

# Optimizing Breeding Programs

## COST-BENEFIT

Armidale Animal Breeding Summer Course 2014

# Cost - Benefit of breeding programs

## Cost of breeding programs for genetic improvement

Fixed costs (logistics, scientists etc. etc.)

Cost related to breeding strategy

- cost of phenotyping
- cost of genotyping
- cost of reproduction

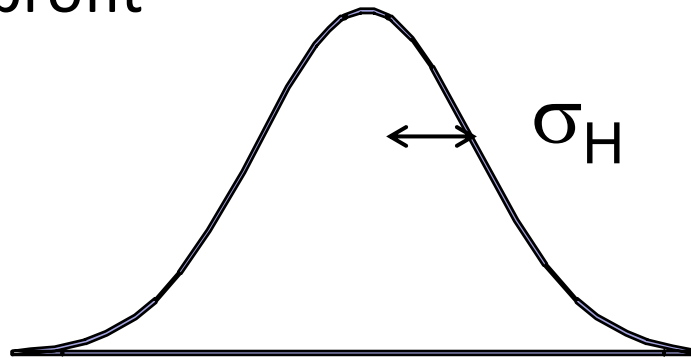
## Benefits

Benefit of more genetic gain

Market share

# Benefits of genetic gain

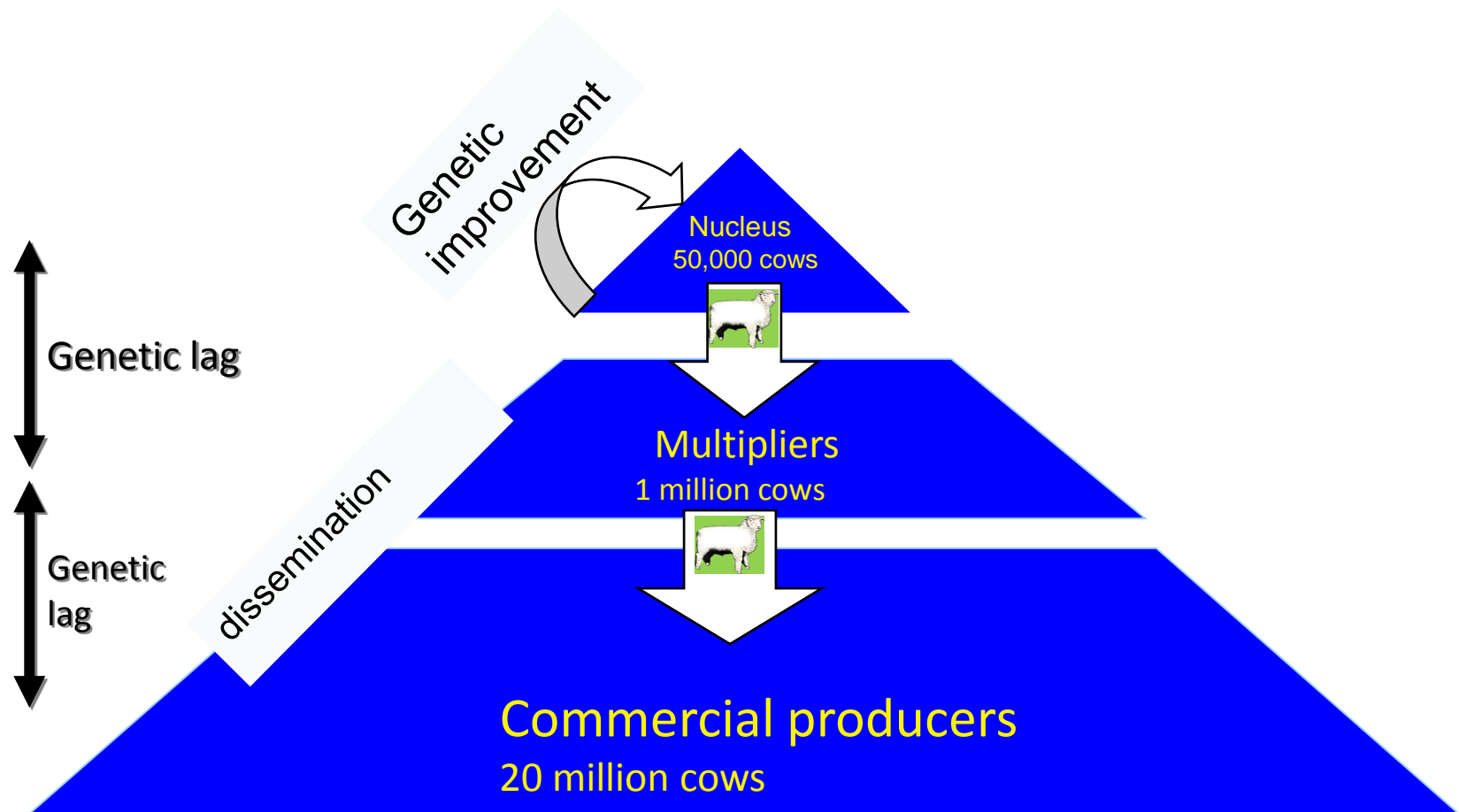
- Assuming the benefit is expressed by the breeding objective (economic values of trait improvement)
- Variation in breeding objective is variation in genetic merit for profit



