



# Genomics, EPDs and their application to beef herds in Florida

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# Outline



- Revisiting some basic genetic concepts
- Beef cattle – traditional selection
- Genomic selection - practical questions
  - Dairy Industry as a genomic selection success story
  - Beef Industry as an “opportunity for improvement”
- What genomic tests are available?
- What do the results mean?
- Current and future status of this technology

# Basic Selection Principles

# Selection and Prediction



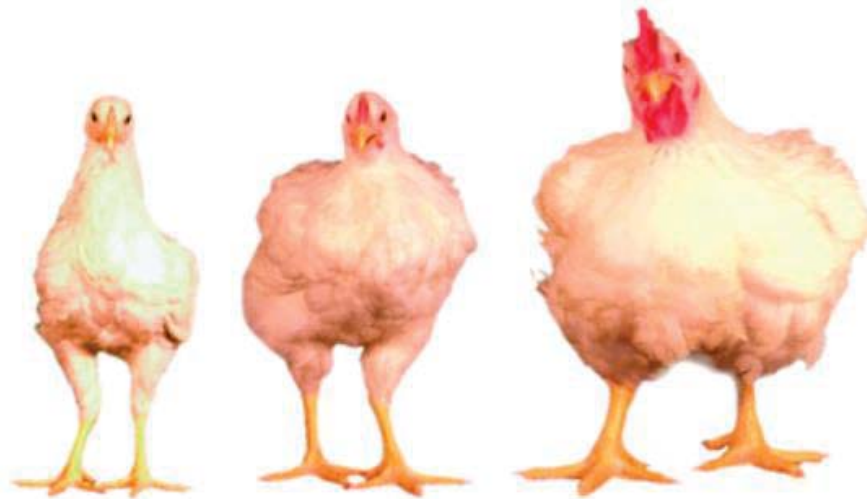
- Genetic change - use animals better than the average as parents of the next generation
- Key to genetic change: selection
  - Estimate **genetic merit** for animals in a population
  - Select **superior animals** as parents of future generations

EBV, EPD

# Traditional Animal Breeding



- Selective breeding for economically important traits
- Traditionally based on phenotypic recording
  - Estimation of breeding values from phenotypic records and pedigrees
  - Knowledge of heritability of each trait
- Successful
  - but slow process especially for certain traits



Fed identical diets, kept in similar conditions for 56 days.

Strains:	<b>1957</b>	<b>1978</b>	<b>2005</b>
56-d weight:	<b>905g</b>	<b>1,808g</b>	<b>4,202g</b>

# Impact of innovation in US



1977

**7 cows today  
yield the beef of  
10 cows in 1977**



Today



Mid 1940's

**Cows yield  
4 times the milk  
of cows in the  
mid 1940s**



Today

1. J.L. Capper. "The environmental impact of beef production in the United States: 1977 compared with 2007." Journal of Animal Science, 2011

2. J.L. Capper, R.A. Cady, and D.E. Bauman. "The environmental impact of dairy production: 1944 compared with 2007." Journal of Animal Science, 2009

# Slow / Difficult to improve traits



- Traits measured in **only one sex** (milk yield)
  - Need phenotypic records on relatives (progeny)
- Traits measured **late** in life (longevity) or **after death** (meat quality)
- Measuring the traits is **expensive** or **difficult** (feed efficiency, disease resistance)

# Rate of genetic change



- Depends on 4 factors:

- Selection **intensity**

- How choosy we are in selecting individuals as parents
    - Can improve (increase) through management

- **Accuracy** of genetic prediction

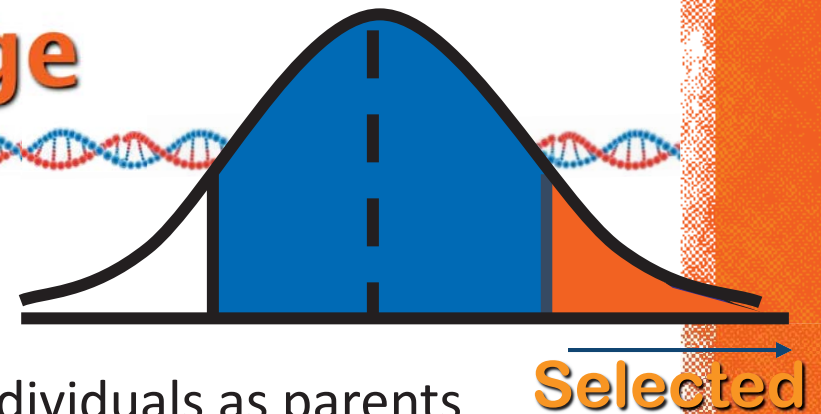
- How close the EBV is to the true BV
    - Can improve (increase) through more/better records

