

The implementation of modern genetic improvement methods in wildlife

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The implementation of modern breeding methodologies in wildlife



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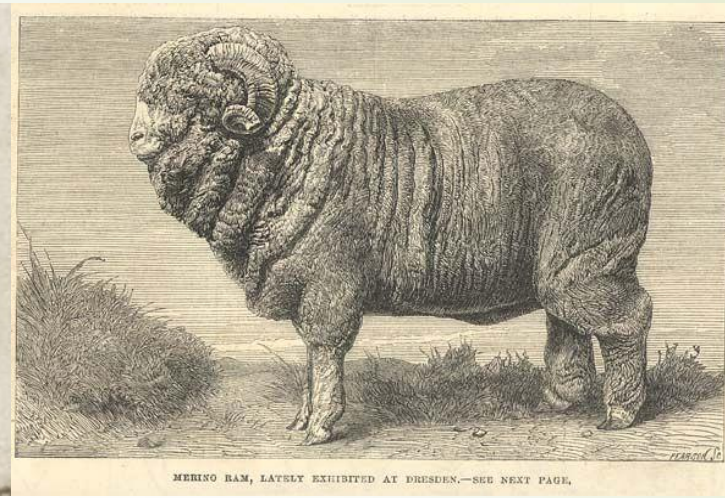
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**Selection is the most
powerful tool in any
breeders armour**

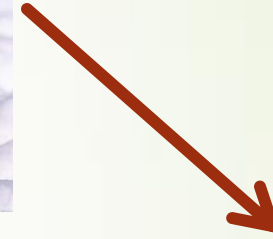


A breed is not a static entity but a process

(Dobzhansky 1959)



Genetic Variation - Dogs



vs

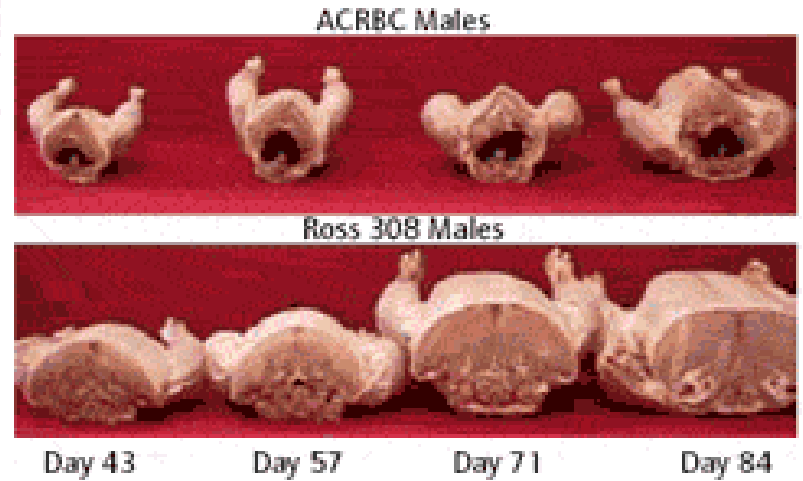


Genetic Variation - Horses



Thumbeline (42cm) vs Clydesdale (2.7m)

Figure 2 - Broiler carcasses from Ross 308 in 2001 compared to unselected controls representing performance in 1957 (Source: G.B. Havenstein 2006, Lohmann Information)

