



Pfizer Animal Health
Animal Genetics

The Genetic Selection Toolbox – Feed Efficiency

**Mark Allan
Ronnie D. Green**

**NCBA Cattlemens College
Phoenix, AZ
January 2009**



GeneSTAR[®]



Pfizer Animal Health
Animal Genetics

GeneSTAR®

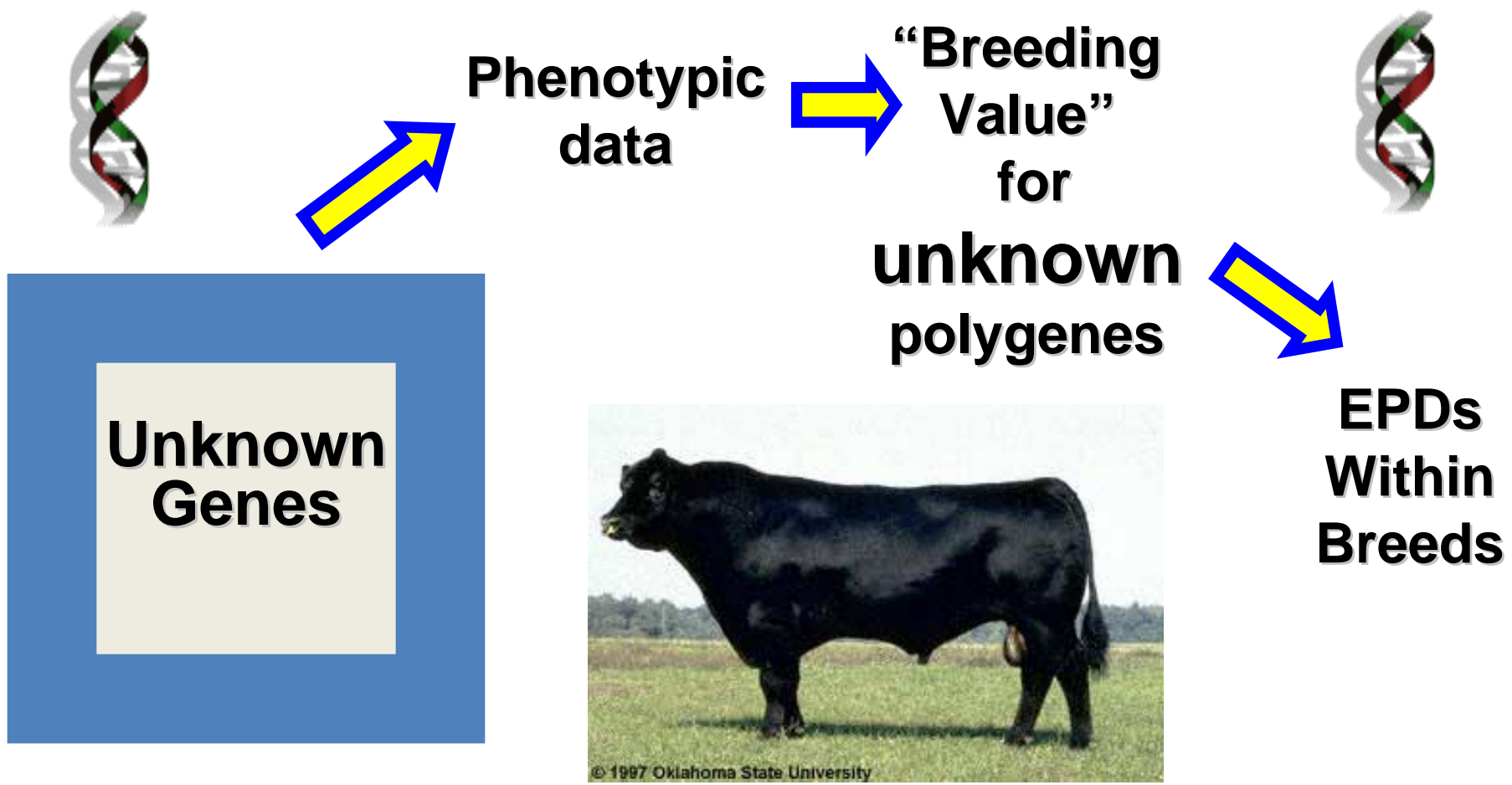
Presentation Road Map

- **Historical Overview of Genetic Selection**
- **Genomics Technology Coming of Age**
- **Technology – Additions to the Breeder's Toolbox**



Quantitative Genetics

Reduced Animal Model



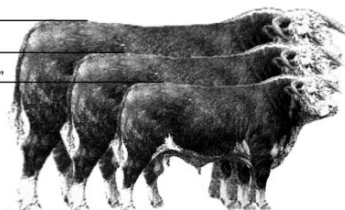
Traditional Selection Works Well



Modern 7 ~ 60"

Classic 2 ~ 50"

Miniature 000 ~ 40"



