

#### "Integrating DNA Information into Beef Cattle Production Systems"

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#### http://animalscience.ucdavis.edu/animalbiotech

Van Eenennaam 1/11/2011

Animal Genomics and Biotechnology Education



## **Expert Predictions**

"We believe DNA marker profiles will become widely used in livestock in the near future as the cost decreases and the benefits increase. In fact, a major research objective may be to make best use of this DNA data in commercial animal production"

Goddard, M. E., and B. J. Hayes. 2007. Genomic selection. Journal of Animal Breeding and Genetics 124: 323-330.



#### "How do you make cost-effective use of DNA information in commercial animal production?"

**GOAL:** Determine how DNA-based information is best incorporated into commercial cattle production systems

- 1. Which of several incorporation methods is best?
- 2. Which is feasible for commercial ranches to use?
- 3. Which provides the most/any economic benefit?
- Research objectives: Determine association between breedassociation genetic predictions (EPDs), and DNA-based genetic predictions (stars, scores, MBVs, MVPs, GEPDs) and evaluate their ability to predict the genetic potential of 125 commercial sires based on the performance and carcass records of their offspring
- Extension objectives: Develop and deliver educational materials to a national audience on the integration of DNA information into beef cattle selection programs.





Ranch resources/collaborators on "Integrating DNA information into beef cattle production systems"

Four ranches on this project (UC Davis and<br/>3 commercial cooperators in Siskiyou Co.)- Cowley 900 (550 Spring; 350 Fall)45- Kuck 500 (200 Spring; 300 Fall)16- Mole-Richardson 700 (Fall)40- UC Davis 300 (Fall)26

Approximately 125 Angus bulls, and 2,400 cows per year on project

#### Happy Cows come from Siskiyou County

## **Cowley Ranch**

## Kuck Ranch

#### Mole-Richardson Farms

A





## Work flow and collaborators

- DNA on all bulls goes for 50K whole genome scan collaboration with Jerry Taylor (MO) and John Pollak (Meat Animal Research Center (NE)
- Molecular breeding value (MBV) prediction of genetic merit based on MARC training data set – collaboration with Dorian Garrick (IA) and Mark Thallman, U.S. Meat Animal Research Center (NE)
- Ranch data including sire groupings, birth dates and weaning weights on all calves, all EIDed, and "DNAed" for parentage determination – collaboration with Dan Drake and producers (CA)
- Steer feedlot in weights, treatments, and carcass traits (Hot weight, grading information and meat sample collected in the processing plant – collaboration with Harris Ranch (CA)
- Compile data and compare three sources of genetic estimates: breed EPDs (bEPDs), commercial ranch EPDs (rEPDs), and MBVs, Kristina Weber, UC Davis, PhD student





## **Commercial ranch applications**



#### **Problems experienced included**



## Tail should be here



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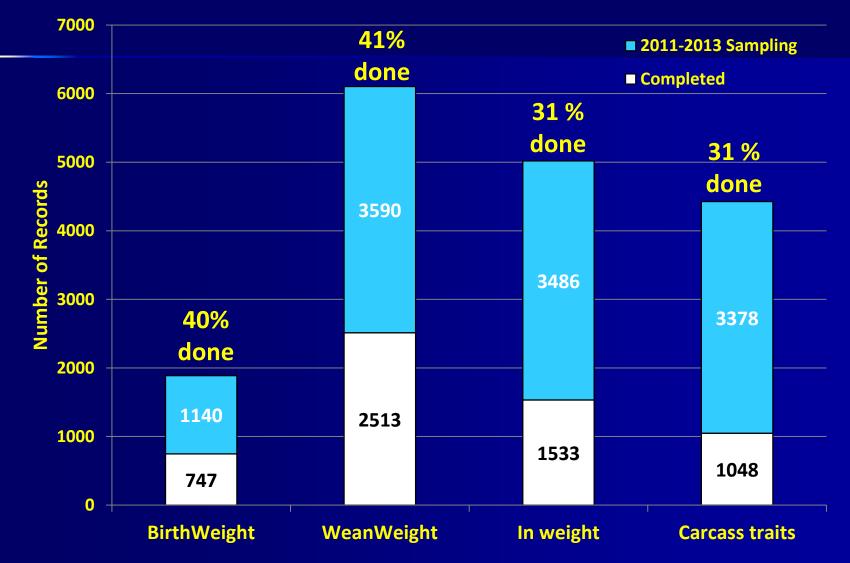
#### **Sampling Summary by Herd: Data Collected to date**

Herd	Season	Birth Weight	WeanWeight	In weight	Carcass traits
Cowley	1/4/2006				Completed
	9/1/2006				Completed
	1/1/2007				Completed
	9/1/2007		Completed	Completed	Completed
	1/1/2008		Completed	Completed	Completed
	9/1/2008		Completed		Completed
	1/1/2009		Completed		Scheduled 1/24/2011
	9/1/2009		Completed	Completed	
	1/1/2010		Completed		
	9/1/2010				
	1/1/2011				
	9/1/2011				
Kuck	1/1/2009	Completed	Completed	Completed	
	10/1/2009	Completed	Completed		
	1/1/2010	Completed	Completed		
	9/1/2010				
	1/1/2011				
	9/1/2011				
Mole-Richardson	9/1/2009		Completed	Completed	
	9/1/2010				
	9/1/2011				
UC Davis	8/15/2009	Completed	Completed		
	8/15/2010				
	8/15/2011				