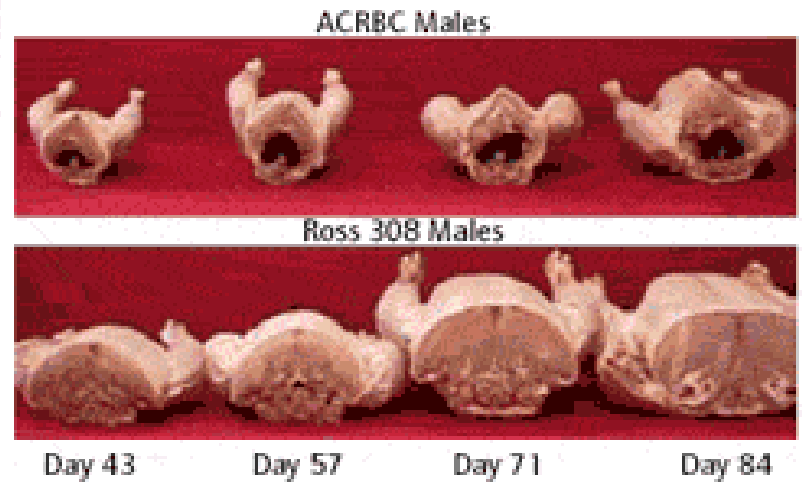


← Unselected

← Selected

- Average body weight of commercial broiler at 56 days of age:
 - 809 g in 1957
 - 3 946 g in 2001

Figure 2 - Broiler carcasses from Ross 308 in 2001 compared to unselected controls representing performance in 1957 (Source: G.B. Havenstein 2006, Lohmann Information)



← Unselected

← Selected

- ➔ Utilizing feeds typical for both years, it can be shown that 85-90% of this 4.87-fold improvement was due to genetics while the remaining 10-15% was due to nutrition.



What is needed for effective selection?

Accurate:

- definition of breeding goals
- identification of superior animals that will be parents in the next generation



