+1)$-ES is attributable to the additional exploration brought forth by the recombination operator. Local search methods do not display enough exploratory capabilities when stuck in a plateau of the search space. Recombination allows for a more diverse sampling in such cases, thereby resulting in a faster exploration through plateaus. The GA is expected to benefit from this as well. The cause of its poor performance is not well understood.

13.4   Conclusions

Pervasive computing applications often need to broadcast grouped data objects such that the elements in the group satisfy a user specified ordering constraint. In addition, objects not served within a specific window may end up having near zero utility. In this chapter, we introduce a method of utility accrual for grouped data objects and use it to evaluate the effectiveness of a schedule. We argue that evolution strategy is a viable methodology to maximize the utility of broadcasts given the run time constraints of the application. We investigate three different methods – a simple stochastic local search using $(1+\lambda)$-ES and $(1,\lambda)$-ES, a $(2+1)$-ES with a modified Syswerda recombination operator, and a genetic algorithm. Our experiments suggest that the generation of an optimal schedule when the most requested group has the highest number of data items is a difficult problem to solve, often requiring a longer duration of search. However, recombination based ES appears to be particularly effective. The $(2+1)$-ES with the proposed recombination operator demonstrates the potential to generate better schedules

without engaging in too many function evaluations, thereby providing a fair trade-off between the run time and utility objectives of a scheduler.

The problem considered in this chapter assumes that data items are unique across different data groups. However, there exists other problems in pervasive environments where the data groups can have common data items. We can investigate if the scheduling strategy identified here works equally well in such problem domains.

CHAPTER 14

Scheduling in Multi-Layered Broadcast Systems

Most of the earlier works in designing broadcast schedulers assume a centralized system where the broadcast server has local access to the data items [2, 7, 55, 107, 187]. This simplifies the problem since the broadcast scheduler need not take into consideration the time required to retrieve the data item while making scheduling decisions. However, a number of application domains exist where such centralized storage of data are not possible. Consider the example of a company that provides investment consulting and management services to its clients (a company like Morningstar®). One service provided by such a company is real-time, on-demand information on stocks, mutual funds etc., in the form of market indices. Typically, such a company does not generate this information itself but fetches it from other companies (Morningstar®, for example, gets a significant portion of this data from the company Interactive Data Corporation[SM]). With the current ubiquity of wireless computing devices, we can imagine that clients of the company seek and receive periodic market data on their mobile phones, PDAs and notebooks, and analyze the data to determine which investments are rewarding. Since multiple clients may seek the same data, it makes good business sense for the company to broadcast this data. The clients also perform financial transactions using such devices. Market indices change throughout the day and it is important to analyze such trends along multiple dimensions in order to perform trading. Thus, although queries from clients are implicitly

associated with deadlines, these can be considered soft. Even if the queries are served after their deadline, they still have some value. The overall goal of the company will be to satisfy as many requests as possible in a timely manner, keeping in mind that the data may need to be fetched from other sources.

Decentralized storage of data introduces a novel problem to the broadcast scheduler, specifically if the data items must be retrieved from a data server prior to broadcast. Typically, the scheduler has complete knowledge on the time required to broadcast an item and uses this information in deciding a schedule. However, when data have to be fetched from a data server, this knowledge assumes a stochastic form. A data server will usually be serving multiple broadcast servers following its own request management policy. Thus, the time required to fetch a data item is not known apriori. A broadcast scheduler then has to build a schedule based on stochastic information available about the retrieval time of data items.

In this chapter, we visit the problem of data broadcast scheduling under such a scenario. We model the problem as a case of *stochastic scheduling* and explore the performance of a number of heuristic based schedulers. We argue that heuristics used in deterministic scheduling may not perform well in a stochastic problem of this nature. We show how probability estimates on a data server's response times can be used in the design of better schedulers. To this end, we propose two heuristics – the *Minimum Deadline Meet Probability* and the *Maximum Bounded Slack Probability* heuristics – that are based on the request completion time distributions rather than their exact values. Further, we augment our observations with an analysis to understand the underlying principles of a good scheduler. Our results demonstrate that a better performing broadcast policy exploits factors such as bandwidth utilization, data sizes and access frequencies. This is often as a result of how the specific heuristic has been formulated.

While considerable attention has been paid to designing heuristics for broadcast scheduling in centralized data systems, heuristics for non-local data models are rare to find. Data staging concerns in broadcast scheduling were first highlighted by Aksoy et al. [8]. The authors focus on a decision problem where the data items are not readily available for broadcast. An *opportunistic scheduling* mechanism is introduced where data items avail-

able for broadcast are chosen instead. Further, *hint-based cache management* and *prefetching* techniques are proposed in order to decrease the fetch latency of a data item. However, these techniques do not utilize any information available on the fetch time of data items while determining the broadcast schedule. The focus in our work is to supplement such data staging mechanisms by informing the scheduler on the workload present on a data server and make decisions accordingly.

Traintafillou et al. study a similar problem in the context of disk access latencies [171]. The primary objective of their work is to study the interplay among different components of a broadcast system, namely the broadcast scheduler, disk scheduler, cache manager and the transmission scheduler. However, disk access latencies are typically much smaller than data retrieval latencies from a secondary source. In effect, the impact of data fetch times is amplified in decentralized storage systems. Further, unlike as in non-local data systems, the heuristics proposed in their work assume that disk service times are known apriori.

Hierarchical data dissemination is used by Omotayo et al. in the context where data updates are frequently pushed from one server to another [138]. The scheduling problem is explored at the *primary* server which needs to decide how frequently to broadcast updated data items to *secondary* servers. Since the primary server does not maintain any incoming request queue, there is no stochasticity involved on part of the secondary servers. Notably, stochastic scheduling has been explored in depth for job shop scheduling problems where execution time of jobs are not known apriori [50, 125].

The remainder of the chapter is organized as follows. Section 14.1 presents the broadcast architecture used in this study. The formal statement of the problem is also outlined here. The heuristics experimented with are discussed in Section 14.2. Section 14.3 introduces the two heuristics proposed in this chapter for stochastic broadcast scheduling. Section 14.4 outlines the details of the experimental setup and the synthetic data set used to evaluate the performance of the heuristics. Section 14.5 presents the results obtained and provides an extensive discussion on the performance of the heuristics. Finally, Section 14.6 summarizes the chapter.

14.1 Problem Modeling

Stochastic scheduling refers to the class of scheduling problems where the time required to complete a particular task is modeled as a random variable. Compared to deterministic scheduling where the time of completion of a task can be computed before it begins execution, the actual completion time in the stochastic case is known only after the task finishes execution. However, a probability distribution of the completion time is known (or can be computed) for such tasks. Data broadcast scheduling transforms into a stochastic problem when the data items to be broadcast are not available locally at the broadcast server. In such a scenario, the broadcast server has to retrieve the data items prior to initiating the broadcast, from *data servers* distributed over a network. The data servers on the other hand will be receiving multiple requests from other broadcast servers and will have their own policy to manage the serving of accumulating data requests. The serving of requests on part of the data server may be accomplished by a unicast, multicast or a broadcast mechanism. Hence, the time required by a broadcast server to fetch a data item will not be known until the item is actually fetched. Scheduling decisions on the broadcast server that must be made prior to fetching any data item will therefore be inaccurate. As an alternative, a broadcast server can use *response time* probability estimates of the data servers. The response time is the time elapsed between making a request and getting the request served. The probability estimates would present a broadcast server with likely durations of time it would require to retrieve a data item, thereby introducing the stochastic broadcast scheduling problem. A stochastic scheduler can use the probability estimates as prior knowledge on the problem instance, but does not utilize anticipated information from the future (such as a possible value for the response time) while generating a scheduling policy. Often times, the policy itself may undergo revisions as data items are actually fetched and their response times are known.

14.1.1 Broadcast server model

Fig. 14.1 depicts the broadcast architecture used in this study. Clients request different data items from a broadcast server using the uplink channel. Each request Q has an arrival time, a data item index and an absolute deadline, given by the tuple $\langle arr_Q, d_Q, dln_Q \rangle$.

Figure 14.1: Broadcast server architecture in non-local data model.

Although deadlines can be explicitly specified by the client, it is not a strict restriction. For example, the broadcast server can assign deadlines to the scheduler based on the current status (say a geographic location) of the client making the request. A single request involves only one data item. If multiple data items are required by a client, then multiple requests are formulated to retrieve them, with no constraint on the order in which they are to be served.

Data items are distributed across multiple data servers. The broadcast server has fast dedicated connections to these servers. The data servers have their own policies to serve incoming requests. The only knowledge the broadcast server has about the data servers are the data items they host and a probability mass function (PMF) of the response times of these servers. The PMFs can either be sampled from a known probability distribution function, or constructed by observing the response times of the data servers over a period of time.

The broadcast server reads the requests in the queue and invokes a scheduler to determine the order in which the data items are to be broadcast. Once a schedule is built, the broadcast server requests the corresponding data item from the respective data server and stores it in a buffer. For the sake of simplicity, we do not assume a system where the fetching of a data item and its broadcasting happens in parallel. Such an approach can raise synchronization issues which then have to be handled with multiple levels of buffering. Pre-fetching of data items (retrieving multiple items and holding in local storage) is also discarded in this model since the scheduler may decide to change the schedule to

accommodate more urgent requests, in which case pre-fetched data items may have to be discarded. Also, the application domains we consider are where data items may undergo frequent updates at the data server. Nonetheless, all these factors can be accommodated in the broadcast server for added complexity. The broadcast server initiates the broadcast for the data item in a downlink channel of bandwidth b_{bs} as soon as the entire data item has been received from the data server. Interested clients retrieve the data item from the downlink channel. A request is served at the time instance, called the *completion time* of the request, when the entire data item is received by the client.