Difference between best and worst is about  $6 \sigma_H$

# Benefits of genetic gain

- Benefit is transmitted is multiplied over many animals



# Benefits of genetic gain

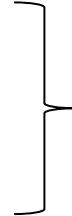
- Benefit is cumulative

[illegible]

# Benefits of genetic gain

Benefit is

- Cumulative
- Multiplied over many



Benefits can be expected to be large

• But:

- Are they achieved?
- Who gets the benefit?
  - Breeders, Producer? Retail? Consumer?

# Economic value of genetic improvement

- Value difference between two bulls
- Value of selecting better bulls
  - Bulls sold to Commercial
  - Bulls used in Stud
- Value of genetic improvement – whole herd

## Two Commercial Bulls

### EBV YWT

Bull 1: Kevin +10 kg

Bull 2: Tony +15 kg

Nr Progeny: 100

Value of 1 kg YWT \$4

Difference in progeny 2.5 kg

Difference in value:

as commercial bulls

= \$1000.-

5\*\$4

Selection  
Difference

\* 100

Nr of  
Progeny

\* 0.5

Expression  
per progeny



## Two Commercial Bulls

\$Index

Bull 1: Kevin

+190

Bull 2: Tony

+180

Nr Progeny:

100

Difference in progeny

\$5

Difference in value:  
as commercial bulls

\$10

\* 100

\* 0.5

Selection  
Difference

Nr of  
Progeny

Expression  
per progeny

= \$500.-

# Selecting Better Bulls

		<u>\$Index</u>	
Average of 100 rams sold:	With Genomics	+182	
	No Genomics	+180	
Nr Progeny:	100 per bull		
Difference in progeny	\$1.0		
Difference in value:	\$2	* 100	* 0.5
as commercial bulls	Selection Difference	Nr of Progeny	Expression per progeny
= \$100.- * 100 rams = \$10,000.			

# So principles are

## Value of a superior bull

= Selection Difference \* Nr.Progeny \* expressions per progeny

We look at all expressions in commercial progeny

To evaluate benefit we need to predict

- the extra Selection Difference we can get  
*this will depend a lot on extra accuracy*
- the number of expressions