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#### Sampling Summary by Herd: Total Number of Records







#### "Integrating DNA information into Beef Cattle Production Systems" USDA Integrated Grant Collaborators

- Dr. Darrh Bullock, Extension Professor, University of Kentucky, KY
- Dr. Leslie "Bees" Butler, Extension Marketing Specialist, UC Davis, CA
- Dr. Daniel Drake, University of California Cooperative Extension Livestock Advisor, CA
- Dr. Dorian Garrick, Professor, Iowa State University, IA
- Dr. John Pollak, Professor, Cornell University, NY
- Dr. Mark Thallman, US Meat Animal Research Center, Clay Center, NE

#### Graduate Students

- Kristina Weber, Ph.D. Candidate, UC Davis, CA and Krista Cooprider, MS Candidate, UC Davis, CA
   Producer Collaborators:
- Jack Cowley, Cowley Rancher, Siskiyou County, CA
- Dale, Greg, and Richard Kuck, Kuck Ranch, Siskiyou County, CA
- Matt Parker and Scott Dutcher, Mole-Richardson Farms, Siskiyou County, CA
   Processor Collaborators:
- Harris Ranch Beef Company, Coalinga, CA
- Los Banos Abattoir, Los Banos, CA

#### Software Collaborators:

- Jim Lowe, Cow Sense Herd Management Software, NE
- Other Contributors/Collaborators
- Dr. Jerry Taylor, University of Missouri, MO
- Dr. Mike Goddard, University of Melbourne and Victorian DPI, Australia



## **DNA-based tests for cattle**



#### What is working well

- Identification of genetic defects
- Parentage

#### What is not working well (at present)

- Genetic markers for quantitative traits
- Genomic selection in beef cattle



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## **Genetic Defects in Cattle**

#### Images from an article by David S. Buchanan, Department of Animal Sciences, North Dakota State University

http://www.ag.ndsu.edu/williamscountyextension/livestock/genetic-defects-in-cattle



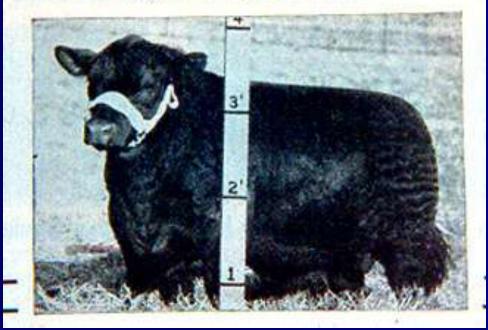


# Compare dwarfism response in th 50s to the response to curly calf (AM)



#### A Triple Compounded EILEENMERE

Look at the Blood Line Wealth concentrated in one animal. "SHORT SNORTER"



An early '50's advertisement that superimposed a measuring stick in the picture of this bull who was nicknamed "Short Snorter." Based upon his height and age, he was less than a frame score 1. The choice of a nick-name in this instance was unfortunate because "snorter" dwarfism would soon devastate the purebred beef industry.

Image from https://www.msu.edu/~ritchieh/historical/shortsnorter.jpg



In 2008, September 8 – November 3 researchers identified genetic problem causing curly calf, developed AM test prototype, and released carrier status of 736 bulls by the end of the year!

- In the 10 months following the release of the test, the AAA posted the results of tests for AM on about 90,000 cattle.
- Of these, almost 5,000 bulls and more than 13,000 heifers have tested as carriers of AM. That leaves more than 22,000 bulls and more than 50,000 heifers which tested as free of AM.
- These tests generally cost less than \$30. While the total cost is substantial it is miniscule when compared with the cost of simply eliminating all descendants of the widely-used GAR Precision 1680 (AA Reg. No. 11520398), and his grandsire Rito 9J9 of B156 7T26, (AA Reg. No. 9682589) from the Angus breed.



# AM test can be done at the following labs

#### The following labs are authorized for AM AgriGenomics 2399 N. 1000 E. Rd.

Mansfield, IL 61854 217-762-9808 http://www.agrigenomicsinc.com

#### Click here for Important Notice about Genetic Testing with MMI MMI Genomics 1756 Picasso Avenue Davis, CA 95618 (800) 311-8808 ext 3016

http://www.mmigenomics.com/AM.html

#### **Pfizer Animal Genetics**

333 Portage Road Kalamazoo, MI 49007-4931 1-877-BEEF DNA 1-877-233-3362 Fax: 269-833-1197 http://www.pfizeranimalgenetics.com