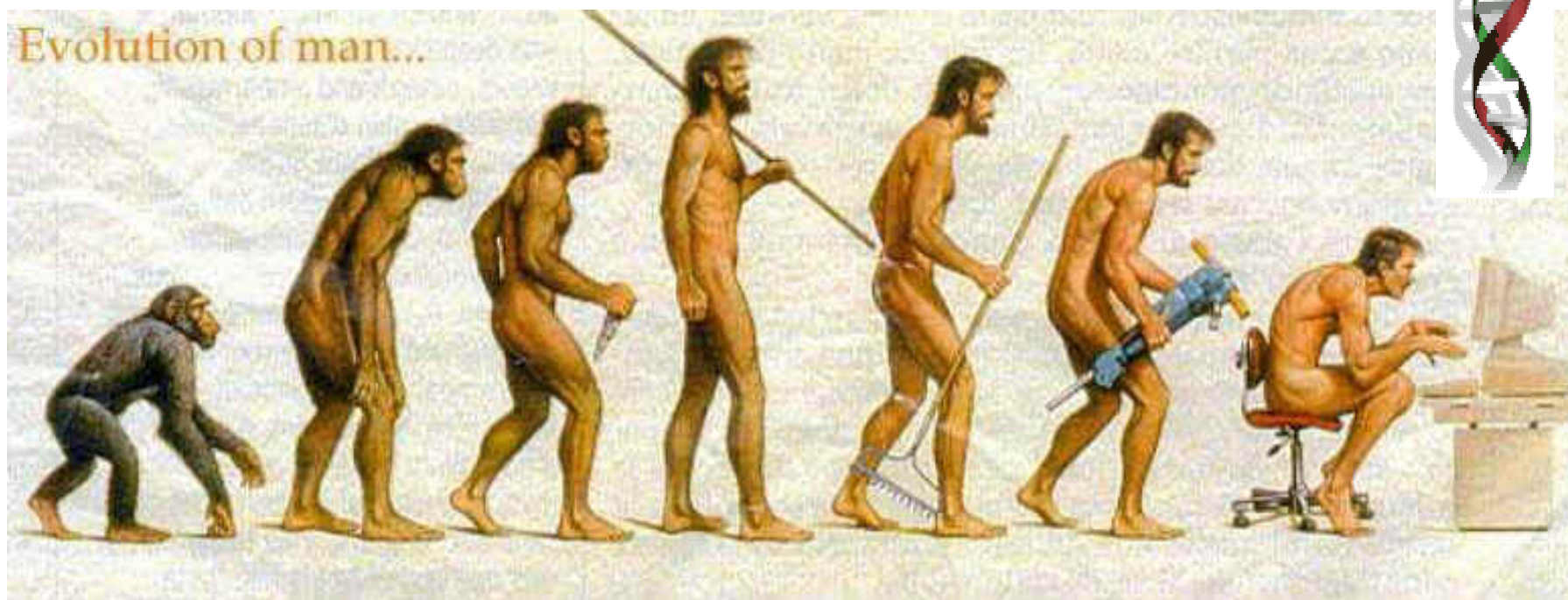
- Visual
- Performance Data **
- **EPDs****** AI use
- Pedigree
- **DNA Marker Information**
- Economic Indexes

Production						Maternal					
CED Acc	BW Acc	WW Acc	YW Acc	YH Acc	SC Acc	CEM Acc	Milk Acc	MkH MkD	MW Acc	MH Acc	\$EN
+11	+1.0	+48	+82	-.5	-.76	+6	+13	2			+10.31
.65	.85	.85	.78	.53	.66	.22	.48	3			

Carcass					Ultrasound				
Cwt Acc	Mrb Acc	RE Acc	Fat Acc	Grp Prog	%IMF Acc	RE Acc	Fat Acc	Grp Prog	
+1	+.09	+.29	-.009	1	-.05	+.09	+.018	36	
.10	.11	.10	.09	1	.74	.75	.75	117	