# How about selection of stud bulls?

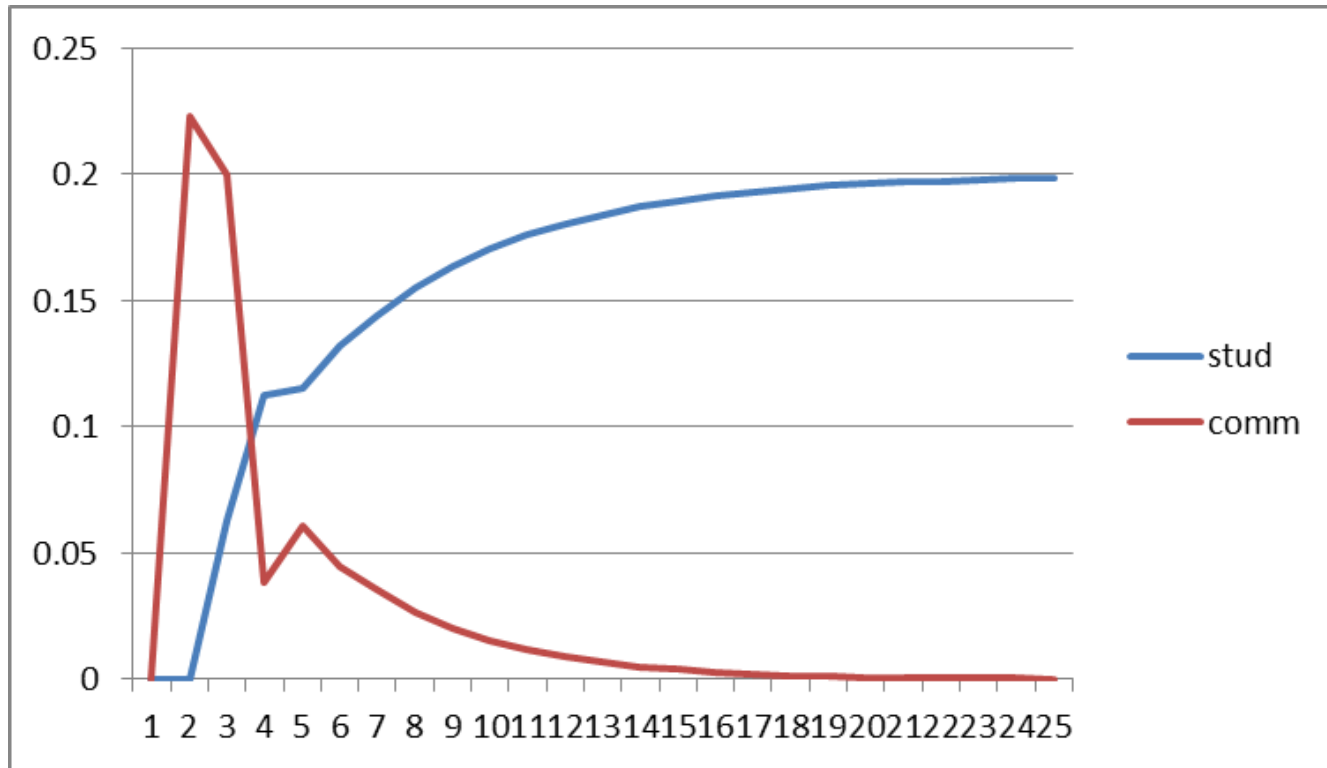
## Value of a superior bull

= Selection Difference \* Nr.Progeny \* expression per progeny



Progeny in commercial, so for a stud bulls  
these are actually  
grand progeny,  
great grand progeny, etc

# (allele) frequency of one unit of superiority as expressed in commercial herd



The fate of superiority from commercial bull vs a stud bull

Noting that a commercial bull also transmits the superiority from a stud bull ?!

# GENEFLOW

males to males	females to males
males to females	females to females

Donors of genes

		Sires of Nucleus					Dams of Nucleus									
P matrix		1	2	3	4	5	1	2	3	4	5	6	7	8	9	10
Recipients of genes	1	0	0.5	0	0	0	0	0.166667	0.166667	0.166667	0	0	0	0	0	0
	2	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	3	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
	4	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
	5	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0
	6	0	0.5	0	0	0	0	0.166667	0.166667	0.166667	0	0	0	0	0	0
	7	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0
	8	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0
	9	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0
	10	0	0	0	0	0	0	0	0	1	0	0	0	0	1	0

# GENEFLOW

Donors of genes

Recipients of genes

Sn<Sn	Sn<Dn	Sn<Sc	Sn<Cm	Sn<Cf	Sn	Sires of Nucleus		
Dn<Sn	Dn<Dn	Sf<Sc	Dn<Cm	Dn<Cf	Dn	Dams of Nucleus		
Sc<Sn	Sc<Dn	Sc<Sc	Sc<Cm	Sc<Cf	Sc	Stud born males to sire commercial		
Cm<Sn	Cm<Dn	Cm<Sc	Cm<Cm	Cm<Cf	Cm	Commercial born males		
Cf<Sn	Cf<Dn	Cf<Sc	Cf<Cm	Cf<Cf	Cf	Commercial born females		