#### IGENITY

4701 Innovation Drive, CB 101 Lincoln, NE 68521 1-877-IGENITY 1-877-443-6489 http://www.igenity.com

#### GeneSeek

4665 Innovation Dr. Suite 120 Lincoln NE 68521 402-435-0665 www.geneseek.com

The following groups collaborate with GeneSeek, Inc., to collect and provide samples for AM testing:

 SEK Genetics Don Coover
 9525 70th Rd.
 Galesburg, KS 66740
 don@sekgenetics.com
 Phone: 800-443-6389

- Stockman's Resource Center LLC 2371 330th Street Eddyville, Iowa 52553 <u>stockmansresource@hotmail.com</u> <u>www.stockmansresource.com</u> Office phone: 641-969-4111 Mobile: 641-660-0771
- Genex Cooperative, Inc. Headquarters: 100 MBC Drive Shawano, WI 54166 Phone: 888-333-1783 Fax: 715-526-3219 info@crinet.com



#### From a breeding standpoint there are different possible scenarios when considering this mutation

If both parents are carriers (AMC) AMC x AMC = 1/4 affected (AMA): 1/2 carriers(AMC): 1/4 AM free (AMF)

If only one parent is a carrier, then all of the offspring will be normal appearing, but half of them will be carriers AMC x AMF =  $\frac{1}{2}$  carriers (AMC):  $\frac{1}{2}$  AM free (AMF)

If neither parent is a carrier, AMF x AMF = all AM free (AMF) even if have Precision in pedigree!



# Early extension education about dwarfism explaining carriers and inheritance

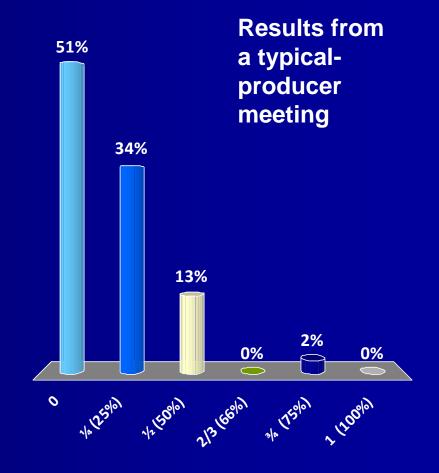


Image from Special CollectionsUniversity Libraries, Virginia Tech: <a href="http://spec.lib.vt.edu/imagebase/agextension/boxseven/screen/AGR3618.jpg">http://spec.lib.vt.edu/imagebase/agextension/boxseven/screen/AGR3618.jpg</a>



If you breed a curly calf carrier cow (AMC) to an curly calf free bull (AMF), what is the chance that the offspring will be stillborn as a result of being curly calf?

1. 0 2. 1/4 (25%) 3. 1/2 (50%) 4. 2/3 (66%) 5. <sup>3</sup>/<sub>4</sub> (75%) 6. 1 **(100%)** 





## **DNA-based tests for cattle**



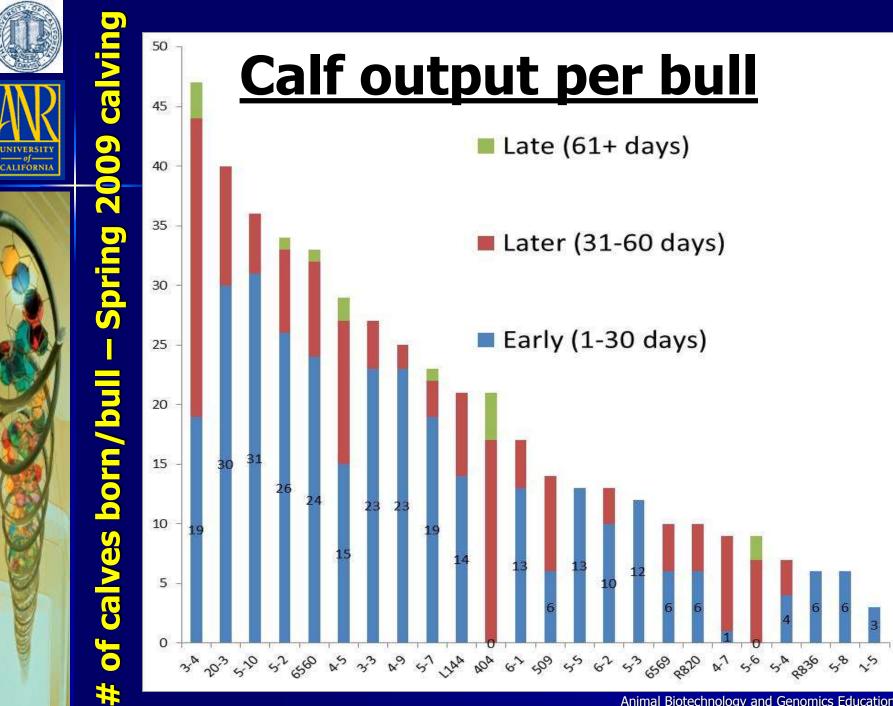
# What is working well Identification of genetic defects Parentage What is not working so well (at present)