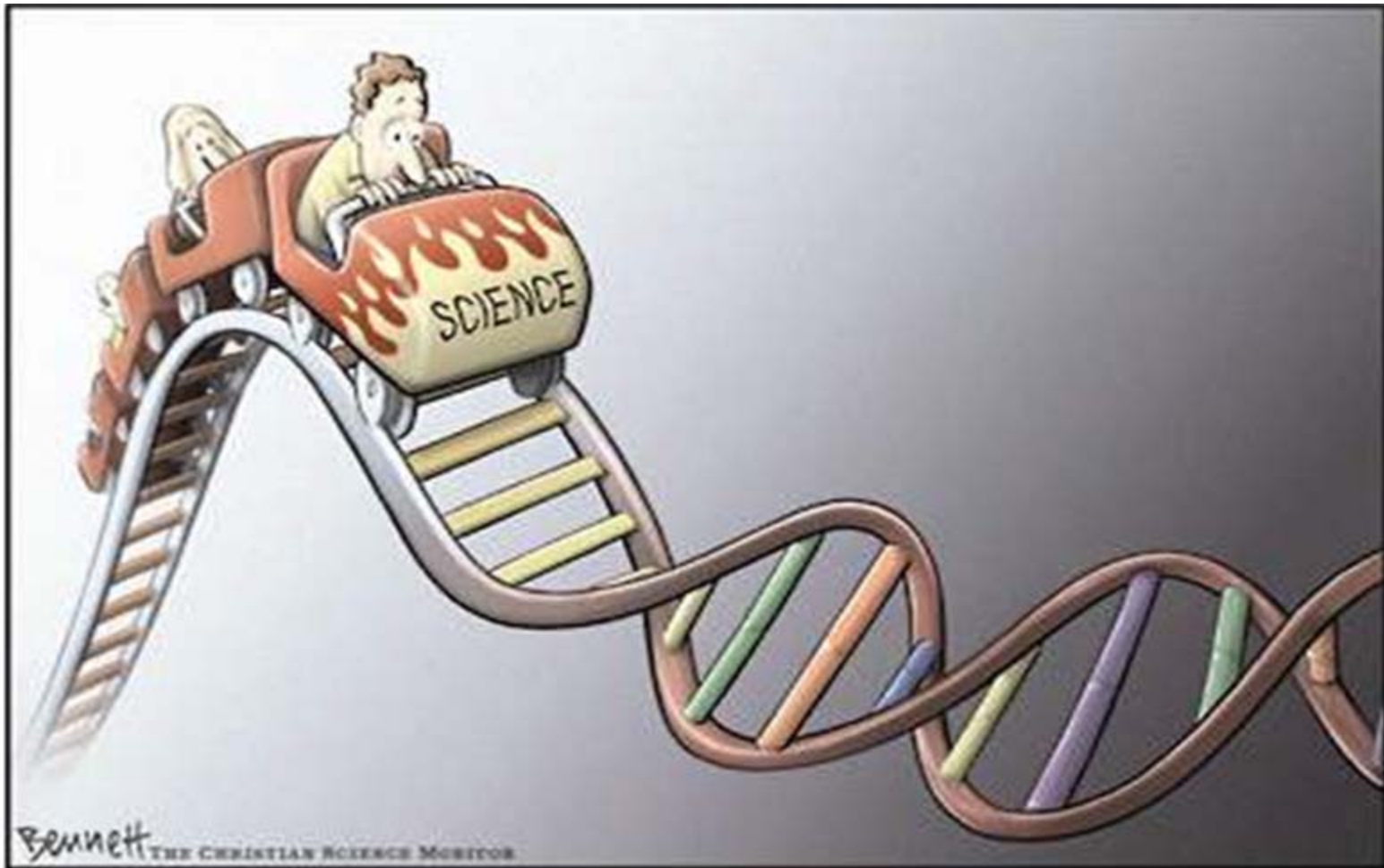
\$Values					
<u>\$W</u>	<u>\$F</u>	<u>\$G</u>	<u>\$QG</u>	<u>\$YG</u>	<u>\$B</u>
+27.96	+22.58	+6.25	+3.73	+2.52	+26.07

We Are Entering Major Transition



<http://www.naute.com/images/evolutionofman.jpg>

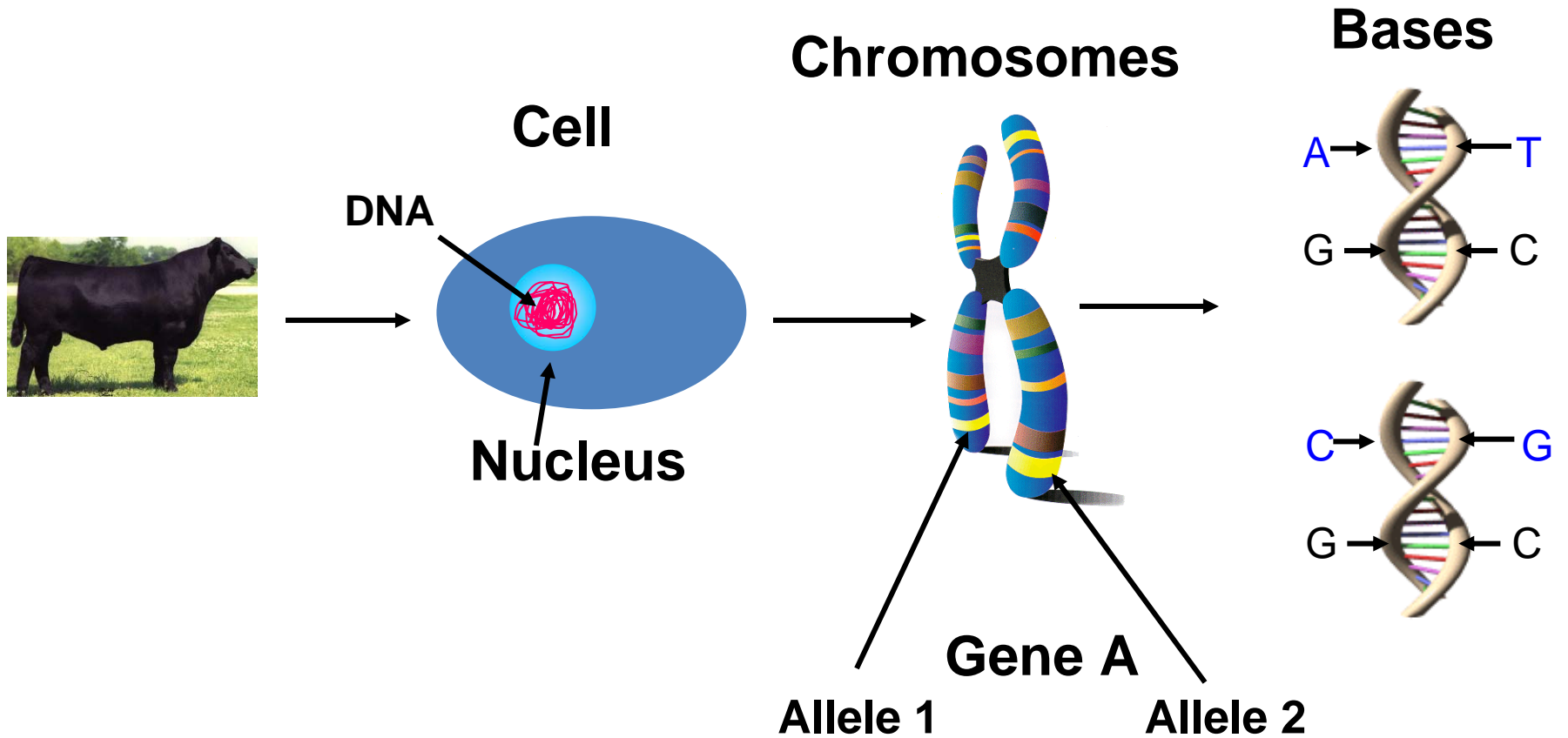
We are on an exciting ride....