The scheduler is invoked every clock tick. During periods when the broadcast server does not receive any new request, the scheduler revisits the current schedule at specified time intervals. This is done since the scheduler gains precise information on the response times of the requests with each completed broadcast and may decide to change the schedule (generated with stochastic information) over time.

14.1.2   Data server model

A data server hosts a number of data items and maintains dedicated connections to multiple broadcast servers. It is also possible that the data server does not locally host the requested data item, but just acts as a designated channel to retrieve the item. In such a case, the data server has to fetch the item from another server. This facet can introduce multiple layers in the data fetching scheme. However, in our model, such multi-layered fetching schemes remain transparent to the broadcast server. This is because the broadcast server can only acquaint itself with response time PMFs about the data server. When such a scheme is in place, the response time PMFs will appropriately reflect the delay introduced. In this study, we assume that data items are locally hosted at the data servers.

A data server is modeled similar to a $M/G/1$ queuing system with the processor sharing (PS) discipline. The $M/G/1$-*PS* queuing model assumes that requests arrive at a data server following an exponential distribution (Poisson in this case) and has arbitrary *service times*. The service time of a request is the time required to complete the request had it been the only one in the system. Completion of a request in this case implies

Figure 14.2: Data server architecture following a $M/G/1$-PS model.

transmission of the entire data item from the data server to the broadcast server. The bandwidth available at the data server is distributed uniformly to serve requests in a round-robin fashion. Hence, if there are n requests ready to be served in the queue and b_{ds} is the bandwidth available at the data server, then each request receives a quanta $\frac{b_{ds}}{n}$ of the bandwidth every clock tick. A request that gets completely served in the quanta leaves the system; otherwise it is cycled back into the queue for another quanta in the next clock tick. Fig. 14.2 depicts this architecture. Note that the quanta received by a request will vary every clock tick as new requests keep entering the system and completed requests leave.

14.1.3 Formal statement

A data server DS_i is a collection of N_i data items $D_i = \{d_1^i, d_2^i, \ldots, d_{N_i}^i\}$ such that $D = \bigcup_i D_i$ and $D_i \cap D_j = \phi$ for $i \neq j$. The null intersection enforces the condition that the same data item is not hosted by two different data servers, i.e., no replication. A broadcast server maintains a dynamic queue with requests arriving as a tuple $\langle arr_Q, d_Q, dln_Q \rangle$, where arr_Q is the arrival time of a request Q, $d_Q \in D$ is the data item requested and $dln_Q > arr_Q$ is the absolute deadline for the request. Further, let b_{bs} and b_{ds} be the bandwidth available at the broadcast and data server respectively.

At each scheduling instance, the scheduler first removes all requests from the queue that will be served by the current broadcast and generates a schedule for all remaining requests Q'. A schedule is thus a total ordering of the data items in $\bigcup_{Q'}\{d_{Q'}\}$. Let $ft(d)$ be the time required to fetch the data item d from the corresponding data server. If t_{ready} is the ready time of the broadcast channel (the time instance when any ongoing

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broadcast ends), $d_{i_1} \rightarrow d_{i_2} \rightarrow \cdots \rightarrow d_{i_p}$ the broadcast schedule and s_d be the size of data item d, then the broadcast of an item d_{i_k} ends at time $t_{d_{i_k}} = t_{ready} + \sum_{j=1}^{k} [ft(d_{i_j}) + \frac{s_{d_{i_j}}}{b_{bs}}]$. All requests Q' served by this broadcast then has a completion time $ct_{Q'} = t_{d_{i_k}}$ and response time $rt_{Q'} = (ct_{Q'} - arr_{Q'})$. The objective of the scheduler is to generate a schedule such that the response time of the requests do not exceed the limit set by the deadline, i.e., $rt_{Q'} \leq (dln_{Q'} - arr_{Q'})$, or $ct_{Q'} \leq dln_{Q'}$.

14.1.4 Performance metrics

We shall use three metrics – *deadline miss rate*, *average stretch* and *utility* – to evaluate the performance of different schedulers. Measurements are taken when all requests in the experimental data set have been served.

14.1.4.1 Deadline Miss Rate (DMR)

Let $\mathcal{Q}$ be the set of all requests served by the broadcast server at an instance of time. The deadline miss rate (DMR) is the fraction of requests in $\mathcal{Q}$ which missed their deadlines, given as

$$DMR = \frac{|\{Q \in \mathcal{Q} | rt_Q > (dln_Q - arr_Q)\}|}{|\mathcal{Q}|}. \tag{14.1}$$

14.1.4.2 Average Stretch (ASTR)

The stretch of a request is the ratio of its response time to its service time. Recall that the service time of a request is the time it would take to serve the request had it been the only one in the system. In a non-local data model, the service time should also include the retrieval time of the data item requested. Hence, if the request Q involves a data item of size s, to be retrieved from a data server with bandwidth b_{ds} and broadcasted over a channel with bandwidth b_{bs}, then the service time of Q is given by $st_Q = s(\frac{1}{b_{ds}} + \frac{1}{b_{bs}})$. The stretch of the request is then given as $STR_Q = \frac{rt_Q}{st_Q}$. A low stretch value indicates closer proximity of the response time to the minimum time needed to serve the request.

Intuitively, it seems that the response time of a request can never be less than the service time. Hence, the stretch of a request should always be greater than or equal to 1.0. However, when data fetching times are significant, the stretch can be less than one. This

can happen when a new request for a data item next scheduled for broadcast, and being currently retrieved from a data server, enters the broadcast server. In such a situation, the request realizes a retrieval time less than $\frac{s}{b_{ds}}$ and gets served as soon as the retrieval ends. Therefore, the response time becomes less than the service time. The average stretch (ASTR) is given as

$$ASTR = \frac{\sum_{Q \in \mathcal{Q}} STR_Q}{|\mathcal{Q}|} = \frac{1}{|\mathcal{Q}|} \sum_{Q \in \mathcal{Q}} \frac{rt_Q}{st_Q}. \tag{14.2}$$

Ideally, a low ASTR signifies that most requests are served with minimal deviation from their service times. Hence, most requests will meet their deadlines as well, resulting in a lower DMR.

14.1.4.3 Utility (UT)

The performance level depicted by DMR is useful when the deadlines imposed are hard. However, when soft deadlines are in place, a request being served after its deadline still holds some utility. Thus, the utility (UT) metric uses a function that maps the response time of a request to a real number. We use the following function in this context.

$$U_Q = \begin{cases} 1 & ,rt_Q \leq (dln_Q - arr_Q) \\ e^{-\alpha_Q(rt_Q - dln_Q + arr_Q)} & ,otherwise \end{cases}, \tag{14.3}$$

where $\alpha_Q = \ln 0.5/(dln_Q - arr_Q)$. Thus, the utility of a request is 1.0 if served by its deadline, otherwise decreases exponentially depending on the relative deadline of the request. The fractional utility attained in the system is then given as

$$UT = \frac{\sum_{Q \in \mathcal{Q}} U_Q}{|\mathcal{Q}|}. \tag{14.4}$$

An UT value of 1.0 indicates that all requests are served by their deadline; otherwise the closer it is to 1.0, the lesser is the time by which requests overshot their respective deadlines. Note that a high UT does not necessarily indicate a low DMR.

14.2 Schedule Generation

A major challenge in the generation of optimal schedules for data broadcasting is the lack of a formal theory underpinning the statistical characteristics of a "good" broadcast schedule. While queuing theory provides preliminary grounds for such analysis, no

attempt is known to have been made to understand a broadcast system that reflects a one-to-many relationship between servings and arrivals in the queue. Hence, a significant amount of focus is concentrated in designing heuristic methods that employ intuitive perceptions of good broadcast mechanisms. Much of this is also due to the additional constraint of scheduling time imposed by the real time requirement in on-demand systems. A typical workload condition may prohibit the use of time consuming methods in order to avoid long accumulation of requests and increases in their response times. Long scheduling times may also keep valuable broadcast bandwidth idle, resulting in inefficient utilization of available resources. To this end, heuristic driven schedulers are often preferred as fast decision makers.

Most of the existing heuristics in broadcast scheduling have been evaluated assuming systems with local data availability. For this study, we adopted four well-performing heuristics – RxW, MAX, SIN-α and $PRDS$ – proposed for such systems and observe their performance in non-local data availability systems. While no intrinsic property in these heuristics prohibit them from use in our problem model, modifications are made if required.

14.2.1 RxW

The RxW heuristic [7] combines the benefits of the MRF (Most Requested First) and FCFS (First Come First Serve) heuristics in order to provide sufficient weight to both heavily requested and long awaited data items. Owing to its simplicity, the RxW heuristic has a low overhead in terms of scheduling time. RxW schedule data items in decreasing order of their $R \times W$ values, where R is the number of pending requests for a data item and W is the time for which the oldest pending request for the data item has been waiting. Such a broadcast mechanism gives preference to data items which are either frequently requested or has not been broadcast in a long time (with at least one request waiting for it). This approach aims at balancing popularity and accumulation of requests for unpopular items. Hence, although the heuristic does not have any implicit factor that considers the deadline of requests, deadlines can be met by not keeping a request too long in the request queue.