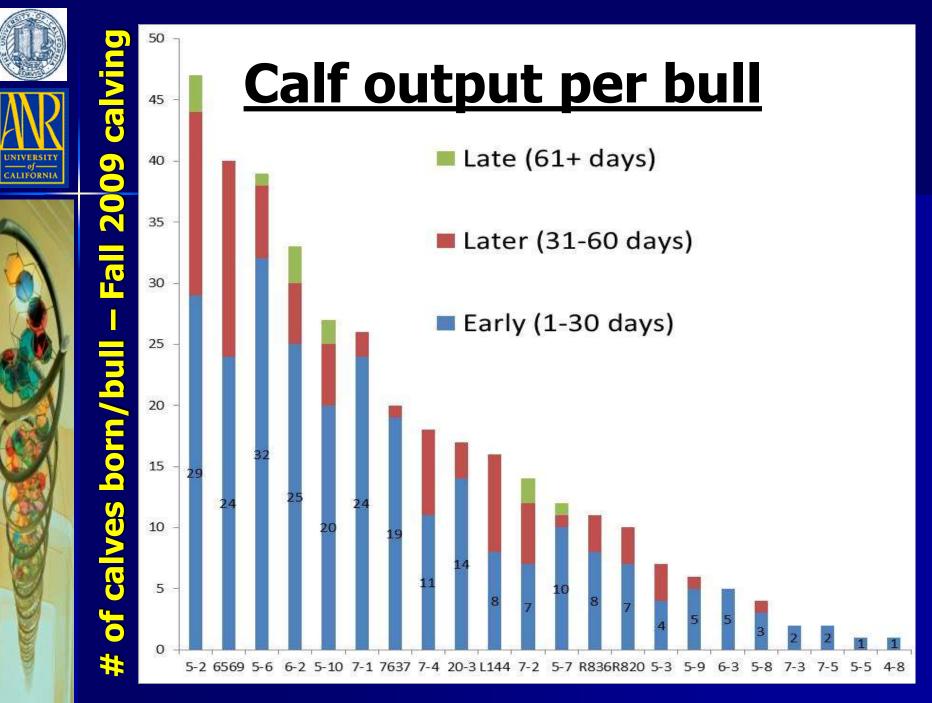
- Genetic markers for quantitative traits
- Genomic selection in beef cattle