Model

$$P = G + E$$



Model

$$P = G + E$$

Phenotype = Genotype + Environment



Model

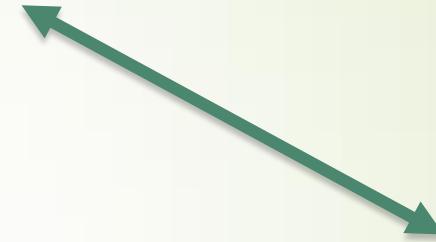
$$P = G + E$$



Phenotype



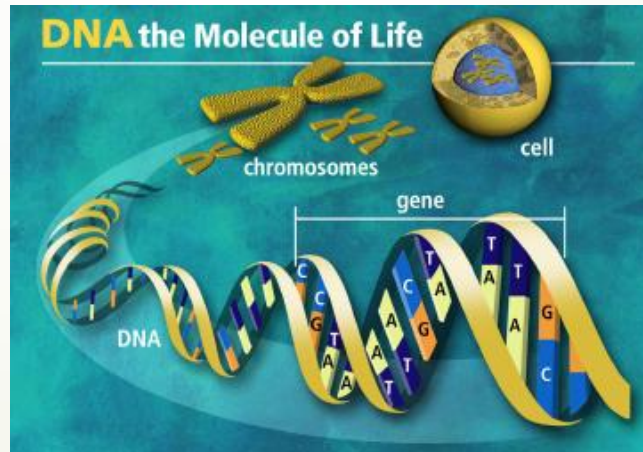
Genotype



Environment



=

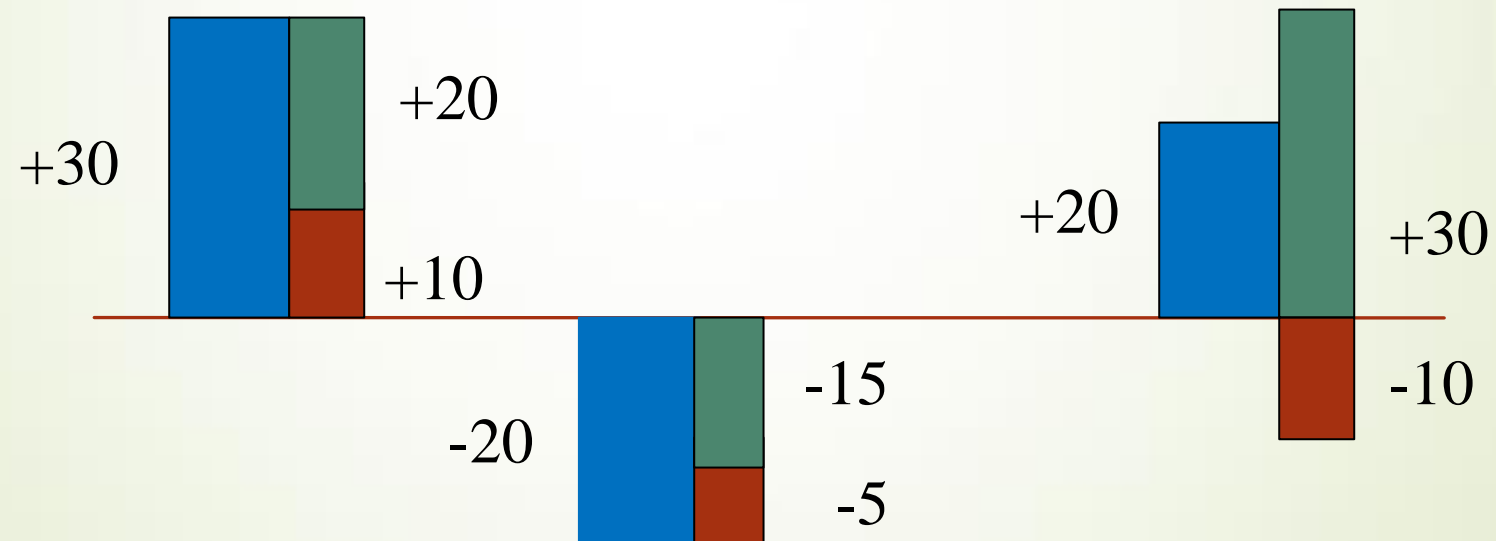


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Genetic and Environmental contributions to weaning weights

$$P = G + E$$






What is BLUP?



- *Best Linear Unbiased Prediction* of **breeding values**
 - Statistical model
 - Uses all information from all animals
- 



What is a breeding value?

- Genotypic values transmitted from parents to offspring
 - Sum of the independent gene effects
 - Value of an animal as a parent
- 

Breeding value (BV): Example

Assumption: Contribution of gene B  +10g
gene b  -10g

Then **BV** for:

$$BB = 10 + 10 = 20g$$

$$Bb = 10 + (-10) = 0g$$

$$bb = -10 + (-10) = -20g$$

➤ **BV** of offspring = $1/2$ **BV** of Sire + $1/2$ **BV** of Dam



How does it work?

- Solving mixed model equations
- *Mixed* – models contain both environmental and genetic effects
- *Animal model* – Combination of own and all relatives' performances




Attributes of BLUP

- Accounts for genetic level of contemporary groups or “competition”
- Accounts for genetic trend
- Uses all data on all relatives
- Can be multiple-trait
 - Utilises correlations between traits
- Accounts for non-random mating
- Accounts for culling for poor performance



Applications of BLUP

- Genetic relationships between herds – comparable
 - Whole breed analysis
 - True genetic progress in breed or herd
 - Environmental trends – Management decisions
 - Inbreeding monitoring
- 



What is needed for BLUP

- Full pedigree records
- Complete performance records
- All non-genetic factors that play a role in the animals' performance:
 - Herd
 - Birth year
 - Sex
 - Age of dam
- **Whole herd recording**



Pitfalls in Large-Scale Genetic Evaluation

Faulty data:

- Pedigrees
- Performance records
- Data used in adjustment / Inaccurate adjustment of data
- Contemporary groups
- Lack of connectedness
- Genotype by environment interaction
- Selective recording




Danger of
single trait
selection
(horns and head
shape)

Trait categories

- Functional / fitness
- Production / growth
- Input (feed efficiency)
- Product / quality
- Behavioral
- Type (current emphasis in game improvement)





Number of Buffalo and Sable animals on data base

| | Buffalo | Sable |
|---------|----------------|--------------|
| Years | 22 | 23 |
| Males | 620 | 1264 |
| Females | 1138 | 1898 |
| Total | 1758 | 3162 |