14.2.2 MAX

The MAX heuristic [2] first assigns a hypothetical deadline to each request based on the maximum stretch value observed in the already served requests. For a request Q arriving at time arr_Q and with a service time st_Q, the hypothetical deadline is calculated as $(arr_Q + st_Q \times S_{max})$, where S_{max} is the maximum stretch observed till now in the served requests. Once the hypothetical deadline for all outstanding requests have been assigned, the MAX heuristic uses EDF (Earliest Deadline First) – the closer the hypothetical deadline to the current time, the higher the preference – to generate the schedule. Following the suggestions in the original work, the hypothetical deadline for a request is not changed once assigned even if the maximum stretch S_{max} is updated over time.

The MAX heuristic effectuates a scheduling mechanism that is targeted towards minimizing the maximum stretch value in the system. By doing so, MAX tries to maintain an optimal response time for requests, taking into account that different requests typically involve data items of different sizes. Note that the hypothetical deadline is not related to the actual deadline imposed on a request. Nevertheless, if MAX manages to generate schedules that prohibit the maximum stretch from increasing drastically, then requests will be served with minimal deviations from the minimum time required to serve them. Assuming that actual deadlines are imposed reasonably, the requests are then likely to complete within their deadlines.

14.2.3 SIN-α

The SIN-α (Slack time Inverse Number of pending requests) heuristic [187] integrates the "urgency" and "productivity" factors into a single metric called *sin.α*. The intuition behind the heuristic is explained by the authors using the following two arguments.

- Given two items with the same number of pending requests, the one with a closer deadline should be broadcast first to reduce request drop rate;

- Given two items with the same deadline, the one with more pending requests should be broadcast first to reduce request drop rate.

Based on these two arguments, the $sin.\alpha$ value for a data item (requested for by at least one client) is given as

$$sin.\alpha = \frac{slack}{num^\alpha} = \frac{1stDeadline - clock}{num^\alpha}, \qquad (14.5)$$

where *slack* represents the urgency factor, given as the duration from the current time (*clock*) to the absolute deadline of the most urgent outstanding request (*1stDeadline*) for the data item, and *num* (≥ 1) represents the productivity factor, given as the number of pending requests for the data item. The parameter α (≥ 0) can amplify or reduce the significance of the productivity factor while making scheduling decisions. With $sin.\alpha$ values assigned to the data items, the schedule is created in increasing order of the $sin.\alpha$ values. Note that the SIN-α heuristic does not take into account the different sizes of the data items involved in the requests. Hence, the heuristic does not differentiate between equi-probable data items with very different sizes. The α value in our experiments is set at 2 based on overall performance assessment presented in the original work. We shall henceforth refer to this heuristic as SIN2.

14.2.4 NPRDS

Apart from the two arguments provided for SIN-α, the PRDS (Preemptive Request count, Deadline, Size) heuristic [107] incorporates a third factor based on data sizes.

- Given two data items with the same deadline and number of pending requests, the one with a smaller data size should be broadcast first.

We modify PRDS into a non-preemptive version and call it NPRDS. Preemptive schedules are usually expected to perform better than non-preemptive ones. However, preemption has been discarded in this study to facilitate a fair comparison. Besides, preemptive scheduling in a non-local data model would either require the broadcast server to repeatedly request the same data item for the same set of requests (thus adding to the response time of the requests), or has to buffer all preempted data items. NPRDS works by first assigning a priority value to each data item (with at least one request for it), given as $\frac{R}{dln \times s}$, where R is the number of pending requests for the data item, dln is the earliest feasible absolute deadline (the broadcast of the item can serve the request before its deadline) of

Procedure 14.1 MDMP()

Output: A broadcast schedule.

1: $t_{cb} =$ time when the currently ongoing broadcast started
2: $s_{cur} =$ size of the item being broadcast
3: Initialize X_{bdst} such that $Pr(X_{bdst} = t_{cb} + \frac{s_{cur}}{b_{bs}}) = 1$. If no broadcast is currently taking place, then the initial PMF for X_{bdst} contains a single impulse at the current time, i.e. $Pr(X_{bdst} = current\ time) = 1$
4: **repeat**
5:    **for all** (pending request Q) **do**
6:       $X_Q = (X_{bdst} \odot X_{D_S(d_Q)}) \oplus \frac{s_{d_Q}}{b_{bs}}$
7:    **end for**
8:    Choose the request Q^* such that $Pr(X_{Q^*} \leq dln_{Q^*}) = \min\limits_{Q} Pr(X_Q \leq dln_Q)$ {*ties are broken by selecting the request which involves a data item that can serve a higher number of requests*}
9:    Schedule d_{Q^*} as the next broadcast item
10:    Mark all requests that would be served by d_{Q^*} as *"served"*.
11: **until** (all pending requests are marked *"served"*)

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outstanding requests for the data item, and s is the size of the item. A higher value of this priority estimate indicates that the data item is waited for by more number of requests, can help attain a tighter deadline and will not take much time to broadcast. Hence, the NPRDS schedule is generated in decreasing order of the priority values. The NPRDS heuristic aims at providing a fair treatment to different data sizes, access frequency of data items and the deadline of requests.

14.3 Scheduling with Stochastic Information

The heuristics outlined in the previous section do not utilize any information available on the response times of the data server. This information is assumed to be available in the form of a probability distribution. Next, we propose two novel heuristics that utilize such stochastic information to compute a completion time distribution for data items. Thereafter, scheduling decisions are made based on the probability of serving a request within its deadline.

14.3.1 MDMP

The MDMP (Minimum Deadline Meet Probability) heuristic can be considered a stochastic version of EDF. Under MDMP, the highest preference is given to the data item

corresponding to the request that has the minimum probability of meeting its deadline. In order to compute this probability, we first have to determine the completion time distribution of broadcasting a data item. In a situation where the response time of fetching a data item is deterministic, the completion time of a broadcast would simply be the addition of the ready time of the broadcast channel, the response time of fetching the data item and the time required to broadcast the item over the channel. A similar method is used for the case when the response time of fetching an item is stochastic. We begin with the response time PMF of the data server where the data item resides and combine it with the ready time distribution of the broadcast channel. Let $D_S(d)$ denote the data server where the data item d resides and X_k denote the random variable representing the response time of fetching a data item from data server k. If X_{bdst} denote the random variable representing the ready time distribution of the broadcast channel, then the completion time distribution of broadcasting data item d of size s_d is given by $X_{bdst} + X_{D_S(d)} + \frac{s_d}{b_{bs}}$. Hence, determining the completion time distribution requires us to find a way of adding random variables. This can be achieved by the convolution, denoted by $\odot$, of the PMFs of the corresponding random variables. Addition of a scalar to a random variable, denoted by $\oplus$, is equivalent to shifting the time axis of the random variable's PMF by the scalar amount. With these two operations, we can define the MDMP heuristic as shown in Procedure 14.1.

Note that the convolution of X_{bdst} with the data server response time PMFs can be precomputed before step 6 in order to reduce the time complexity of the heuristic. Also, proper resolution of ties may serve to be crucial when there exists multiple requests in the queue that have already missed their respective deadlines, i.e. $Pr(X_Q \leq dln_Q) = 0$. Hence, instead of arbitrarily choosing a data item from equal deadline meet probability requests, we choose the one that can serve a higher number of requests and reduce request accumulation.

14.3.2 MBSP

The MBSP (Maximum Bounded Slack Probability) heuristic can be visualized as the dual of MDMP with slight modifications. *Slack* in this case is defined as the duration from

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Procedure 14.2 MBSP()

Output: A broadcast schedule.

1: Initialize X_{bdst} as in Procedure 14.1
2: **repeat**
3:   **for all** (pending request Q) **do**
4:     $X_Q = dln_Q \ominus [(X_{bdst} \odot X_{D_S(d_Q)}) \oplus \frac{s_{d_Q}}{b_{bs}}]$
5:   **end for**
6:   Choose the request Q^* such that $Pr(X_{Q^*} \leq \mathcal{S}_{dev} \cdot [dln_{Q^*} - arr_{Q^*}]) = \max_{Q} Pr(X_Q \leq \mathcal{S}_{dev} \cdot [dln_Q - arr_Q])$ {*ties are broken by selecting the request which involves a data item that can serve a higher number of requests*}
7:   Schedule d_{Q^*} as the next broadcast item
8:   Mark all requests that would be served by d_{Q^*} as "*served*"
9: **until** (all pending requests are marked "*served*".)

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the deadline of a request to the time when it gets served. This slack value is positive if the request is served before the deadline, otherwise negative. MBSP schedules data items based on a lower bound for the slack value of requests. Since negative slack requests have already missed their deadlines, it might seem reasonable to give preference to ones which can still meet their deadlines. However, such a strategy can result in certain requests being pushed too far away from their imposed deadlines. Hence, MBSP employs a *slack deviation* parameter $\mathcal{S}_{dev}$ to extend the deadline of a request by a variable amount and measures the slack from this extended deadline. This is equivalent to imposing a lower bound on the slack value of requests. If a request misses this extended deadline too, then one can say that the request has been ignored for a long duration of time (after it missed the deadline) and should now be served. Thus, MBSP gives preference to requests with the maximum probability of missing the deadline extended by the slack deviation parameter, or in other words, to requests with the maximum probability of going below the lower bounded slack value. The heuristic can be computed as shown in Procedure 14.2.