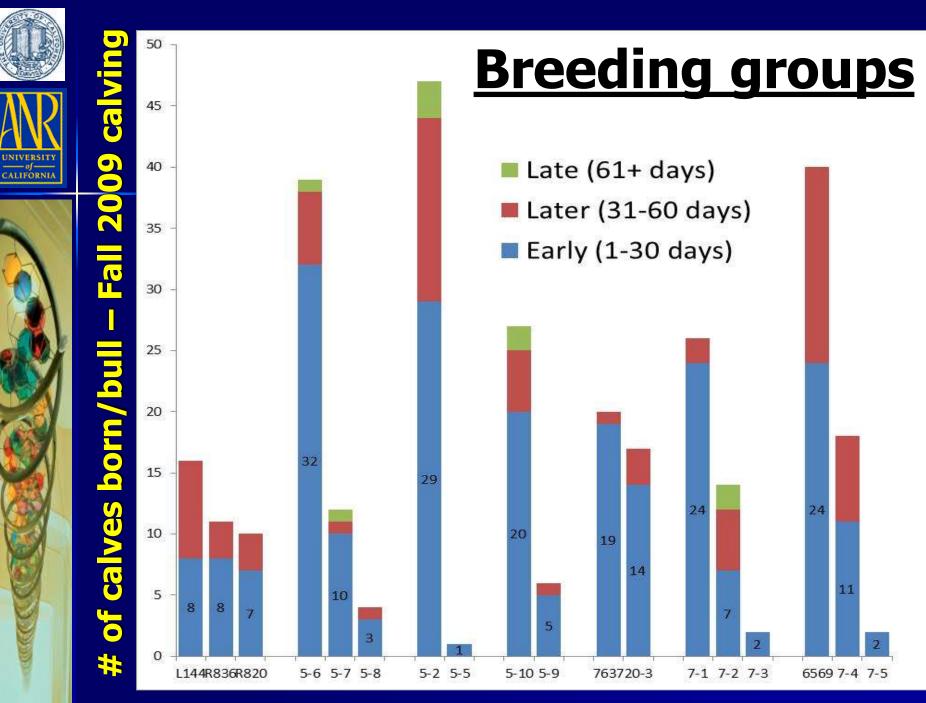


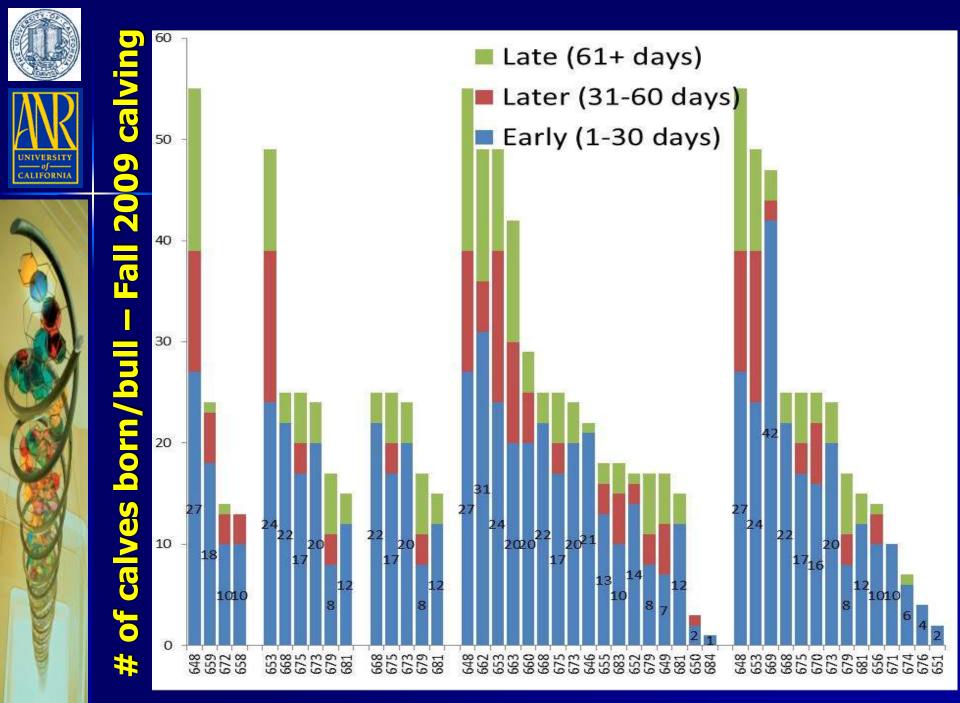
## **Benefits of DNA-based parentage identification**

- Correct pedigree errors so improve the rate of genetic gain
  Enables the use of multi-sire breeding pasture
  - Higher fertility
  - Elimination of sire failure
  - Tighter calving season
- Reduces the need for different breeding pastures
  - Allows for better pasture management
  - Less sorting and working of animals into different groups
- Reduces the need to disturb newborn animals
  - Labor savings so can focus on concentrate on offspring survival
  - Worker safety improvement
  - Better bonding of offspring with dam
  - Can determine which bull is causing calving problems
- Enables the development of commercial-ranch genetic evaluations





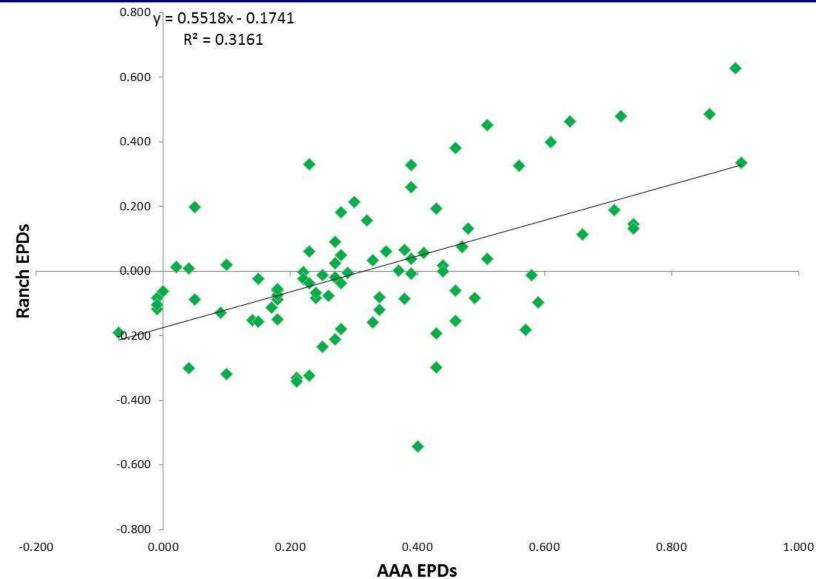




## **Preliminary correlation of EPDs**

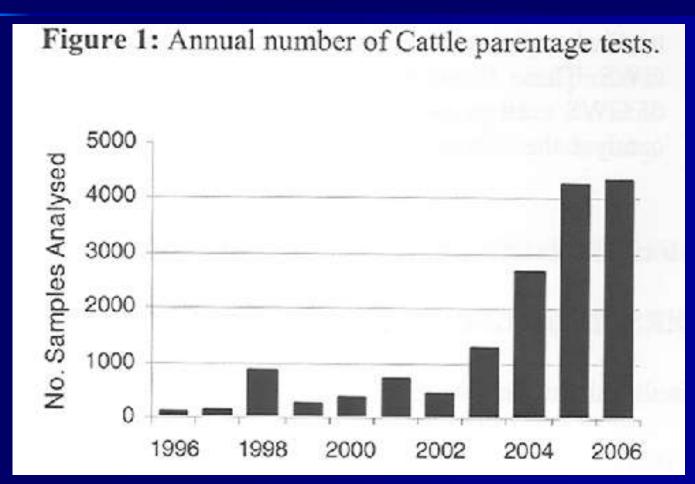
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#### In excess of 30% of animals in the sire breeding tier of some livestock industries in New Zealand are currently being DNA tested for parentage