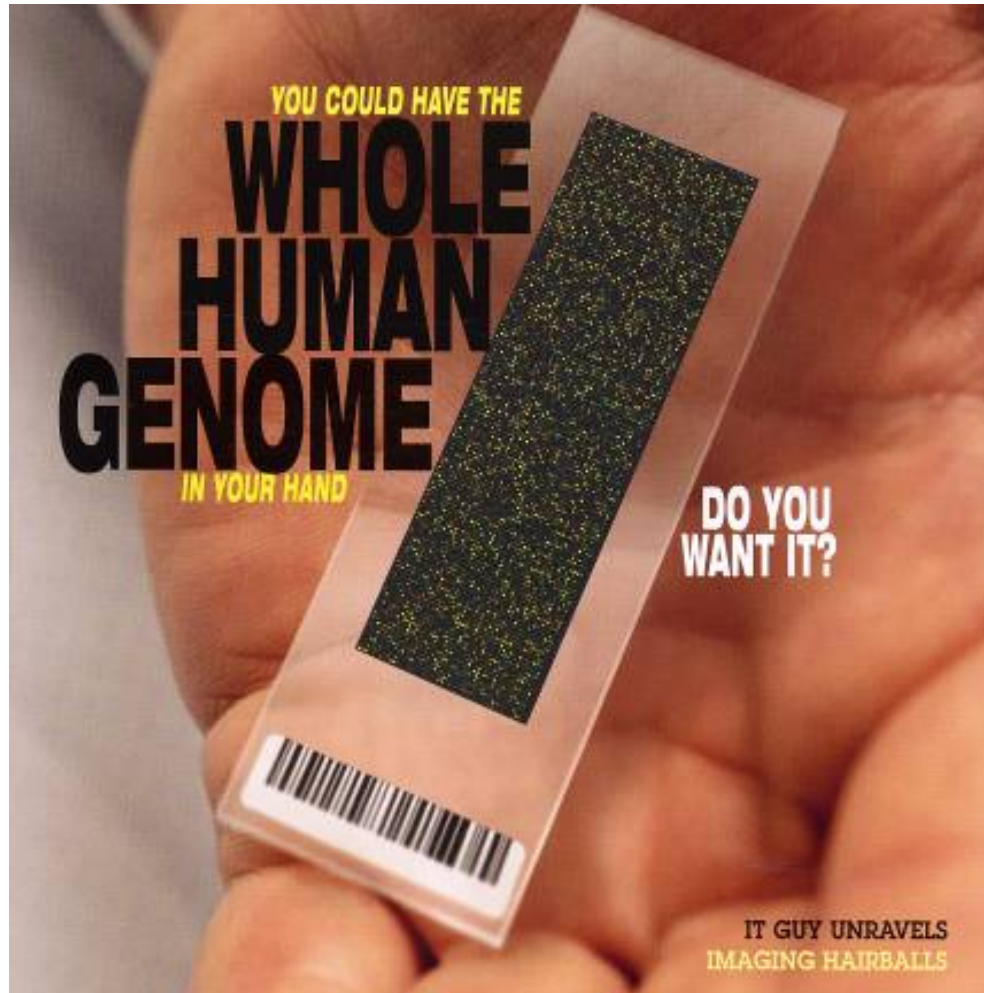
Pfizer Animal Health
Animal Genetics

GeneSTAR[®]

DNA - Technology



Driving Force.....



Pfizer Animal Health
Animal Genetics

GeneSTAR[®]

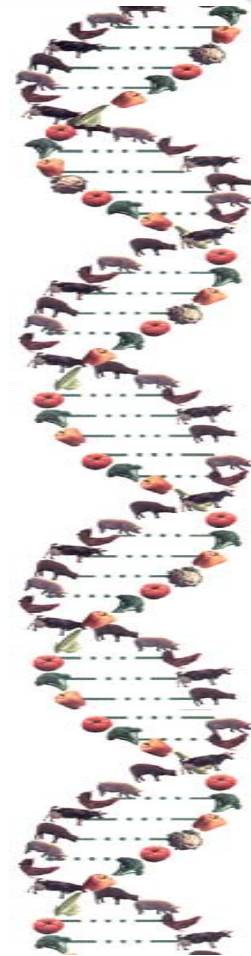
Genome Sequence Assemblies



2004



2005



2007



2007

2008-9



2010

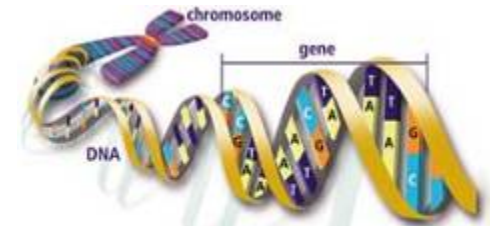


Pfizer Animal Health
Animal Genetics

GeneSTAR[®]