The $\ominus$ operator in step 4 signifies the subtraction of a random variable X from a scalar quantity q_{scalar}. This operation results in a random variable $X' = q_{scalar} \ominus X$, such that $Pr(X' = q_{scalar} - x) = Pr(X = x)$. Note that if $\mathcal{S}_{dev} = 0$, then MBSP simply chooses the request with the maximum probability of missing its deadline. We have used the relative deadline of a request as the factor to scale in deciding the extended deadline. This is primarily based on an understanding of the utility derived from serving a request that

has already missed its deadline. Since we have assumed here that the drop in utility is related to the time overshot from the deadline, we adhere to this value in calculating the extended deadline as well. Other representations of utility may signify a different formulation to be more appropriate.

The novelty in MDMP and MBSP lies in the treatment of the stochastic information available about retrieval times of data items. A typical scheduler can compute the expected value from the response time distribution and use it as the time required to fetch any data item from the data server. Therefore, the scheduler assumes that a data item of any size can be fetched in the same amount of time, and ignores the varying workload in the data server. MDMP and MBSP work directly with the probability distribution and hence utilize convolution, instead of direct sums of broadcast and expected retrieval times, to find completion time distributions. The potential advantage of obtaining this completion time distribution lies in the fact that a scheduler can now compute the exact probability with which a particular request will be completely served within a given time period. To our knowledge, using the probability of successfully completing a broadcast to build a schedule has not been attempted earlier, and is an approach that diverges from typical deterministic methods.

14.4   Experimental Setup

The experimental setup used in this study consists of a single broadcast server and four data servers. Requests at the broadcast server arrive at a rate of $\lambda_{bs} = 5$ requests per second following a Poisson distribution. Each request consists of an arrival time, a data item number and an absolute deadline. A total of $100,000$ such requests are generated for inclusion in a synthetic data set. A data item is chosen for a request based on its size and popularity, as discussed in Sections 14.4.1 and 14.4.2.

We have written our custom tool to simulate the broadcast and data server interactions. The tool is a sequential control flow program that switches between a *broadcast server module* and a *data server module*, and measures the various time related statistics required in the performance metrics. Note that although our implementation has a sequential control flow, the execution of the data server module is designed in a way that

the data servers appear to be running in parallel to the broadcast server. Details of this implementation is given in Section 14.4.7. Owing to the nature of the implementation, a single machine is sufficient as a host for the tool to measure the relevant time statistics (completion time of requests). Different data servers are only instantiations of the data server module.

14.4.1 Data item popularity

Data items are requested following the Zipf-like distribution with a characterizing exponent of $\beta = 0.8$. A larger exponent means that more requests are concentrated on a few hot items. Breslau et. al found that the distribution of web requests from a fixed group of users can be characterized by a Zipf-like distribution with the exponent ranging from 0.64 to 0.83 [27]. They also found that the exponent centers around 0.8 for traces from homogeneous environments. In order to use such a distribution, data items are first assigned *ranks* which are then used in deciding the probability of requesting a data item of a particular rank. The probability of choosing a data item of rank k is given as

$$Pr(Rank = k) = \frac{1/k^\beta}{\sum_{n=1}^{N}(1/n^\beta)}, \tag{14.6}$$

where N is the total number of data items. A total of 400 data items is used here. Ranks are assigned to these data items and the probability distribution is used to determine which item is requested as part of a request. The rank 1 item is the most frequently requested, while the rank 400 item is least requested. Note that we still need to assign a size to each data item.

14.4.2 Relating data item size to popularity

Data item sizes vary from $5KB$ to $1MB$ in increments of $50KB$. The data sizes are set based on the assumption that most data items appearing in relevant application domains will typically contain raw data or encoded using some markup language. A data item of $500KB$ in itself can contain significant amounts of information in this case. We do assume that some large files may exist and hence set the upper limit at $1MB$. These values also reflect the numbers used in related literature. Assignment of sizes to data items of different ranks is done using a hybrid distribution [15]. The body of this distribution
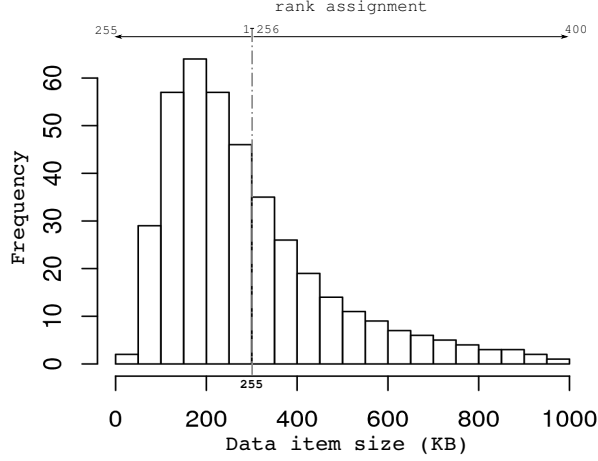
Figure 14.3: Data item size distribution and rank assignment.

follows *lognormal* characteristics while the tail follows a heavy tailed *Pareto* distribution, given as

$$Pr(Size = s) = \begin{cases} \frac{1}{s\sigma\sqrt{2\pi}}e^{-(\ln s - \mu)^2/2\sigma^2} & ,s \leq s_{break} \\ \alpha k^\alpha s^{-(\alpha+1)} & ,s > s_{break} \end{cases}, \tag{14.7}$$

where the parameters are set as follows: $\mu = 5.5$, $\sigma = 0.6$, $\alpha = 1.4$, $k = 175$ and $s_{break} = 500$. The parameters are set so that the lognormal distribution spans over the $0 - 500KB$ range and smoothly transitions into the Pareto distribution from $500KB$ onwards. Fig. 14.3 shows the data item size distribution and the rank assignment scheme on the items. As also observed in web traces used by Barford and Crovella, about 82% of the data items lie in the body of the distribution with the used parameter settings [15]. Note that an actual workload may demonstrate slightly different values for these parameters depending on the number of items, minimum and maximum data size, access frequencies of items, etc. However, the overall distribution is expected to remain consistent with the one used here. The performance results are therefore not restricted to the used parameter settings, but are rather dependent on the effectiveness of the underlying distributions in capturing real workload characteristics.

Each unique data item with size from $5KB$ to $255KB$ are assigned ranks from 1 to 255 in descending order of the sizes. This ranking scheme continues from 256 to 400 with data items of size higher than $255KB$ in ascending order. The rank assignment makes sizes in the range $[5KB, 255KB]$ more frequently requested, while bigger sizes get a below average

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request frequency. The average file size under this distribution is $s_{avg} = 205KB$. This setup follows the general observation made by Breslau et. al that the average size of less frequently accessed data items is comparatively larger than that of frequently accessed items [27].

14.4.3 Assigning deadlines to requests

Expected response times are assigned to requests from a normal distribution with mean $60s$ and standard deviation $20s$. This response time is added to the arrival time of a request to get the absolute deadline for the request.

14.4.4 Varying workloads

One way to simulate different workload conditions is by changing the request arrival rate. Higher request rates result in a larger number of pending requests to schedule at a given time instance. The scheduler's performance is then gauged by how fast the request queue can be cleared by the schedule built by it. This assessment assumes a fixed (and reasonable) bandwidth available for broadcast. An alternative to this, and the one adopted here, is to modulate the bandwidth *utilization factor* of the broadcast server (UF_{bs}) and keep the request rate fixed. By changing the utilization factor we can effectuate different bandwidths available at the broadcast server. This in turn changes the rate at which a particular schedule can be executed, thereby modulating the number of pending requests and simulating different workload conditions. The bandwidth is thus assigned as $b_{bs} = \lambda_{bs}s_{avg}/UF_{bs}\ KB/s$. Given a fixed bandwidth b_{bs}, the request rate is directly proportional to the utilization factor.

14.4.5 Generating requests at data servers

The data servers are simulated as $M/G/1$-PS systems running in parallel with the broadcast server. The bandwidth available at a data server is fixed at $b_{ds} = 5760KB/s$ (a T3 connection). The 400 data items are randomly assigned to the four data servers. Every request coming to a data server has an associated service time (time required to serve the request had it been the only one in the system). For requests originating at the broadcast server, this service time is equal to the size of the data item to be retrieved divided by

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the total bandwidth at the data server. The data servers are also accessed by clients other than the broadcast server in our simulation setup. Service times for requests from such clients are generated as follows. First, a sample of 10 million requests is generated for the broadcast server. For each data server, the sizes of data items in the sample that would be served by this server are then used to generate a size distribution for the server. The service time distribution for the server follows this size distribution, with service times given as the size values divided by the bandwidth. Every time a new request is to be generated at the server, the service time distribution is sampled and the value is used as the service time of the request. The size distribution also lets us calculate an expected data item size that is requested at the server. Requests arrive at a data server at the rate of $\lambda_{ds} = 16$ requests per second according to a Poisson distribution. This, coupled with the expected data item size information, generates a 70% bandwidth utilization in the data servers.

14.4.6 Estimating data server response time distributions

A data server distributes its available bandwidth equally to all pending requests in a round-robin fashion. Hence, depending on the number of requests pending at a server, the time required to retrieve a data item from the server, i.e. the response time of the data server, would typically be higher than the service time of the request. Application of the proposed methods will require an approximation of the probability distribution of data retrieval times. We do not assume any known probability distribution. Rather, we suggest that the approximation be obtained by sampling the data server, and representative points be used to obtain the approximated PMF. In order to generate the response time PMFs of the data servers, each server is run independently and the response times of 10 million requests are noted. This gives us an estimate of the response time distribution at the server, from which 100 points are chosen uniformly to construct the PMF.