Crawford, A. M., R. M. Anderson, and K. M. McEwan. 2007. Uptake of DNA testing by the livestock industries of New Zealand. Proceedings of the New Zealand Society of Animal Production 67:168-174 Animal Biotechnology and Genomics Education



## **DNA-based tests for cattle**



#### What is working well

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#### The Power of the IGENITY® profile for Angus

The American Angus Association® through its subsidiary, Angus Genetics Inc.® (AGI), has a vision to provide Angus breeders with the most advanced solutions to their genetic selection and management needs.

Genomic-enhanced Expected Progeny Differences (EPDs) can now be calculated for your animals using the highly predictable American Angus Association database along with IGENITY\* profile results to provide a more thorough characterization of economically important traits and improved accuracy on young animals.

Using the IGENITY profile for Angus, breeders receive comprehensive genomic results for multiple, economically important traits.

- 1. Dry Matter Intake
- 2. Birth Weight
- 3. Mature Height
- 4. Mature Weight
- 5. Milk
- 6. Scrotal Circumference
- 7. Weaning Weight
- 8. Yearling Weight
- 9. Marbling
- 10. Ribeye Area
- **11. Fat Thickness**
- 12. Carcass Weight
- 13. Tenderness
- 14. Percent Choice (quality grade)
- **15. Heifer Pregnancy**
- **16. Maternal Calving Ease**
- **17. Direct Calving Ease**
- **18. Docility**
- 19. Average Daily Gain
- **20. Feed Efficiency**
- 21. Yearling Height



## Lead Today with 50K

- 1. Birth weight
- 2. Weaning weight
- 3. Weaning maternal (milk)
- 4. Calving ease direct
- 5. Calving ease maternal
- 6. Marbling
- 7. Backfat thickness
- 8. Ribeye area
- 9. Carcass weight
- 10. Tenderness
- 11. Postweaning average daily gain
- 12. Daily feed intake
- 13. Feed efficiency (net feed intake)





Pfizer Animal Health Animal Genetics 50K SNP chip assays 50,000 SNPs spread throughout genome



#### Black Angus Sire G A R Predestined



Reg. No.: 13395344 Calved: 8/16/1999 Tattoo: 5899 Semen: \$25 Certificates: \$20 Spring 2010 EPD

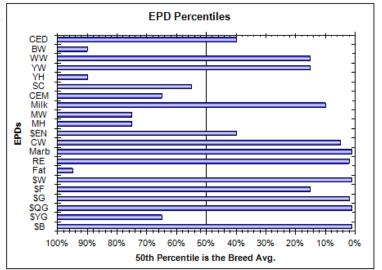
#### G A R Predestined:

From start to finish--conception to carcass--no other bull in the beef business today adds as much real value to cattle as Predestined. Ranking as the #1 bull for \$B in the breed--our customers tell us that their Predestined-sired cattle return the most dollars to their pockets--they know that \$B works. Unlike any other 036 son, Predestined tones down size, adds depth of flank, superior feet and legs and a pleasant disposition to his offspring. His conception rate is high and he's been a standout in timed-Al programs. His progeny look good--his bulls are thick and his heifers are fancy--and they always display additional shape and capacity. He ended 2006 as our top-seller and rightfully so--Predestined's many talents for creating value are for real.

		Produ	ction			Maternal							
CED Acc	BW Acc	WW Acc	YW Acc	10000000000	SC Acc	CEM Acc	Milk Acc	MkH MkD	MW Acc	MH Acc	ENS		
+7 .84	+4.1 .97	+53 .96	+99 .94	+0 .96	+.31 .95	+6 .80	+28 .85	345 1135	+13 .81	+.2 .81	+5.24		

Carcass					Usnd	<b>S</b> Values								
CW Acc	Marb Acc	RE Acc	Fat Acc	Grp Prog	UGrp UProg	Wean	Feedlot	Grid	SQG	\$YG	Beef			
+26 .82	+1.07 .84	+.59 .82	+.046 .81	47 261	4269 11990	37 <b>.39</b>	37.08	38.21	35.04	3.17	69.78			

8	QG1	na	QG2	па	QG3	па	QG4	na	QG GPD	
IIII	T1	*	T2	0	Т3	0	-	<u></u>	T GPD	-0.35
×,	FE1	na	FE2	na	FE3	па	FE4	na	FE GPD	10 10



#### Current Sires Percent Breakdown

Regi	Tend	FalT	Yield	Ribey	Carc	Pero	Marb
Registration	ndern	Thick	ld Gra	eye A	rcass	rcent (	Marbling
ŧ III	ess	ness	qe	rea	Weight	Choice	

As of 03/22/2010

EPDs (CW, Marb, RE, Fat) are enhanced by genomic profiles generated by 🕲 igenity.