International Bovine Genome Sequencing Project

- **NIH / NHGRI -- \$25M**
- **USDA -- \$11M**
- **State of Texas – \$6M**
- **Genome Canada -- \$5M**
- **U.S. Beef Councils -- \$1.2M**
- **Australia -- \$1M**
- **New Zealand -- \$1M**



Project Total ~ \$50M

Advancements In Technology -- SNPs



Single Nucleotide Polymorphism

- Advantages

Stable (low mutation rates)

Frequent (1 every 300 - 1,000 bases)

Relatively uniformly distributed

Amenable to automation and high-throughput

Costs dropping



ATGCAATTGCC**C**ACGTTGCAAT

ATGCAATTGCT**T**ACGTTGCAAT

ATGCAATTGCC**C/T**ACGTTGCAAT

“50K” Bovine BeadChip - Discovery

~ 54,008 **SNP** markers
across the bovine genome

- On average **SNP** every
<67,000 base pair

- Discovery SNP includes
many breeds

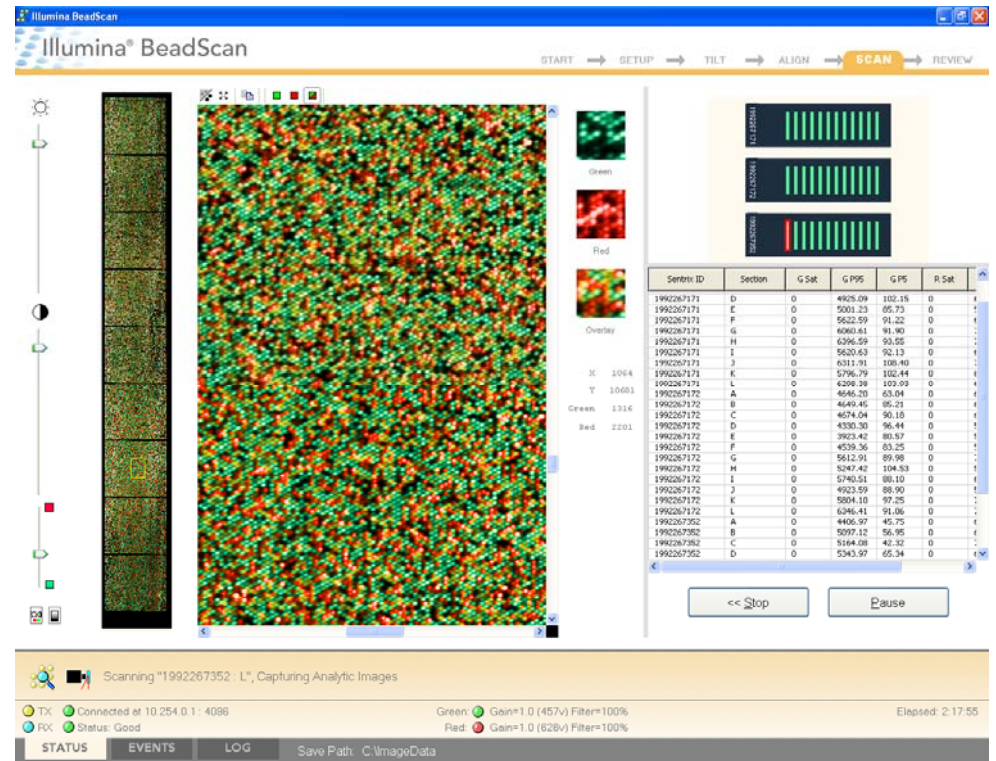
USDA - BARC

US MARC

University of Missouri

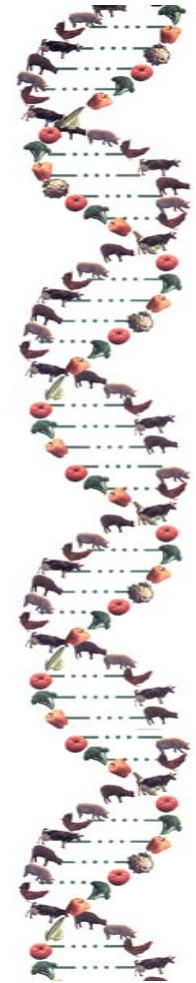
University of Alberta

(Van Tassell et al., 2008 Nature Methods)



Real-world applications.....

- *Trait Prediction – Genetic*
- *M-Inverse Enhancement*
- *Parentage / Com Ranch GE*
- *Precision Management*
- *Supply Chain Management*