- **Generation interval**

- Time between 2 generations
    - Can improve (decrease) through management or genomic selection

- Amount of **genetic variation** in the trait

- Genetic variation in a population (constant over short period of time)



# How to achieve high accuracy?



Historically:

- Number of records: large number of animals and high-quality phenotypic records
- Trait is highly heritable

## Slow/Difficult to improve traits



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# MAS – marker assisted selection



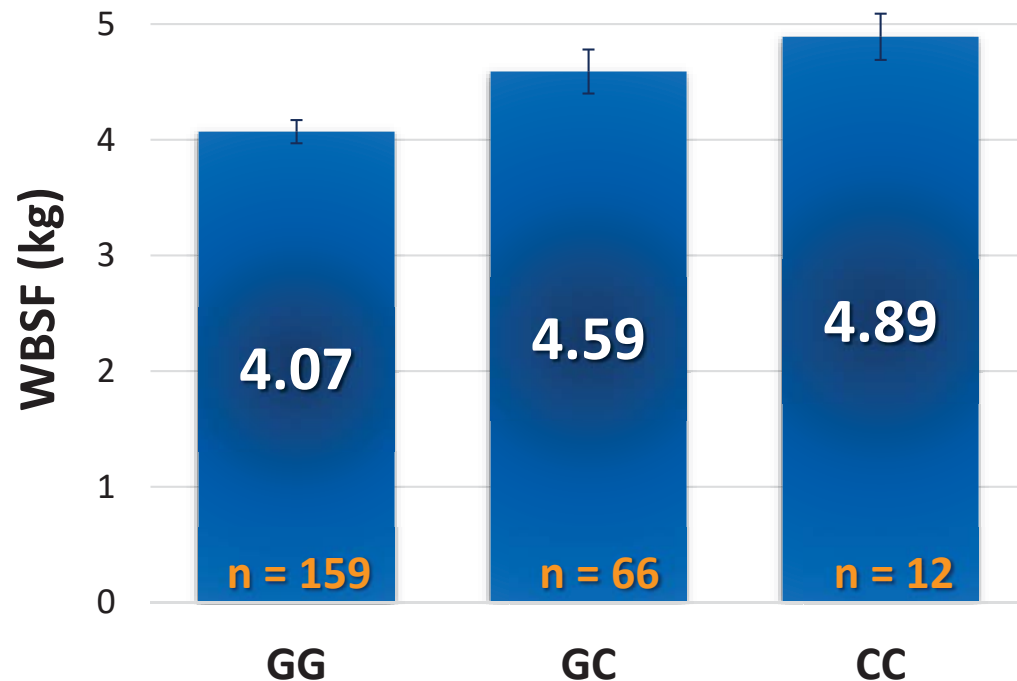
- Since 1990s – DNA information can increase the rate of genetic improvement.
- Challenges:
  - The effect of individual markers (QTLs) on complex traits is small
  - A large number of markers (QTLs) are necessary to explain the genetic variation
  - Marker information in outbreeding species is limited – linkage phase between marker and QTL (gene) must be established for every population

# Tenderness – calpain

GAGTGG AACG **G** CGTGG ACCCT  
GAGTGG AACG **C** CGTGG ACCCT

- CAPN1-316 = marker for tenderness
- One of the SNPs in the **GeneStar** Tenderness test
  - **GG** was **1.10** kg **tougher** than GC (*Pinto et al., 2010*)
  - **GG** was **0.36** kg **tougher** than GC (*Curi et al., 2010*)
  - **CC** is **1.23** kg **tougher** than CG (*UF multibreed pop., Casas et al., 2010*)

	<i>Bos taurus</i>	<i>Bos indicus</i>
<b>GG</b>	tougher	
<b>GC</b>		
<b>CC</b>		tougher



# Potential Benefits of Genomics



- Benefits are greatest for economically important traits that:
  - Are difficult or expensive to measure
  - Measured late in life or after death
  - Not currently selected for because are not routinely measured
  - Have lower heritability
- Benefits:
  - Determine the value of animal at birth
  - Increase accuracy of selection
  - Reduce generation interval
  - Increase selection intensity
  - Increase rate of genetic gain