G A R Prede	G A R Predestined 1339534														
	CED	BW	WW	YW	ADG	DMI	NFI	CEM	MA	CW	FAT	REA	MS	TND	\$B/\$MVP⁼∟
EPD	7	4.1	53	99	-	-	-	6	28	26	0.046	0.59	1.07	-	69.78
ACC	0.84	0.97	0.96	0.94	-	-	-	0.8	0.85	0.82	0.81	0.82	0.84	-	-
EPD % Rank	30	85	15	15	-	-	-	55	10	4	90	2	1	-	1
MVP	13	1.0	37	-	0.45	0.97	0.04	8	33	55	0.07	0.92	1.52	-0.43	243
MVP % Rank	3	70	10	-	30	90	90	4	1	1	90	1	1	80	1



## **DNA tests for selection**



#### **Bad News**

- Tests are breed specific only Angus
- Data reporting is varied and hard to interpret
- No independent estimate of test accuracy

#### **Good News**

- Larger SNP panel (700K) might help tests work across breeds
- DNA information is stating to get integrated into EPDs (Angus)



#### **Beef Improvement Federation (BIF)**

"BIF believes that information from DNA tests only has value in selection when incorporated with all other available forms of performance information for economically important traits in NCE, and when communicated in the form of an EPD with a corresponding BIF accuracy."



<u>Manual</u> - National Beef Cattle Evaluation Consortium (2010) UPDATED 2010

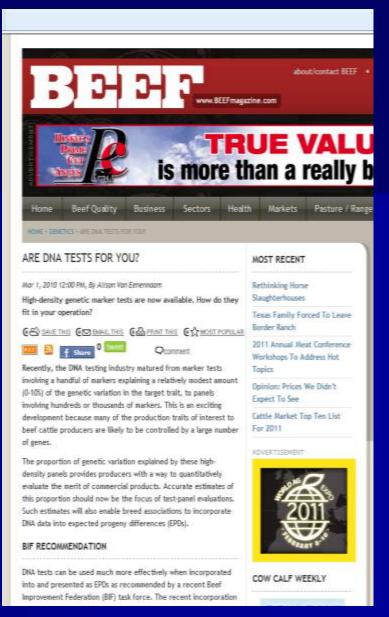
Marker-Assisted Selection **DNA** Companies Commercially-available DNA Tests for Beef Cattle (06/10) Value of DNA Information for beef bull selection (6/10) Are DNA tests for you? Beef Magazine (3/10) DNA markers... Revolution or Evolution? ABS Breeders Journal (Fall/Winter 2009) Do DNA tests work? Beef Magazine (10/09) Basics of DNA Markers and Genotyping (6/09) DNA-Based Progeny Testing (6/09) Fundamentals of Expected Progeny Differences (6/09) Marker-Assisted Selection in Beef Cattle Handout (6/09) The Value of Improving Accuracy of Yearling Bulls (6/09) Validation of Marker Tests (6/09) Whole Genome Selection (6/09) 2009 Beef Improvement Federation Conference Proceedings Curly Calf Syndrome (Arthrogryposis Multiplex (AM)) Update (2/09)

Aarker Assisted Selection

niversity of California, Davis

Cattlemen to Cattlemen streaming video (5/08)

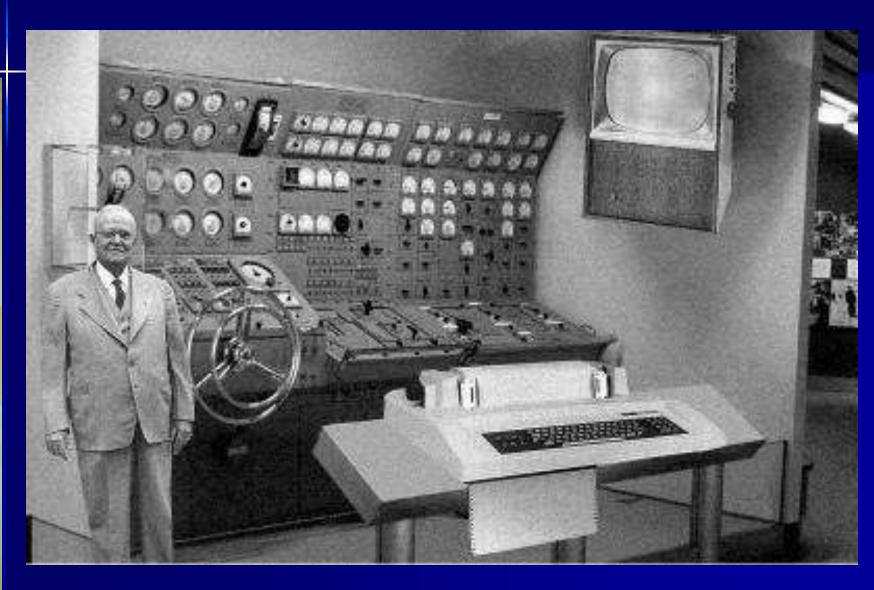
"No Bull" Discussion on Genetic Markers (5/08)