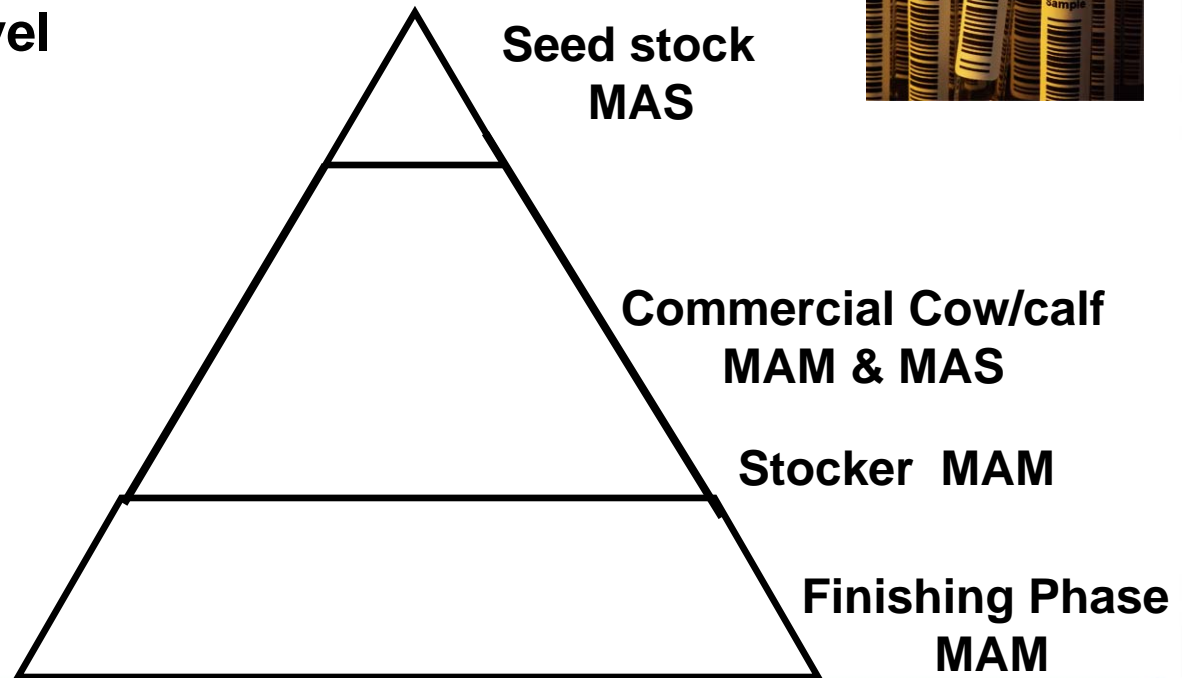
Reach of Marker Technology

Mark-Assisted Selection (MAS)

- At the seed stock/multiplier level

Marker-Assisted Management (MAM)

- At the commercial level



Current Products



GeneSTAR[®]

Feed Efficiency

Marbling

Tenderness



Pfizer Animal Health
Animal Genetics

GeneSTAR[®]

.....Evolving Products.....



- Moving away from individual markers toward panels
- Predictions for genetic improvement or phenotype
 - *“Molecular Value Predictions” with Reliabilities*
- Economic value needs to be demonstrated
- Combining different sources of information (EPD + markers + herd /market analysis)
- Decision support in the production system

“Solutions Beyond Data”



Molecular Value Predictions

MVP → *the sum of individual molecular breeding values for the genotypes of an animal across all marker loci evaluated (N=56).*

- Individual marker effects assumed to be additive, but not necessarily equal
- Reliability maximized by fitting marker additive effects as a regression on the number of favorable alleles