14.4.7 Software implementation

The broadcast server requests an item from a data server only when the current broadcast ends and there is an item waiting for broadcast next in the schedule. We pay nec-

essary attention to maintain the dynamics of the data server during the time when the broadcast server is in the process of broadcasting an item. Our implementation achieves this using sequential programming of two components - (i) the *broadcast server module* (BSM) and (ii) the *data server module* (DSM). A system global *Clock* keeps track of the current time (initialized to zero). The BSM reads the data set file and inserts all requests having an arrival time equal to *Clock* into a request queue. The scheduling algorithm is then invoked to generate a schedule to serve the requests in the queue. *Clock* is incremented by one and portions of the schedule that would get executed within this time period (between the old and new value of *Clock*) is updated. The update is performed as follows. The data server hosting the data item next scheduled for broadcast is determined. The DSM for the corresponding data server is called to compute the retrieval time for the data item. It executes by infusing its request queue according to a Poisson distribution, with service times drawn as described in Section 14.4.5. Each DSM maintains its own local clock that runs at 1000 ticks per *Clock* tick. The data item request of BSM is inserted into the queue only when the DSM's local clock is at the time when the previous broadcast (at the broadcast server) ended. This guarantees that the data server is running even when no request was made by the broadcast server. The DSM returns back to the BSM as soon as the requested data item is completely served. In addition, it saves its current state (queue and local clock) if the local clock time did not surpass *Clock*. Similarly, the BSM considers the broadcast of the data item complete only if the retrieval of the item did not overshoot into the next clock tick. A completed broadcast implies removal of all requests in the queue waiting for the data item. The BSM reads the next set of requests in the data file and repeats the process. DSMs corresponding to the four data servers are initialized by running them for at least 50000 clock ticks before starting the BSM.

14.4.8 Convolution time

Convolution is implemented using an $O(n \log n)$ algorithm with *Fast Fourier Transforms* [142]. Iterative convolution, as in MDMP and MBSP, can increase the convolution time as the number of samples in the generated PMF (X_{bdst}) increases. In general, if PMF f has N_f samples and PMF g has N_g samples, then their convolution contains $N_f + N_g - 1$

samples. When applied iteratively, the PMF size can increase considerably, thereby increasing the convolution time as well. Hence, in order to keep the scheduling time within acceptable limits, we direct the algorithm to generate resulting PMFs with a maximum of 100 samples.

14.4.9   Other specifics

The slack deviation parameter $\mathcal{S}_{dev}$ is set at 0.25, unless otherwise stated. The scheduler is invoked every second until the last request in the data set enters the queue; thereafter, it is invoked every five seconds until the request queue becomes empty. The scheduler is invoked every second since the schedule built by it is based on probabilistic estimates of retrieval times of data items (especially for MDMP and MBSP). The actual retrieval time is known only after a data item is fetched completely from the data server, in which case, the scheduler may require to rearrange the remaining part of the schedule for a better QoS. This is also the reason why the scheduler is repeatedly invoked even if no new request enters the queue. Our implementation operates in a batch mode, meaning, the current schedule is revisited (and reorganized if required) after a few data items (more precisely, as much as that can be fetched and broadcasted in one tick) have been fetched and broadcasted.

14.5   Empirical Results

Performance measurement is taken once all 100,000 requests in the data set are served. The total time spent in scheduling instances between a request's arrival and completion is added to the response time of the request while taking the measurements.

14.5.1   Effectiveness of using distributions

In order to demonstrate the effectiveness of using data server response time distributions in scheduling, we compare the performance of MDMP and MBSP to their non-stochastic versions. The non-stochastic versions use the expected value of the response time distributions as an estimate of the time required to fetch a data item from a server. The non-stochastic version of MDMP serves requests in ascending order of their close-
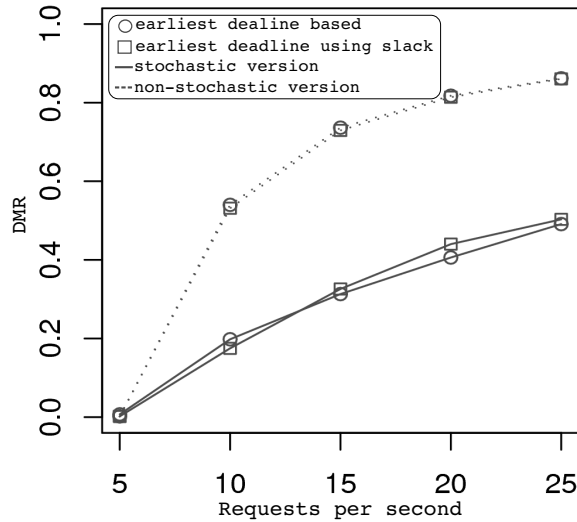
Figure 14.4: Comparative deadline miss rates of identical algorithms in their stochastic (using probability distributions) and non-stochastic (using expected values) versions.

ness to the corresponding deadlines. The expected response time is used to compute when a particular data item broadcast will end, depending on which the earliest deadline request from the pending queue is determined. A similar methodology is used for the non-stochastic version of MBSP, the difference being that the extended deadline (as determined by the slack deviation parameter) is used instead of the actual one. Fig. 14.4 depicts the DMR of the heuristics for varying request arrival rates with a fixed broadcast bandwidth of $1MB/s$. While differences are marginal within the stochastic and non-stochastic versions, significant improvements can be observed when creating schedules based on the probability distributions directly. Deadline misses in the non-stochastic versions are almost double that of the stochastic versions as the workload on the broadcast server increases. This is not a surprising observation since the non-stochastic versions work under the assumption that all data items can be fetched in a fixed amount of time irrespective of the varying workload on the data server. The characteristics of the data item sizes hosted at a data server and the potential impact of the data server workload on their retrieval are more thoroughly maintained in the distributions. The non-stochastic versions do not have a direct method to utilize this information and essentially treats the data servers as a single client secondary storage with constant access time. Note that the expected value is the average response time over a fairly large number of requests made

to a data server. However, requests from a broadcast server to a given data server may be sporadic depending on the popularity of the data items hosted by the server.

14.5.2 Comparative performance

Fig. 14.5 shows the performance metric values for the six heuristics under different workload conditions. Variable workload conditions are simulated by changing the bandwidth of the downlink channel, as given by the different bandwidth utilization factors $UF_{bs} = 0.5, 0.75, 1.0, 2.0, 3.0, 4.0, 5.0$. An utilization factor greater than 1.0 means data requests arrive at a rate higher than that can be handled by the broadcast server bandwidth. With the fixed bandwidth model, these factors translate to workloads generated by more than 5 requests per second with $b_{bs} = 1MB/s$. MDMP and MBSP maintain a DMR of less than 1% as the bandwidth utilization approaches 100%. On the other hand, the DMR of NPRDS increases to 8%. In general, deadline miss rates are less than 10% for utilization factors of less than 1.0. Similarly, all heuristics can generate schedules with more than 95% utility at such utilization factors. All heuristics show an exponential increase in the number of deadlines missed when the utilization factor goes above 1.0. The drop in utility and increase in the average stretch follow similar trends. Both stochastic heuristics, MDMP and MBSP, perform better in DMR and UT over a wider range of bandwidth utilization. MDMP and NPRDS have around 20% deadline misses at $UF_{bs} = 2.0$, compared to 34% of MAX. The utility metric is between 82% (MAX) and 93% (MDMP) at this point. However, as mentioned earlier, the average stretch metric does not reflect this performance. MAX and NPRDS are the better performing ones according to ASTR. We also notice an improvement in the comparative performance of NPRDS as bandwidth utilization goes beyond 300%. The NPRDS heuristic displays better sustenance in all three metrics at such utilizations.

14.5.3 Impact of broadcast utilization factor

Heuristic performance is heavily affected by the workload conditions and bandwidth available at the broadcast server. In conditions of heavy workload, or low bandwidth, the accumulation of requests at the broadcast server dominates the rate at which requests get
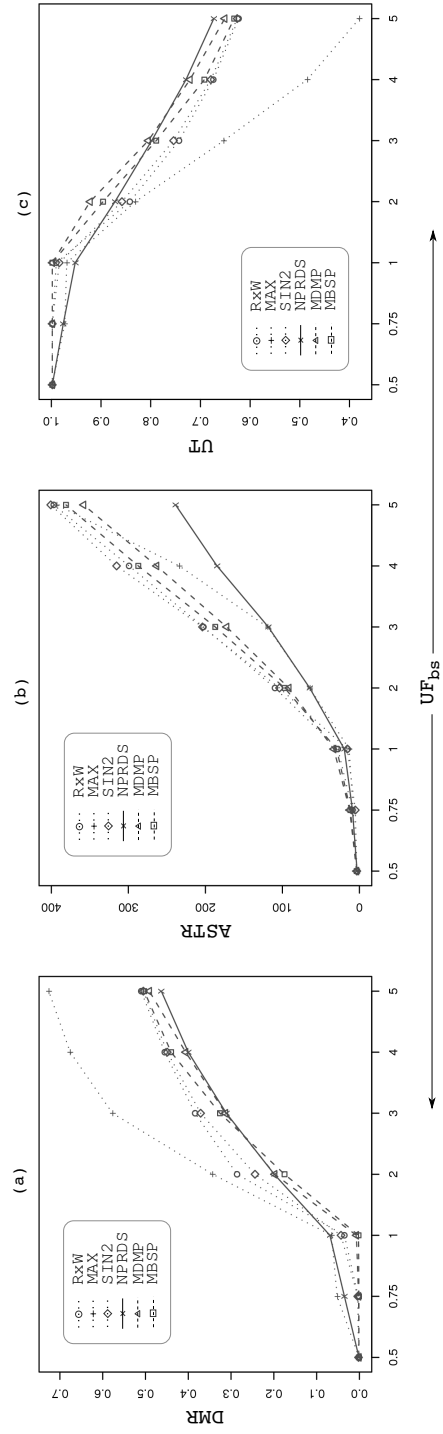
Figure 14.5: Performance metric values for different heuristics under different workloads.

served. If frequently requested items are not regularly broadcast, the request queue gets flooded with such requests. On the other hand, if they are broadcast too often, then the accumulation happens for average and less frequently requested data items. However, we believe the latter factor plays a more prominent role. This is because, although requests for frequent items can get accumulated in the queue, a broadcast policy can let this accumulation happen and reduce the queue size considerably by a single broadcast of the item. This strategy will not work for average and less frequently requested items since the accumulation of such requests will typically take a longer time resulting in a number of them missing their deadline when the broadcast is finally made. In low workloads, most requests are typically for frequently requested items and hence a relatively smaller number of broadcasts can serve most requests in the queue.