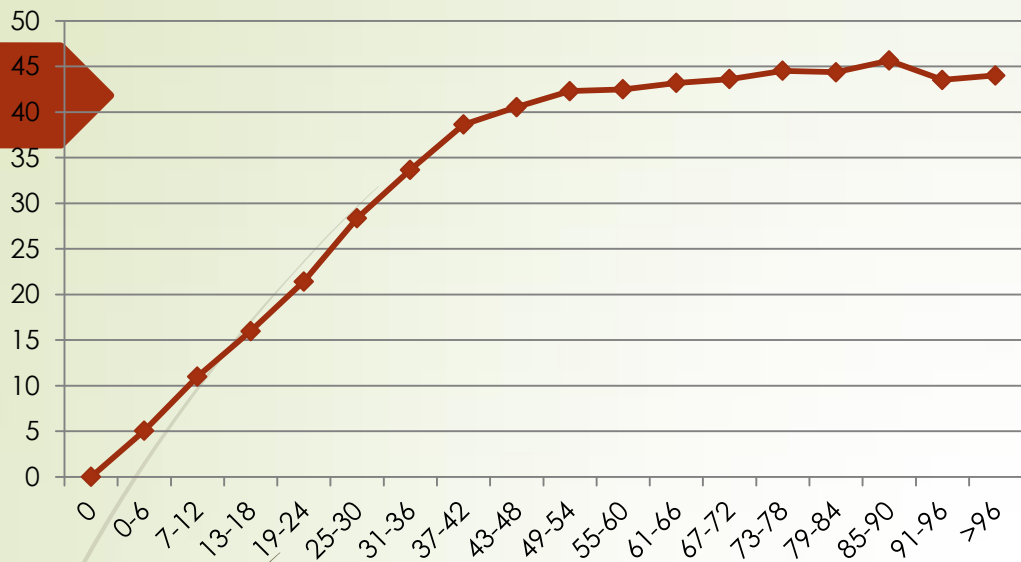
Current buffalo data on system

| | n | Mean | Min | Max | STD |
|-------------------------------|----------|-------------|------------|------------|------------|
| SCI | 570 | 66.11 | 16.75 | 136.50 | 17.26 |
| Tip to Tip | 292 | 29.72 | 4.00 | 97.50 | 20.36 |
| Width of Boss Right | 665 | 9.96 | 4.25 | 39.38 | 3.57 |
| Width of Boss Left | 646 | 9.96 | 4.25 | 38.00 | 3.61 |
| Greatest width outside | 1406 | 27.92 | 2.38 | 625.00 | 21.20 |
| Scrotal circumference | 28 | 21.96 | 1.50 | 49.00 | 15.25 |
| Horn Length Right | 219 | 26.87 | 0.13 | 48.00 | 8.99 |
| Horn Length Left | 191 | 27.16 | 6.50 | 49.50 | 6.50 |