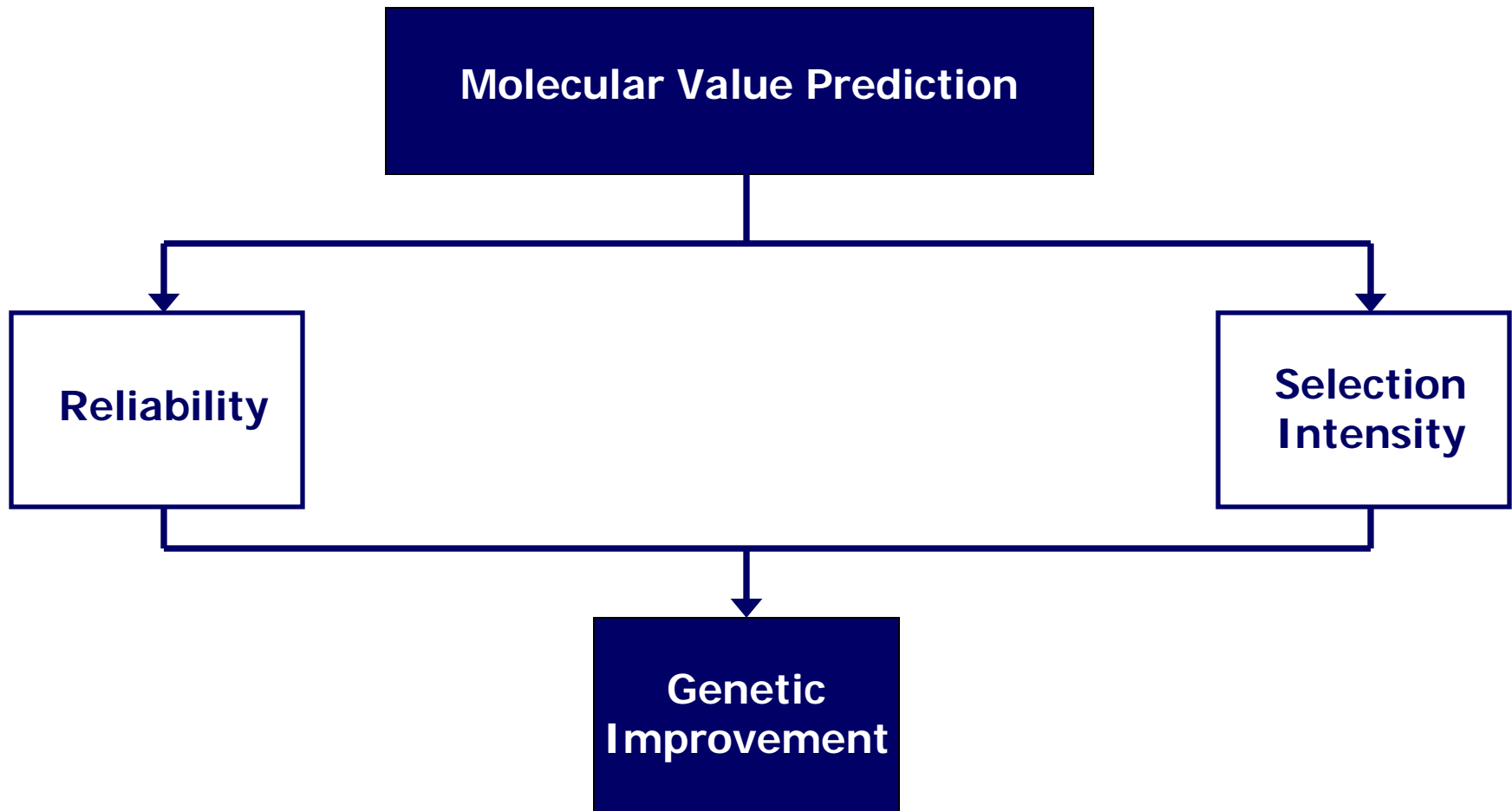


Stars to MVPs



11 individual markers	56-marker panel
Number of stars	Greater selection opportunity
Number of favorable alleles	Collective effects of markers into a trait prediction
Equal effects of each marker	Accounts for individual marker effects
No calculated reliability	% reliability described 26 – 49%
Not benchmarked against breed	Breed average and percentile ranks provided
Good selection tool	A new era in genetic improvement

Driving Genetic Improvement



	MVP Range	% Reliability
Feed Efficiency (lbs.)	★ 5.55	30%



 **PIONEER**
BRAND PRODUCTS
33B51


 **PIONEER**
33H68

 **PIONEER**
33P67

 **PIONEER**
33A72

 **PIONEER**
33R7

Real world example – two bulls

Feed efficiency of two bulls in a commercial cow herd:

- Two Bulls – Genetic Difference
 - Bull A: -0.73 lbs. (30% rank)
 - Bull B: 0.60 lbs. (80% rank)
- } **1.33 lb/day**
- Feeding Period: 150 DOF x 1.33 lbs. = 199.5 lbs. feed
 - Feed Cost: \$200/ton x 199.5 lbs. = **\$19.95/hd**