## Donors of genes

[illegible]

P = matrix describing transmission of genes



# GENEFLOW

- R = a matrix defining gene transmission of some superiority (or particular allele)

- Q= a matrix describing aging

		Sires of Nucleus					Dams of Nucleus			
P matrix		1	2	3	4	5	1	2	3	4
	1	0	0.5	0	0	0	0	0.166667	0.166667	0.166667
	2	1	0	0	0	0	0	0	0	0
	3	0	1	0	0	0	0	0	0	0
	4	0	0	1	0	0	0	0	0	0
	5	0	0	0	1	0	0	0	0	0
	1	0	0	0	0.5	0	0	0.166667	0.166667	0.166667

- P = matrix describing transmission of genes
  - $P=R+Q$

$$m_t = P m_{t-1} + R n_{t-1}$$

- m vector of allele frequency in each age class
- n vector to describe inserting allele or superiority

# GENEFLOW

g1	g2	g3	g4	g5	g6	g7	g8	g9	g10	g11	g12	g13	g14	g15	g16	g17	g18	g19	g20	g21	g22	g23	g24	g25	
0	0.5	0	0.333333	0.083333	0.305556	0.111111	0.273148	0.138889	0.251543	0.156636	0.236368	0.169496	0.225609	0.178498	0.21805	0.184849	0.212718	0.189324	0.208962	0.192477	0.206315	0.194699	0.20445	0.196265	
1	0	0.5	0	0.333333	0.083333	0.305556	0.111111	0.273148	0.138889	0.251543	0.156636	0.236368	0.169496	0.225609	0.178498	0.21805	0.184849	0.212718	0.189324	0.208962	0.192477	0.206315	0.194699	0.20445	
0	1	0	0.5	0	0.333333	0.083333	0.305556	0.111111	0.273148	0.138889	0.251543	0.156636	0.236368	0.169496	0.225609	0.178498	0.21805	0.184849	0.212718	0.189324	0.208962	0.192477	0.206315	0.194699	
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0	0	0	0	0	0	0	0.25	0	0.208333	0.229167	0.104167	0.22338	0.138889	0.218557	0.156057	0.214989	0.170332	0.211072	0.179479	0.208424	0.185905	0.190297	0.204567	0.193304	0.195378
0	0	0	0	0	0.25	0	0.208333	0.229167	0.104167	0.22338	0.138889	0.218557	0.156057	0.214989	0.170332	0.211072	0.179479	0.208424	0.185905	0.190297	0.204567	0.193304	0.203333	0.203333	
0	0	0	0	0	0	0.25	0	0.208333	0.229167	0.104167	0.22338	0.138889	0.218557	0.156057	0.214989	0.170332	0.211072	0.179479	0.208424	0.185905	0.190297	0.204567	0.193304	0.203333	
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0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.25	0	0.208333	0.229167	0.104167	0.22338	

Allele frequency in the limit, from on 'insertion' of superiority (or an allele) =  $1/(L_m + L_f)$

Geneflow mainly useful for initial part of an action, otherwise can use Rendel and Robertson