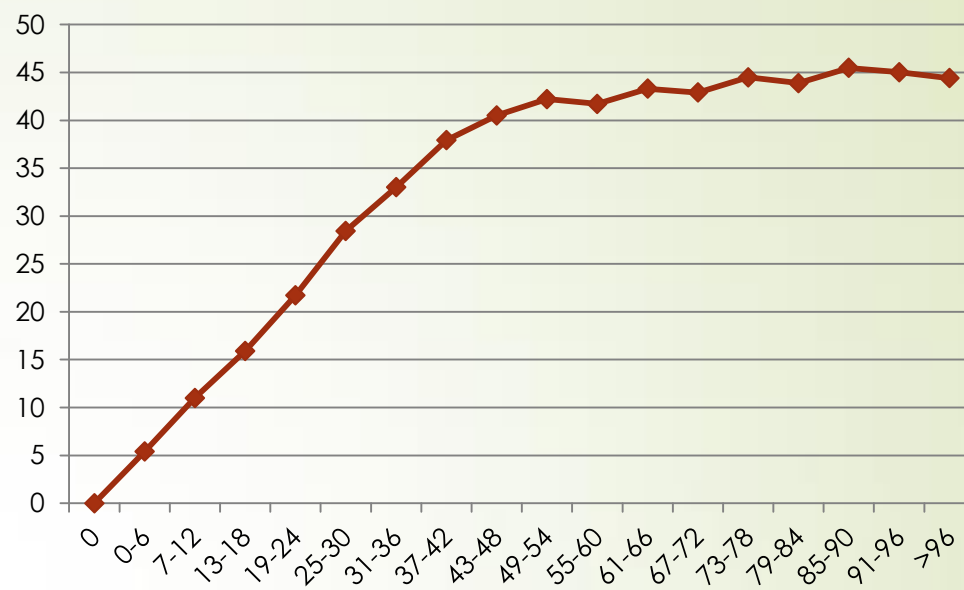
Current Sable data on system

| Trait | n | Avg | Min | Max | SD |
|---------------------------|------|-------|------|------|-------|
| Circumference at base | 1851 | 7.49 | 1.88 | 25 | 1.86 |
| Circumference (right) | 1842 | 7.5 | 1.88 | 40 | 2.05 |
| Horn Length (Left) | 2188 | 25.03 | 1.06 | 94 | 11.73 |
| Horn Length (Right) | 2194 | 24.97 | 1.06 | 94.5 | 11.7 |
| Length of horn from first | 3717 | 9.68 | 1 | 36 | 2.78 |
| Number of Rings (Left) | 1750 | 21.09 | 1 | 57 | 12.79 |
| Number of Rings (Right) | 1728 | 21.18 | 1 | 55 | 12.79 |
| Scrotal circumference | 77 | 20.55 | 5 | 41 | 5.92 |
| Tip to Tip | 1905 | 10.38 | 1 | 49 | 5.01 |

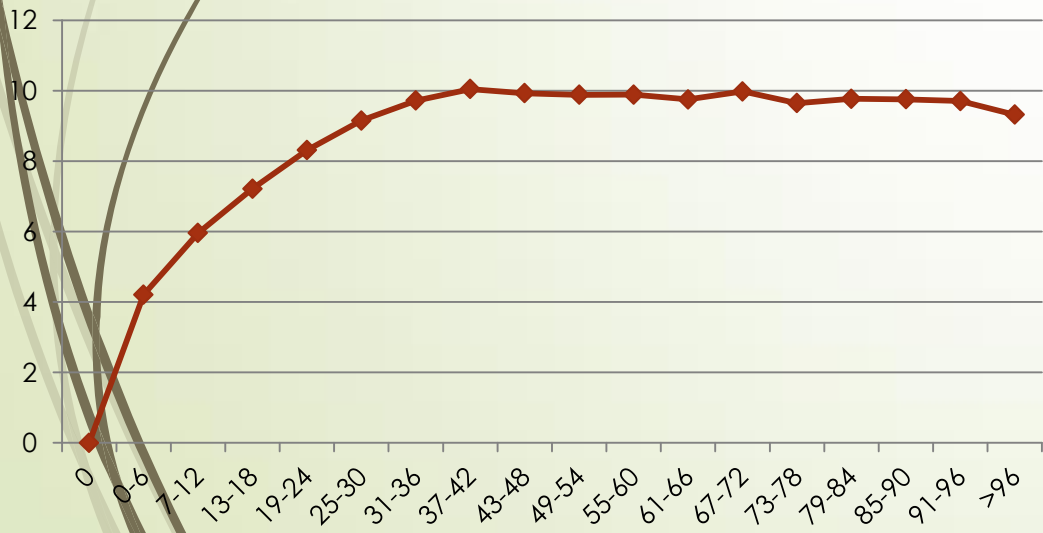
Horn Length Left (inches)