Irrespective of the broadcast policy generated by a heuristic, deadline miss rates can still increase with decreasing bandwidth. This is in general attributable to the higher time required to transmit the data items of even an average size. Thus, requests that are supposed to be served by data items towards the end of the schedule suffer a high response time. The situation worsens if an intermediate broadcast is for a larger item.

MDMP and MBSP maintain lower deadline misses for utilization factors less than 3.0. NPRDS and MAX have the worst DMR for $UF_{bs} \leq 1.0$. Recall that NPRDS is the only heuristic that considers the size of the data item in its scheduling decisions. This makes us believe that the size of a data item is not an important factor to consider when scheduling for a high bandwidth system. Clearly, the consideration of size becomes important when bandwidth is low as depicted by the better performance of NPRDS. Unlike the size of data items, stochastic information utilized by MDMP and MBSP seem to become less and less relevant for higher UF_{bs}. The probability estimates do help in sustaining the performance of the two heuristics beyond the $UF_{bs} = 1.0$ mark (note that NPRDS overtakes most other heuristics at this point). To our understanding, for very low broadcast bandwidth, the retrieval time of data items (carried out through a high-speed connection) becomes negligible in comparison to their broadcast times, which in effect diminishes any improvements obtainable by accounting for the retrieval time in scheduling decisions.

An interesting observation to note is at $UF_{bs} = 2.0$. Although, the DMRs of NPRDS, MDMP and MBSP are quite similar at this point, there is still a significant difference in their UT values. NPRDS and MDMP display a DMR of about 20% and MBSP has a DMR of 17.5%. However, MDMP maintains a higher utility than NPRDS (93% compared to 87%). The requests which missed their deadlines in MDMP did so by smaller durations than in NPRDS. Hence, equivalent deadline misses need not always correlate to similar utilities. Broadcast schedules can be generated that, although misses equal number of deadlines, can maintain closer proximity of the response time of requests to their service times. MBSP, on the other hand, displays a trade-off characteristic where lower DMR has been achieved by a slight reduction in the utility (89.6%).

Based on these observations, MDMP and MBSP are probable choices for low and marginally higher workloads, while NPRDS is the likely candidate for very heavy workloads. One should note that deadline misses will increase exponentially when bandwidth utilization goes higher than 100%. Marginal increases in the utilization factor are acceptable owing to bursts in traffic. However, the range of utilization factors where heuristics such as NPRDS performs better in this problem are very unlikely settings for a realistic broadcast system (a system having almost 50% deadline misses). If a situation does occur where the utilization surpasses the expected range, then the network design is revisited or additional resources are installed to bring down the utilization to the sought range. In this regard, MDMP and MBSP both sustain their better performance up to 300% utilization. The observation helps conclude that MDMP and MBSP are also suitable for handling occasional bursts in traffic.

14.5.4 Impact of slack deviation on MBSP

The MBSP heuristic first schedules the request with the maximum probability of missing the extended deadline decided by the slack deviation parameter $\mathcal{S}_{dev}$. The extended deadline for a request Q is given as $dln_Q + \mathcal{S}_{dev}(dln_Q - arr_Q)$. Table 14.1 shows the variation in the performance metrics across five different slack deviations. Recall that $\mathcal{S}_{dev} = 0$ corresponds to the heuristic that prefers the request with maximum probability of missing the deadline and shows very similar performance to the dual heuristic MDMP (minimum

Table 14.1: Performance metric values for MBSP with $UF_{bs} = 2$ and varying $\mathcal{S}_{dev}$.

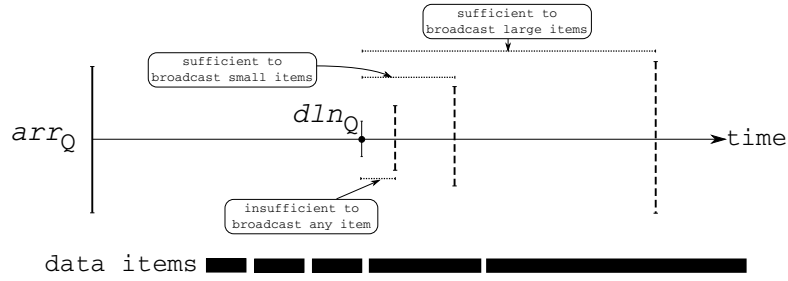| $\mathcal{S}_{dev}$	DMR	ASTR	UT
0.00	0.198	93.0	0.922
0.25	**0.175**	97.0	0.896
0.50	0.236	111.8	0.866
0.75	0.263	124.9	0.84
1.00	0.266	132.5	0.833

Figure 14.6: Impact of $\mathcal{S}_{dev}$ on scheduling decisions.

probability of meeting the deadline). While ASTR and UT show consistent degradation as $\mathcal{S}_{dev}$ is increased from 0.0 to 1.0, DMR displays a change in gradient around $\mathcal{S}_{dev} = 0.25$.

The intuition behind MBSP is to provide "sufficient" slack to the scheduler to satisfy other requests at the expense of ones which are likely to or have already missed their deadlines. This is achieved by informing a longer deadline for requests to the scheduler, instead of the actual one. The slack deviation parameter helps modulate the amount of slack so that exploited requests do not remain in the system for ever. The key to setting the parameter is understanding what serves as "sufficient".

Refer to Fig. 14.6. When the slack deviation is too small, the slack is of little or no help since no data item (to serve other requests) can be broadcast between the actual and the extended deadline. The performance of MBSP would thus be equivalent to that of MDMP. The reason why a value such as 0.25 works best for the tested workloads can be attributed to the size of the most frequently requested items (an item of around $200KB$ in the experimental setup). Assume that a request Q has already missed its actual deadline $t = dln_Q$. Further, let its extended deadline be $t + x$, where x is determined by the slack deviation parameter as $x = \mathcal{S}_{dev}(dln_Q - arr_Q)$. Let the current time be $t + 1$. Since MBSP sees the deadline for Q as $t + x$, it would try to schedule other requests first

(ones which are closer to their extended deadlines) provided the probability of Q missing its extended deadline is not the maximum. If x is small enough (correspondingly a value such as 0.25) that only moderately sized data items can be fit in the schedule before Q's probability of missing the extended deadline becomes the maximum, then there will be a higher likelihood that a more frequently requested item is scheduled next. This can be viewed as a positive effect of using the slack since many requests will get served by this broadcast. However, for larger x, broadcast of a large data item can also be accommodated within the slack period. This is a negative effect of using slack since broadcasting such an item would serve only one or two requests, in addition to delaying the other pending requests further. The sufficiency of the slack is thus related to the size of data items that are more frequently requested. Intuitively, the slack deviation should be set so that the introduced slack is not misutilized in broadcasting data items that would serve very few requests. Of course, this explanation is valid only in the context of our experimental setup. In our setup, the infrequent data items were the ones larger than $500KB$. Hence, higher slack values demonstrated poorer performance. On the other hand, if frequently requested items are indeed the larger data items, then introducing a higher slack would be beneficial.

14.5.5 Effect of scheduling by size

In this section, we shall try to understand the effect of considering the size of data items in scheduling decisions for different workload conditions. NPRDS is the only heuristic we explored that explicitly uses data size in its formulation. Fig. 14.7 illustrates the frequency distribution of broadcasts by the rank of data items. At $UF_{bs} = 0.75$, RxW and SIN2 has a DMR value of less than 0.005, while that of NPRDS is 0.035.

The broadcast policy adopted by RxW and SIN2 results in a frequency distribution similar to the Zipf distribution of data item requests. NPRDS shows peaks in the range of data items that are requested on a more than average basis. As seen in the plot, even closely ranked data items can have very different broadcast frequencies.