# Cumulative Discounted Expressions CDE

Value (V) in year t is worth now  $V.c$  where  $c=1/(1+d)^t$

d = discount rate

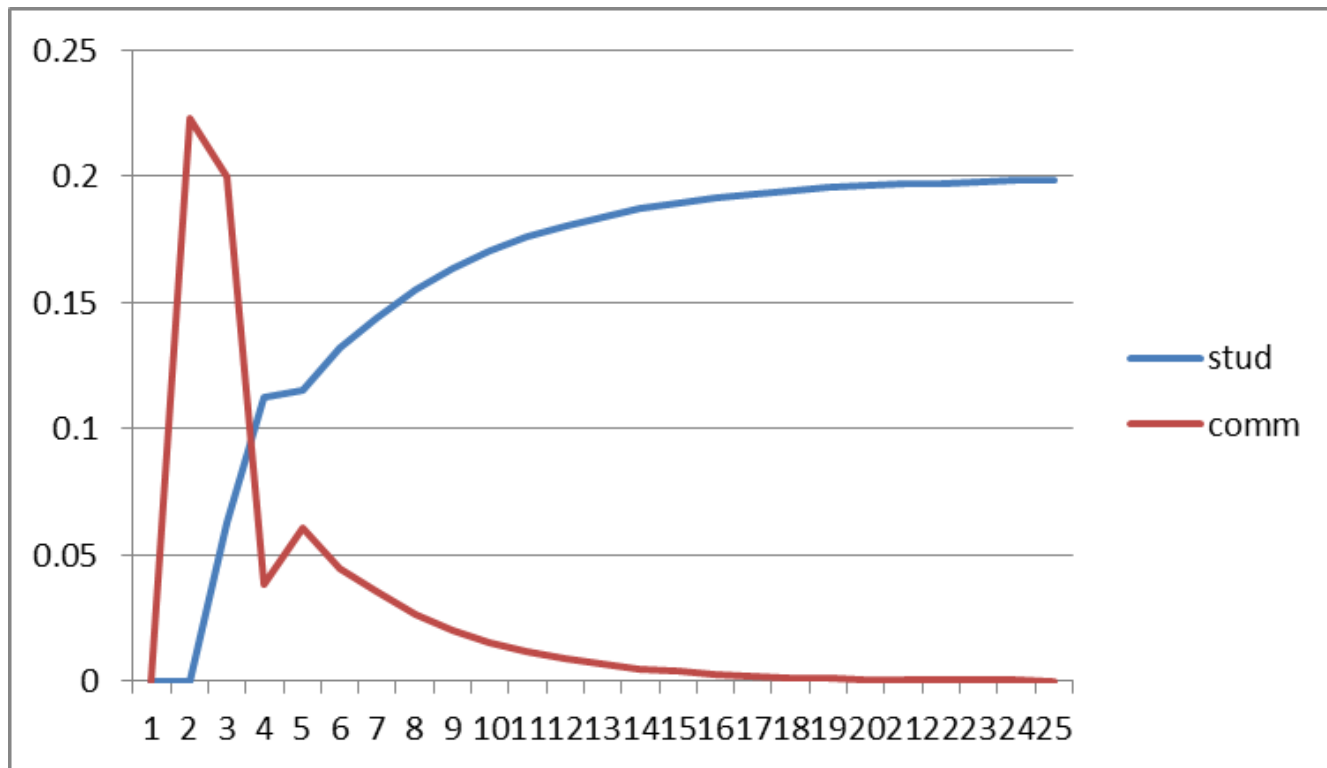
c = discount factor

Expression in age class i in year t is  $m(i)_t = E_{it}$

Net Present Value of Sum of expression over 25 years

$$CDE = \sum_{t=1}^{25} \sum_{i=1}^{nac} E_{it} c_t$$

# (allele) frequency of one unit of superiority as expressed in commercial herd



Discount rate

0

0.05

0.08

CDE comm bulls

0.99

0.78

0.68

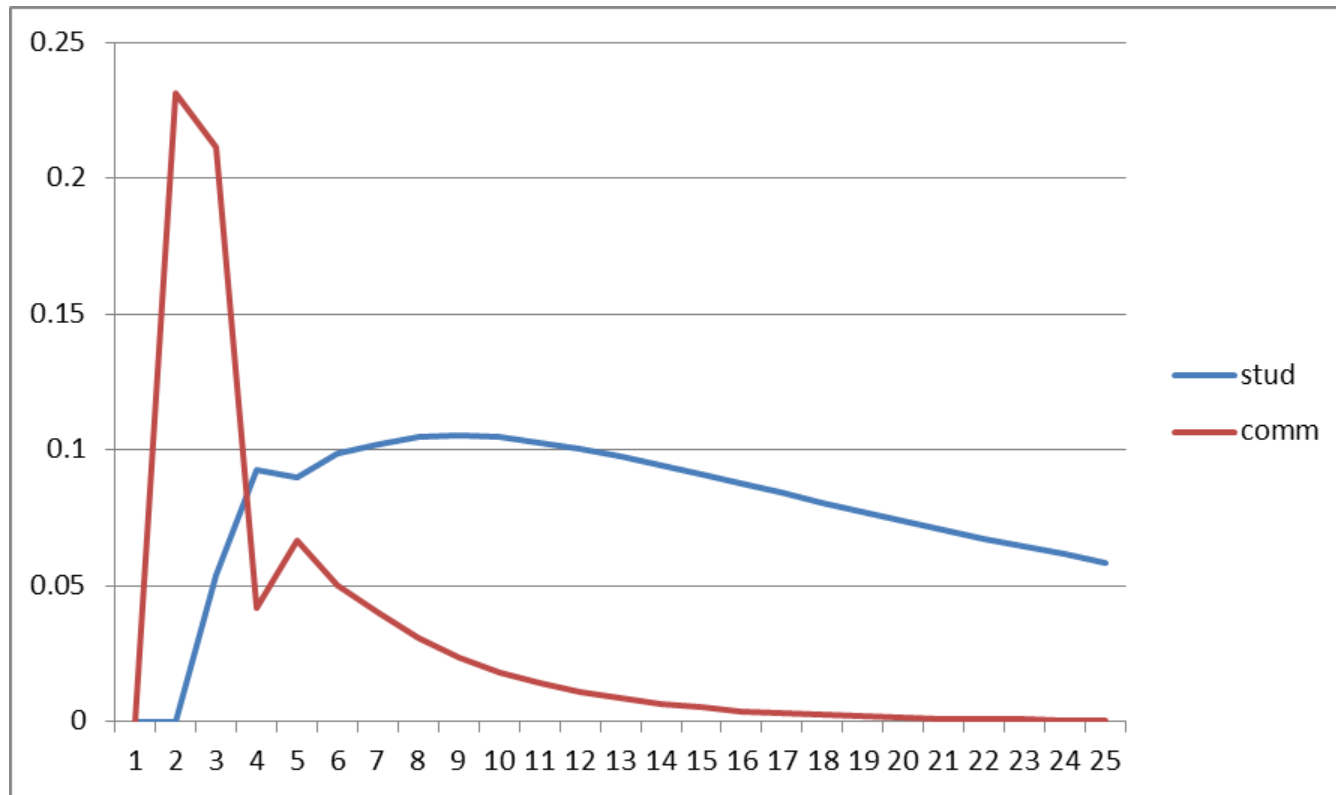
CDE stud sires

3.93

1.96

1.37

# (allele) frequency of one unit of superiority as expressed in commercial herd



Discount rate	CDE comm bulls	CDE stud sires
0	0.99	3.93
0.05	0.78	1.96
0.08	0.68	1.37

# Value of selecting Stud Sires and Comm bulls

## Value of a superior bull

$$= \text{Selection Difference} * \text{Nr.Progeny} * \text{expression per progeny}$$

CDE

Comm bull	+ 1.4	100	0.55	= \$ 77
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Stud Sire	+ 3.0	400	1.35	= \$ 1,620
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# Herd structure

	Nr Cows Commercial Herd	12,000	
	Comm Dams/sire	50	
	Comm Sire replacem. rate	0.33333	
	Comm Weaning rate	1	
Nr new rams needed for comm herd/yr		80	
Nr lifetime Progeny per commercial sire			150
Prop. Stud.Males sold as breeding bull		20%	
	Stud weaning rate	1	
	Stud dams/sire	40	
	Nr stud breeding cows	800	
	Nr. Of stud sires	20	
	Nr of comm bulls sold per year	80	
Nr of commercial bulls sold per Stud male		4	
Progeny receiving genes from a stud male			600

150 prog/comm bull

600prog/stud bull

# Value of selecting Stud Sires and Comm Bulls

Value of a superior sire = Selection Difference \* Nr.Progeny \* CDE

- Selection differential within the cohort: “The result of one round of selection”