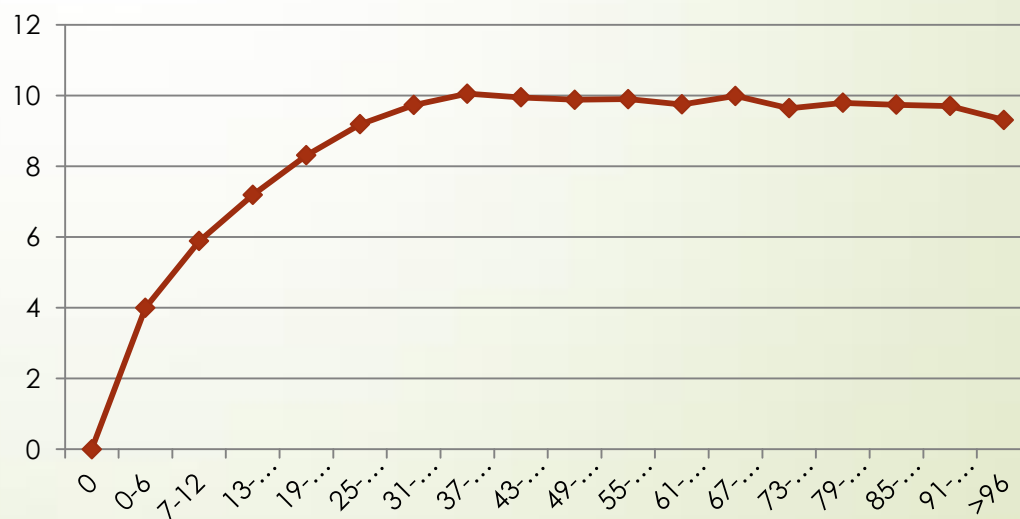
Horn Length Right (inches)



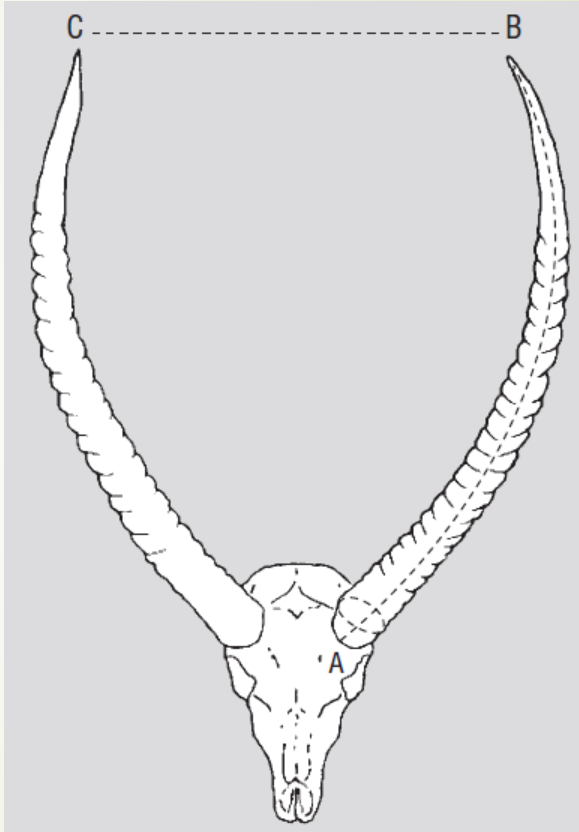
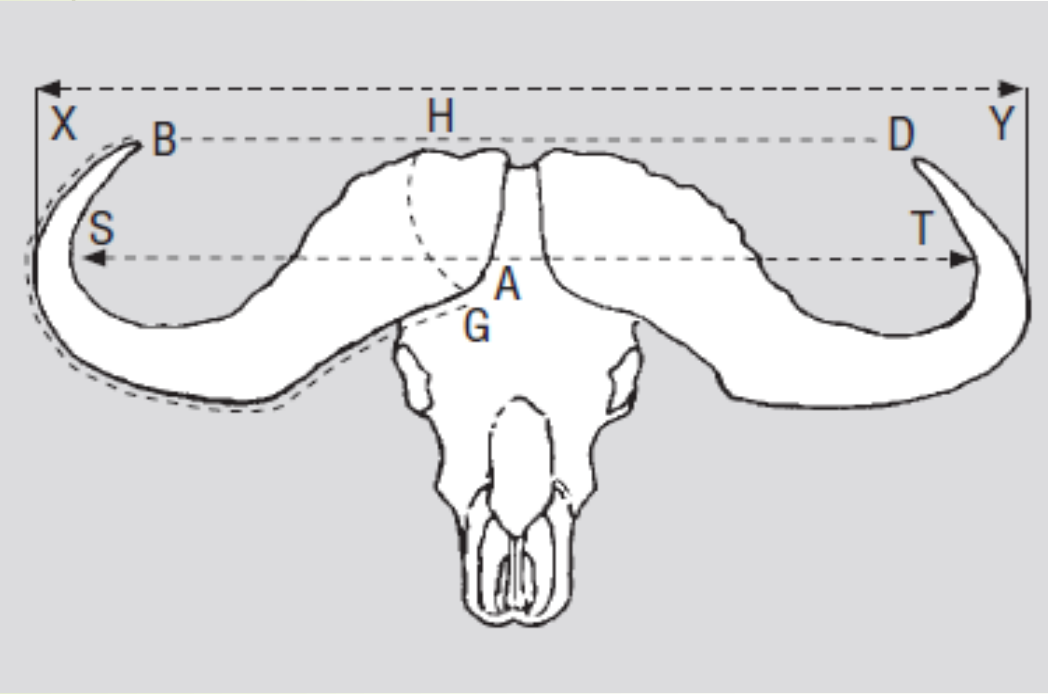
Horn Circumference Left (inches)