Real world example – two bulls

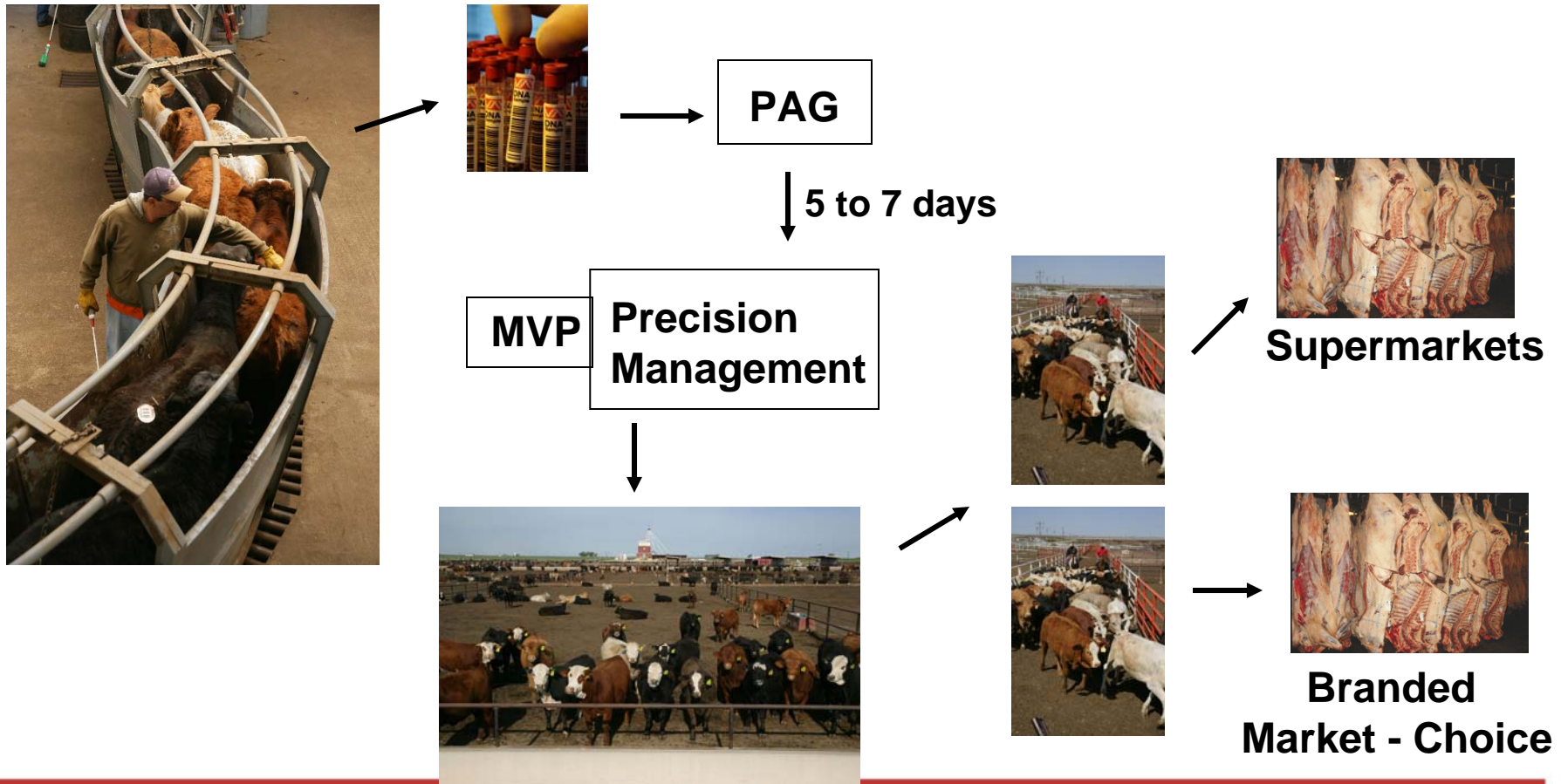


Difference between a low and a high MVP:

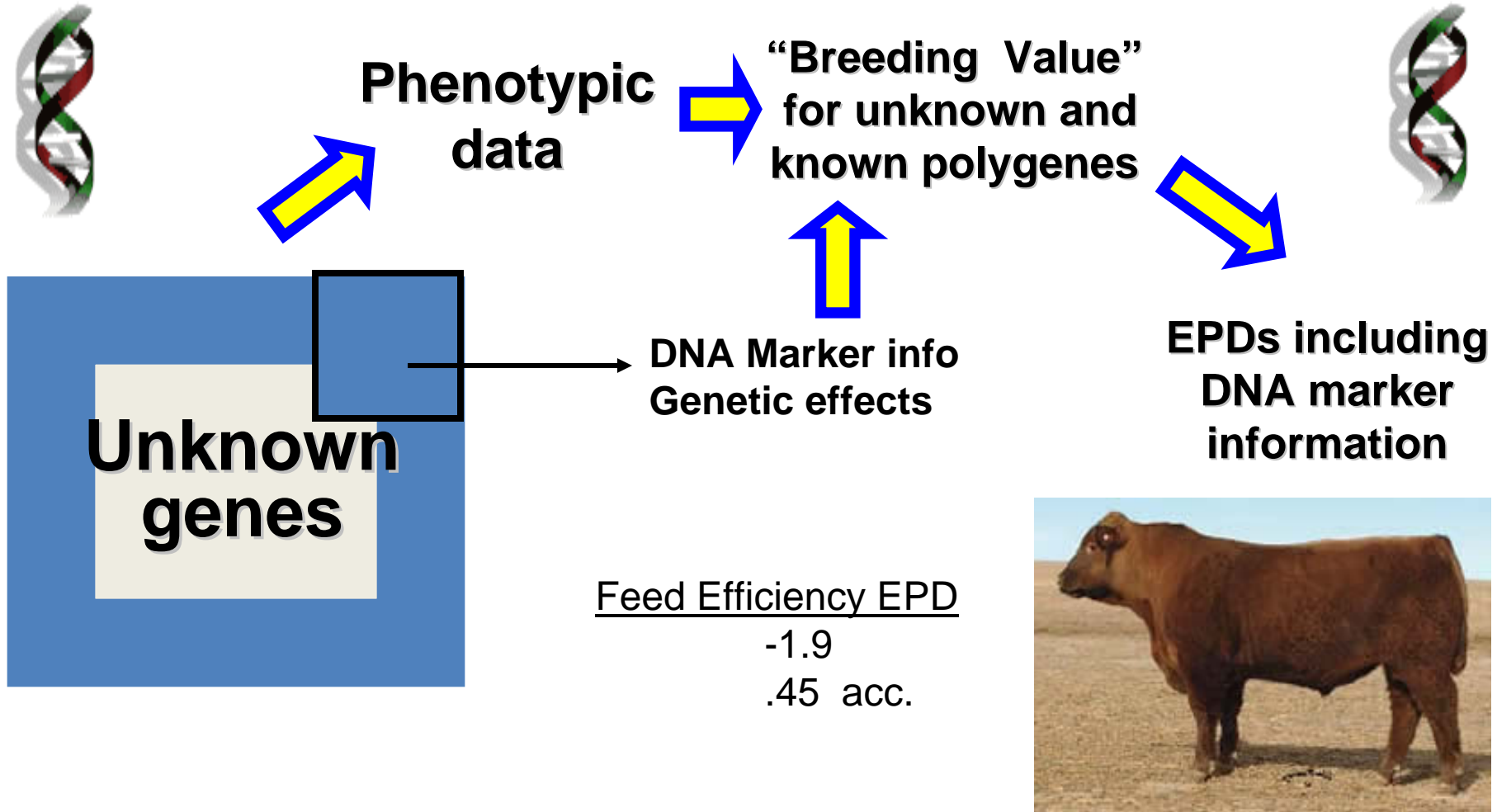
- Average lifetime of 5 years
 - 25 calves per year from average dams
 - 90% survival rate on calves
- } **112 feeder calves**
- Value Difference = $\$19.95 \div 2^* \times 112 =$ **\$1117**

*Divided by two to account for sire influence only in the calf

Example MAM - Quality Grade



Quantitative Genetics- “New” Reduced Animal Model



“Common Sense Interdependencies”





www.pfizeranimalgenetics.com