Breeding performance					
		SD of breeding Objective	10.82		
		Male Selection intensity	2.06		
		Female Selection intensity	0.2		
		Male Selection accuracy <i>without</i> genomics	0.358	increase	
		Male Selection accuracy <i>with</i> genomics	0.432	21%	
		Female Selection accuracy	0.358		
		Generation Interval Stud males	1.53		
		Generation Interval stud females	2.97		
	approximateley	1.90	CDE stud sires	1.90	
			CDE flock sires	0.6	
			no GS	GS	
			Sire superiority	7.979534	9.628934
			Dam Superiority	0.774712	0.774712
			Rate of gain/year	1.945	2.312
					19%



# Comparing geneflow with dG/year method

group	int	acc		Sup	L		dG/year
sires	2.1543	0.53		10.27622255	1.0		3.233294535
dams	0.7979	0.37		2.656955587	3.0		

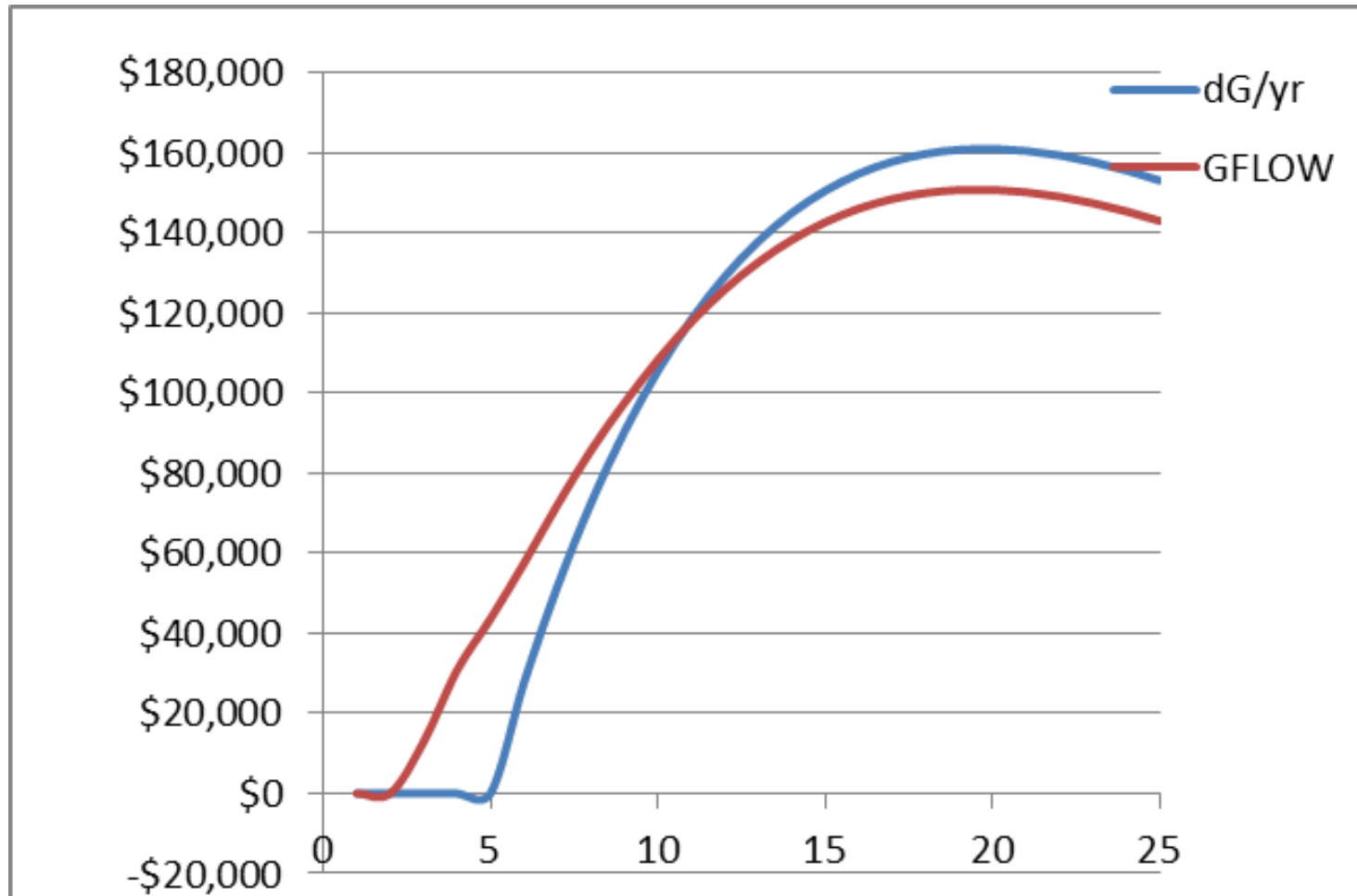
Calculations based on dG/year						calculations based on GFLOW				
		dG/gen					sire selection	dam selection		
		3.23			dG/yr	superiority	10.2762	2.6570		GFLOW
year	disc fact	genetic mean	cum benefit	cost	disc retruns		Expr_SS	Expr_DS	cum benefit	disc retruns
1	1.000	0	\$ -	\$0	\$0		0.000	0.000	\$ -	\$ -
2	0.935	0	\$ -	\$0	\$0		0.000	0.000	\$ -	\$ -
3	0.873	0	\$ -	\$0	\$0		0.119	0.000	\$ 14,694	\$ 12,834
4	0.816	0	\$ -	\$0	\$0		0.174	0.048	\$ 37,679	\$ 30,757
5	0.763	0	\$ -	\$0	\$0		0.131	0.105	\$ 57,158	\$ 43,606
6	0.713	\$3.23	\$ 38,800	\$0	\$27,664		0.157	0.128	\$ 80,610	\$ 57,474
7	0.666	\$6.47	\$ 77,599	\$0	\$51,708		0.185	0.149	\$ 108,155	\$ 72,068