Both RxW and SIN2 consider only the urgency and productivity factors, while NPRDS also considers the size in addition. The scheduling policy used by NPRDS is based on a
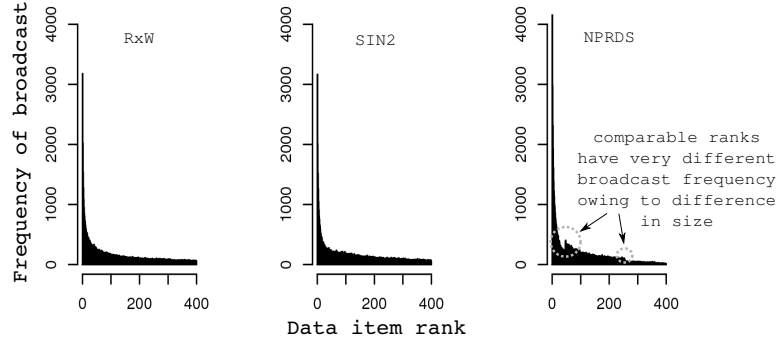
Figure 14.7: Broadcast frequencies of data items by rank at $UF_{bs} = 0.75$.

priority value that is higher for smaller sized data items (assuming similar productivity and urgency). Recall the rank assignment scheme for data items. Most average requested data items under the scheme have lower (but comparative) data size than the frequently requested items (in the range of $155KB$ to $205KB$). Contentions are not unlikely between data items of similar sizes. Consider a case where a request for a data item of size $255KB$ and a request for a data item of size $155KB$ are equally urgent. Let there be 30 requests for the $255KB$ item and 20 for the $155KB$ item. RxW and SIN2 in this case would give preference to the $255KB$ item, whereas NPRDS would choose the $155KB$ item ($\frac{20}{155} > \frac{30}{255}$). Note that the difference in broadcast time of the two items will be very small in a high bandwidth system. Hence, broadcasting the data item that can serve more requests (the $255KB$ item) would be more sensible so that less number of requests (the ones for the $155KB$ item) have to wait further. However, the policy of NPRDS can be useful in low bandwidth situations since broadcast of the smaller item achieves the objectives of inducing lower latencies for other requests and serving a significant number of requests. The better performance of NPRDS in heavy load conditions conforms to this justification. Thus, size considerations are more important when fast servings are required for a large number of requests in a relatively small amount of time.

14.5.6 Stretch and performance

Recall that the MAX heuristic has the worst performance according to DMR and UT across all variations of the utilization factor. This performance indication contradicts the picture presented by the ASTR metric. MAX maintains lower average stretch across a
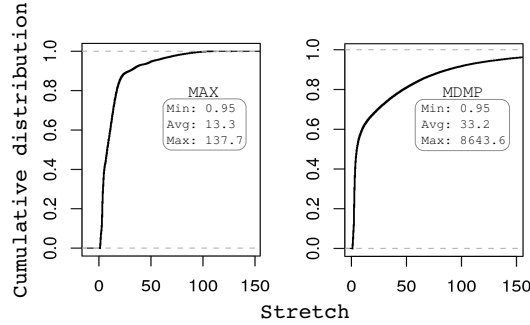
374

Figure 14.8: Cumulative probability distribution of stretch values at $UF_{bs} = 1.0$.

wider range of utilization factors. We also observe that the maximum stretch in a MAX schedule is comparatively much lower than that from the other heuristics. Nevertheless, MAX misses the highest number of deadlines. Ideally, lower stretch values mean that response time of requests are closer to their service times, and hence they should be able to meet their deadlines. To understand this discrepancy, we refer to the distribution of the individual stretch values under MAX and MDMP generated schedules.

Fig. 14.8 shows the cumulative probability distribution of stretch values for MAX and MDMP at $UF_{bs} = 1.0$. At this utilization, MAX has a DMR of 6.5% while that of MDMP is 0.7%. However, the average and maximum values of stretch is much lower in MAX. In fact, MAX demonstrates clear efficiency in keeping the maximum stretch of the system at a minimum at all levels of utilization. The distribution reveals that the stretch values of about 60% of the requests are very similar across the two heuristics. Sharp differences appear in the remaining 40%. These differences shifted the average stretch of the system to a higher value in MDMP. However, it still does not explain why MAX has a higher DMR. Since the stretch values are spread over a much wider range for the aforementioned 40% of the requests in MDMP, the likelihood of missing deadlines is more than MAX for such requests. Thus, whatever performance difference exists between MAX and MDMP must come from requests with small stretch values. We refer to Fig. 14.9 with an intention to observe any such difference.

The fundamental difference becomes clear from Fig. 14.9. The plots show the distribution of slack values (time difference between the completion time of a request and its deadline) for requests that met their deadlines, i.e. $dln_Q > ct_Q$. It seems the poor perfor-
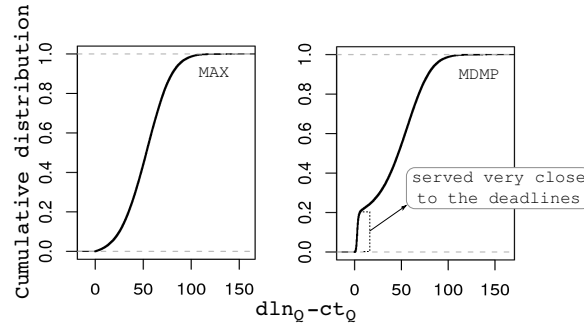
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Figure 14.9: Cumulative probability distribution of $dln_Q - ct_Q$ (positive slack) for requests which are served within their deadlines ($dln_Q > ct_Q$); $UF_{bs} = 1.0$.

mance of MAX is attributable to its inability to utilize the time gaps between a request's arrival and its imposed deadline to serve other pending requests. 20% of the requests in MDMP are served very close to their deadlines (near zero positive slack). This indicates that MDMP manages to utilize the slack between a request's arrival and its deadline in serving more urgent requests. Requests far away from their deadlines will have higher probabilities of meeting their deadlines and hence can be pushed back in the schedule to accommodate requests with lower probabilities of meeting their deadlines. The hypothetical deadline created by MAX cannot capture this logic since it attempts to service a request as close as possible to its service time. An important conclusion from this analysis is that the stretch of a request, at least in its defined form, is not an accurate factor to include in deciding scheduling policies. This is particularly true for deadline based systems where using any introduced slack may be advantageous.

14.5.7 Broadcast policies

The final analysis presented in this section is on the broadcast policies generated by heuristics that result in additional performance improvement. For this we observe the policies generated by MBSP and NPRDS. The performance difference between the two heuristics diminish as the utilization factor increases, with NPRDS gaining at very high utilization. The policies are observed in terms of the request and broadcast frequency of data items. To do so, we divided the data items into four access types: *frequent* (ranks 1 to 20), *above average* (ranks 21 to 45), *average* (ranks 46 to 255) and *infrequent* (ranks 256 to 400). We then observed how the broadcast frequency of data items in the four categories
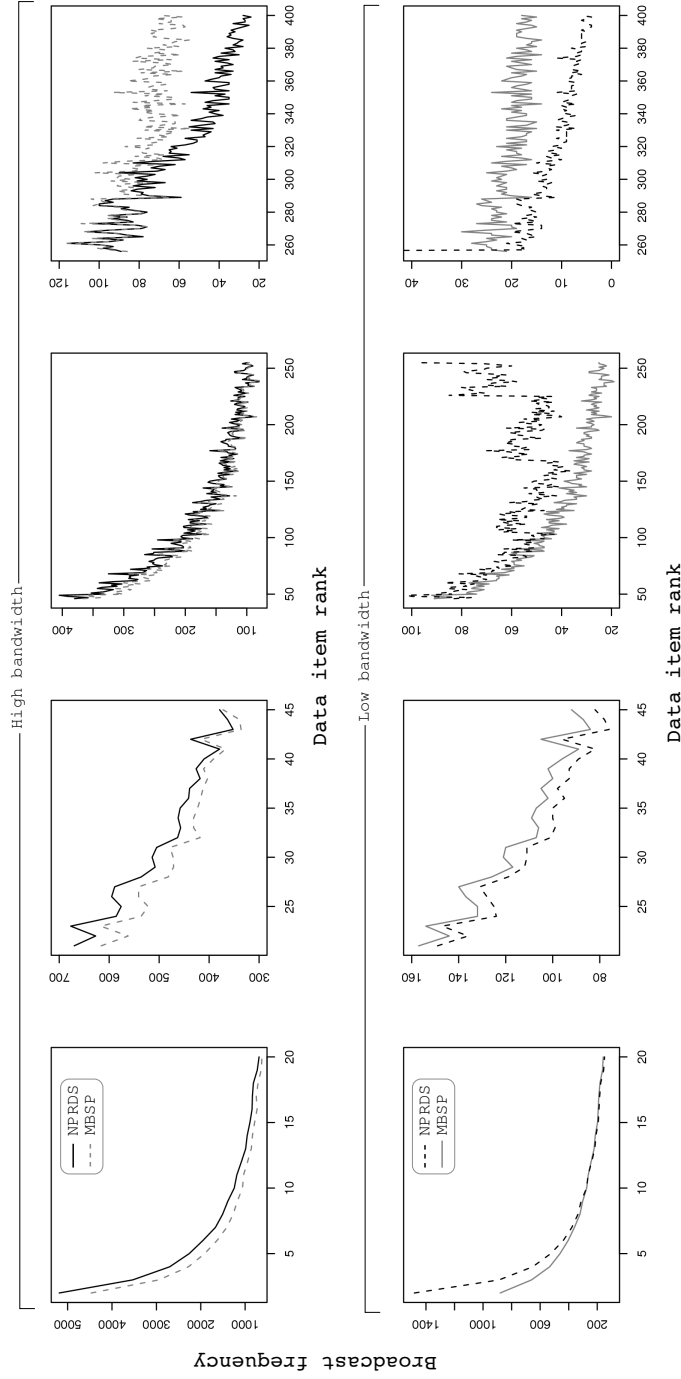
Figure 14.10: Broadcast frequency of different data items by their ranks using MBSP and NPRDS. Top: Low utilization ($UF_{bs} = 0.75$). Bottom: High utilization ($UF_{bs} = 4.0$). The dotted lines signify the better performing heuristic.