Horn Circumference Right (inches)



Standardize Recording



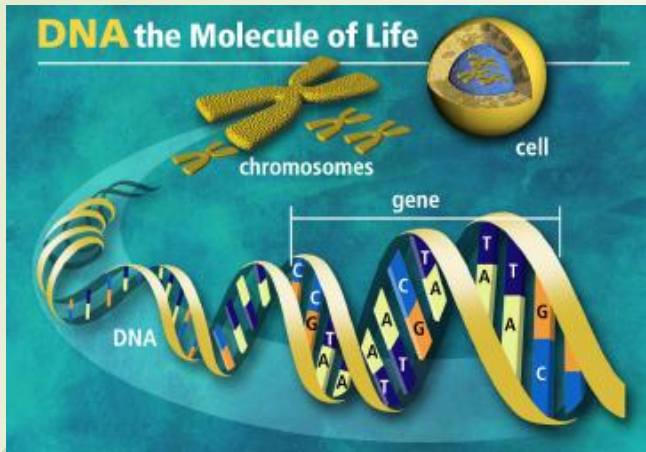
Traits

- Standardize traits
 - Age of animal, and method of measurements
- Many traits determine the value of an animal
- Too many traits in selection program – slow progress
- Choice of traits NB
- Use an economic index:
 - $$I_T = -1.60BW_D - 1.95BW_M + 2.23WW_D + 1.75WW_M - 0.54FW - 2.01MCW - 13.21DC + 4.97BF - 2.36T + 12.66M$$



Take home message

- MEASURE - **Phenotypes** (Traits) with **Pedigrees** of all animals
- DATA - **BLUP for BV's**
- GOALS – Set up a balanced **selection index**
- SELECT – Superior animals to be **parents of next generation** based on BV's
- BREED - Use these animals in a **mating system** to **achieve the goals**



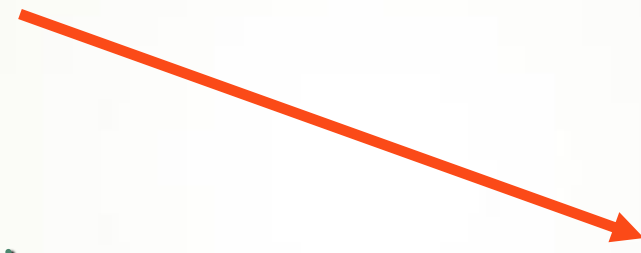
Model

$$P = G + E$$

Genotype

Environment

Phenotype



Thank You