http://animalscience.ucdavis.edu /animalbiotech/Biotechnology/ MAS/index.htm



#### "1954 version of what 'home computers' might look like in 50 years time (i.e. 2004)"





## **Wrong Expert Predictions**

# I think there's a world market for about five computers.

Thomas J. Watson, chairman of the board of IBM. 1943

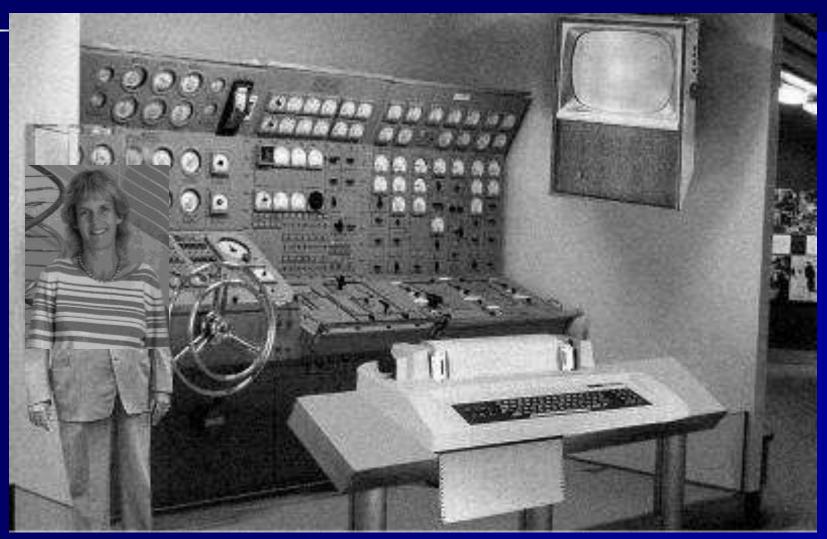
## There is no reason anyone would want a computer in their home.

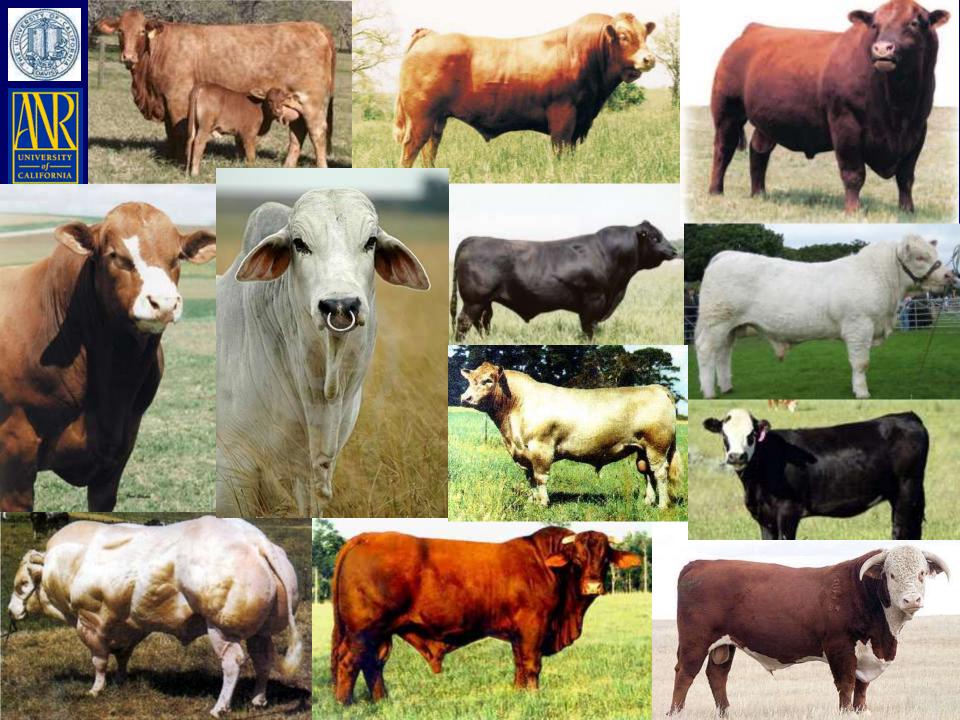
Ken Olson, president of Digital Equipment Corp. 1977

The cost for a "large" genome scan (defined as 18 chromosomes\* 7 chromosome (i.e. 126 markers!) \* \$4/marker) = \$504 Ben Hayes and Mike Goddard, 2003. Evaluation of marker assisted selection in pig enterprises. Livestock Production Science 81:197-211.











United States Department of Agriculture

National Institute of Food and Agriculture

"This project is supported by National Research Initiative Grant no. 2009-55205-05057 from the USDA Cooperative State Research, Education, and Extension Service Animal Genome program."

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2-5 May 2011 The Sebel Albert Park Melbourne

www.smogenomics.org

genomics-conf@jkconnections.com.au



# **Questions?**