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Table 14.2: Relative broadcast frequencies of data items according to access types. Each entry signifies if the frequency in the better performing heuristic is lower, similar, or higher than that in the other heuristic.

Access type	$UF_{bs} = 0.75$	$UF_{bs} = 4.0$
frequent	Lower	Higher
above average	Lower	Lower
average	Similar	Higher
infrequent	Higher	Lower

differ across the two heuristics. Factors such as request accumulation, broadcast time and request rank appear as major aspects underlying the differences in the broadcast policies.

Fig. 14.10 illustrates the broadcast frequencies of the data items according to the four access categories. The better performing heuristic is marked with a dotted line in the plots. Table 14.2 summarizes the broadcast frequencies of the better performing heuristic relative to the other.

In a high bandwidth (or low utilization) situation, it may be reasonable to allow more accumulation of frequent data items. This is not only because more requests can be served by a single broadcast, but also because it helps assign more broadcast time to infrequent items. Assuming that access frequencies for data items are the same across different workload conditions, there will not be many requests for infrequent items in a low workload. Hence, a scheduler should not wait for request accumulation of infrequent items and should broadcast the items on a regular basis. It can happen that such a broadcast serves a single request only. Regular broadcasts of infrequent items (larger sizes in our data set) do not induce much delay for the other requests owing to the high bandwidth available. The broadcast frequencies of MBSP is in conformation to this intuition.

The broadcast policy is quite different when the bandwidth available is low, or there is heavy utilization. The plots clearly show differences in the broadcast frequencies of average and infrequent items. Observe that NPRDS has higher broadcast frequencies for frequent and average data items. Higher broadcast patterns for frequent items is crucial in low bandwidths since the broadcast times of items will be typically high. Allowing too many such requests to accumulate and then broadcasting over a slow channel can

introduce deadline misses for a higher number of requests. On the other hand, accumulation of average items should not be allowed since it would usually take a long time for a significant accumulation to happen. This will only keep such requests pending in the system for a long duration of time. The request accumulation of above average items are somewhere in between frequent and average ones. NPRDS decides to allow the accumulation of these requests more than MBSP to utilize the broadcast bandwidth for frequent and average items. Infrequent items are ignored the most owing to their high broadcast time.

Understanding the broadcast policies in this manner not only reveals what factors are more important in different utilization, but also paves the way to transform a pull-based system to a push-based one with similar performance. However, in addition to relative measurements, such conversions will also require the estimation of absolute frequency in a probabilistic manner.

14.6   Conclusions

In this chapter, we introduce the problem of stochastic scheduling in broadcast systems where data items must be retrieved from different data servers prior to broadcasts. We model the data servers as $M/G/1 - PS$ systems where the available bandwidth is equally divided among pending requests. Schedules are then generated at the broadcast server to serve requests within their imposed deadlines. Two novel heuristics are proposed that use the probability distribution of response times in the data servers to generate schedules. The Minimum Deadline Meet Probability (MDMP) heuristic schedules requests based on their probability of meeting the deadline. The Maximum Bounded Slack Probability (MBSP) heuristic defines an extended deadline for requests as given by a slack deviation parameter, with an intention to use any negative slack for serving other pending requests. The performance of these heuristics is compared to that of four other heuristics – RxW, MAX, SIN2 and NPRDS – that have been proposed for deterministic broadcast scheduling. Performance is evaluated using three metrics - deadline miss rate, average stretch, and utility.

Empirical results on a synthetic data set reveals that both MDMP and MBSP show

superior performance in conditions of both low and high broadcast utilization. In very heavy workloads, the response time of data retrieval is observed to become negligible compared to the broadcast time. NPRDS provides a better performance under such situations. However, such load conditions are often not realistic. Further analysis on the impact of various factors on broadcasting policies reveals that size of data items can often be ignored in scheduling decisions when the broadcast bandwidth is high. The stretch of requests, as typically defined, is also found to be an inaccurate factor to consider while scheduling in deadline based systems. Finally, good broadcast policies can have very different characteristics depending on workload conditions. Request accumulation, data item access frequencies and their broadcast times assume different levels of prominence in different situations, thereby affecting the performance of the heuristics. We also explore the effect of the slack deviation parameter on MBSP and found correspondence of its performance to the amount of negative slack that the heuristic is allowed to exploit. Typically, very low values do not enable any slack utilization, while high values provide too much of it to make the broadcast of larger data items feasible.

In future, we need to explore the scenario when multiple servers can host the same data item. The scheduler then has to build a fetching schedule based on perceived response times and evaluate its effectiveness in combination with the broadcast schedule. Interactions between the broadcast and data server can be made more complex by including concepts such as parallel/pre-fetching and data replication. In general, parallel/pre-fetching will change the probabilities of retrieval times of the data items. The final scheduling approach can therefore remain the same. Data replication will have a deeper impact since the probabilities of fetching the same item within a specific period will be different for different data servers. Increasing the number of data servers and distributing frequently accessed data items across them can also provide lower data retrieval times and can correspondingly impact heuristic performance. These issues require a much extensive analysis and thus forms the basis for our future work.

CHAPTER 15

Dissertation Summary

$\mathbf{M}$odern information management systems are required to satisfy multiple goals in their implementation. Simultaneous adherence to these goals is often found to be difficult owing to their conflicting nature. This dissertation explores three goals laid down for any such system, namely risk minimization, maximal resource utilization and service quality maximization.

Multi-criteria analysis is proposed as a flexible tool to obtain the best balance along the three dimensions – risks, resources and services. Balancing these dimensions is challenging due to the presence of inherent reciprocal influences. We explore three domains in computer science that exhibit the requirement for a multi-criteria analysis. Extensive empirical studies are provided to establish our propositions.

We use disclosure control as the first platform to demonstrate how controlling the exposure of sensitive personal information can negatively impact the service goals of a data broker. Our extensive survey of existing techniques indicate that the current trend is to determine data modifications that preserve a pre-specified level of privacy and induce a minimum loss in information within the constraint. Contrary to this trend, we propose evaluating privacy and data utility within a multi-objective optimization framework. This provides a data publisher the trade-off information required to make an informed decision on what level of privacy can be enforced for a certain level of data utility, and vice

versa. Our proposed techniques also allow a data publisher to limit the exploration of solutions to a subset that closely matches the preferences of the publisher. One of the important outcomes of the multi-criteria analysis is the identification of bias in individual privacy levels. This bias is introduced primarily due to the minimalistic nature of existing privacy models. Towards this end, we redefine the characteristics of an optimal solution and propose a multi-criteria framework capable of identifying solutions with user-defined properties.

The second domain we explore, security risk management, highlights how limited resource availability often becomes a deciding factor while protecting the assets in a network. We move beyond the assumption that an organization has all necessary resources to protect its network, and argue that sections of a network will always remain exposed to an attacker. Towards this end, we propose using cause-consequence attack models to obtain a global picture of residual risks and involved hardening costs. The methodology can also be used to determine optimal points of hardening under other resource constraints such as energy consumption and computational power of a node. We also show that knowledge about the attack difficulties can help identify where resources need to be spent predominantly. This can be done by mitigating vulnerabilities with a high probability of being exploited. Game theory based extensions of the security hardening problem reveal that attacker-defender dynamics cannot be ignored since an optimal solution in the long run is very different from one that provides short term optimality.

Wireless data broadcasting is used as the third domain to exemplify the impact of limited resources on the quality of service that can be delivered by an information system. The QoS criteria to satisfy while maximally utilizing the wireless bandwidth is predominantly based on hard deadlines. We provide justifications to use soft deadlines as an alternative since it allows a broadcast scheduler to relatively determine the utility of a broadcast data item. We explore broadcast systems supporting different data dependencies (single data, transaction data and ordered data) and data retrieval schemes.

Specific future directions have been highlighted at the end of individual chapters. We outline below the general directions for each of the explored domains.

- There is an abundance of privacy models proposed for data anonymization. How-

ever, privacy issues extend much beyond databases. The rapid growth of mobile communication systems, and the application domains thereof, have started to reveal disclosure problems usually unseen in databases. However, a majority of existing models can be transitioned to address these problems. The question that remains open is how these models will affect the usability of a mobile application. Addressing the privacy versus usability problem is an immediate requirement for these mobile information systems.

- Optimal security hardening has been researched with a static perspective for a long time now. Interesting approaches have originated out of this research. Nonetheless, the optimality of a security policy is very easily invalidated by the changing dynamics of a network and growth of attacker capabilities. Given that policy decisions cannot change as easily as the network itself, there is the need to revisit the definition of an optimal security solution. Research is required to tightly integrate the possible evolution of the network and the attacker into this definition. Resource utilization cannot be maximized in this domain without an evolving defender model.

- A number of generic heuristics exist to build broadcast schedules for efficient data broadcasting. However, application requirements are often very diverse in mobile environments. Although utility functions can capture the global characteristics of data usefulness, the quality of service would be sub-optimal unless data utility can be expressed as a function of the underlying application usage. Custom representations of data utility will further influence a number of decisions on where data items should be stored, what replication frequencies should be used, and what update policies will be best for transient data, among others.

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