8	0.623	\$9.70	\$ 116,399	\$0	\$72,487		0.196	0.165	\$ 137,537	\$ 85,651
9	0.582	\$12.93	\$ 155,198	\$0	\$90,327		0.197	0.178	\$ 167,514	\$ 97,494
10	0.544	\$16.17	\$ 193,998	\$0	\$105,522		0.206	0.190	\$ 198,976	\$ 108,230
11	0.508	\$19.40	\$ 232,797	\$0	\$118,342		0.213	0.199	\$ 231,559	\$ 117,713
12	0.475	\$22.63	\$ 271,597	\$0	\$129,034		0.217	0.206	\$ 264,833	\$ 125,820
13	0.444	\$25.87	\$ 310,396	\$0	\$137,820		0.220	0.211	\$ 298,645	\$ 132,602
14	0.415	\$29.10	\$ 349,196	\$0	\$144,904		0.223	0.216	\$ 332,996	\$ 138,182
15	0.388	\$32.33	\$ 387,995	\$0	\$150,471		0.225	0.220	\$ 367,735	\$ 142,614
16	0.362	\$35.57	\$ 426,795	\$0	\$154,690		0.227	0.222	\$ 402,772	\$ 145,983
17	0.339	\$38.80	\$ 465,594	\$0	\$157,713		0.228	0.225	\$ 438,053	\$ 148,384
18	0.317	\$42.03	\$ 504,394	\$0	\$159,678		0.229	0.226	\$ 473,539	\$ 149,910
19	0.296	\$45.27	\$ 543,193	\$0	\$160,711		0.230	0.228	\$ 509,179	\$ 150,648
20	0.277	\$48.50	\$ 581,993	\$0	\$160,926		0.231	0.229	\$ 544,943	\$ 150,681
21	0.258	\$51.73	\$ 620,793	\$0	\$160,425		0.231	0.230	\$ 580,808	\$ 150,092
22	0.242	\$54.97	\$ 659,592	\$0	\$159,300		0.232	0.231	\$ 616,754	\$ 148,954
23	0.226	\$58.20	\$ 698,392	\$0	\$157,636		0.232	0.231	\$ 652,763	\$ 147,337
24	0.211	\$61.43	\$ 737,191	\$0	\$155,508		0.232	0.232	\$ 688,823	\$ 145,305
25	0.197	\$64.67	\$ 775,991	\$0	\$152,984		0.233	0.232	\$ 724,923	\$ 142,916
				NPV	\$2,607,849				NPV	\$ 2,645,255

Expressed in  
12,000 cows

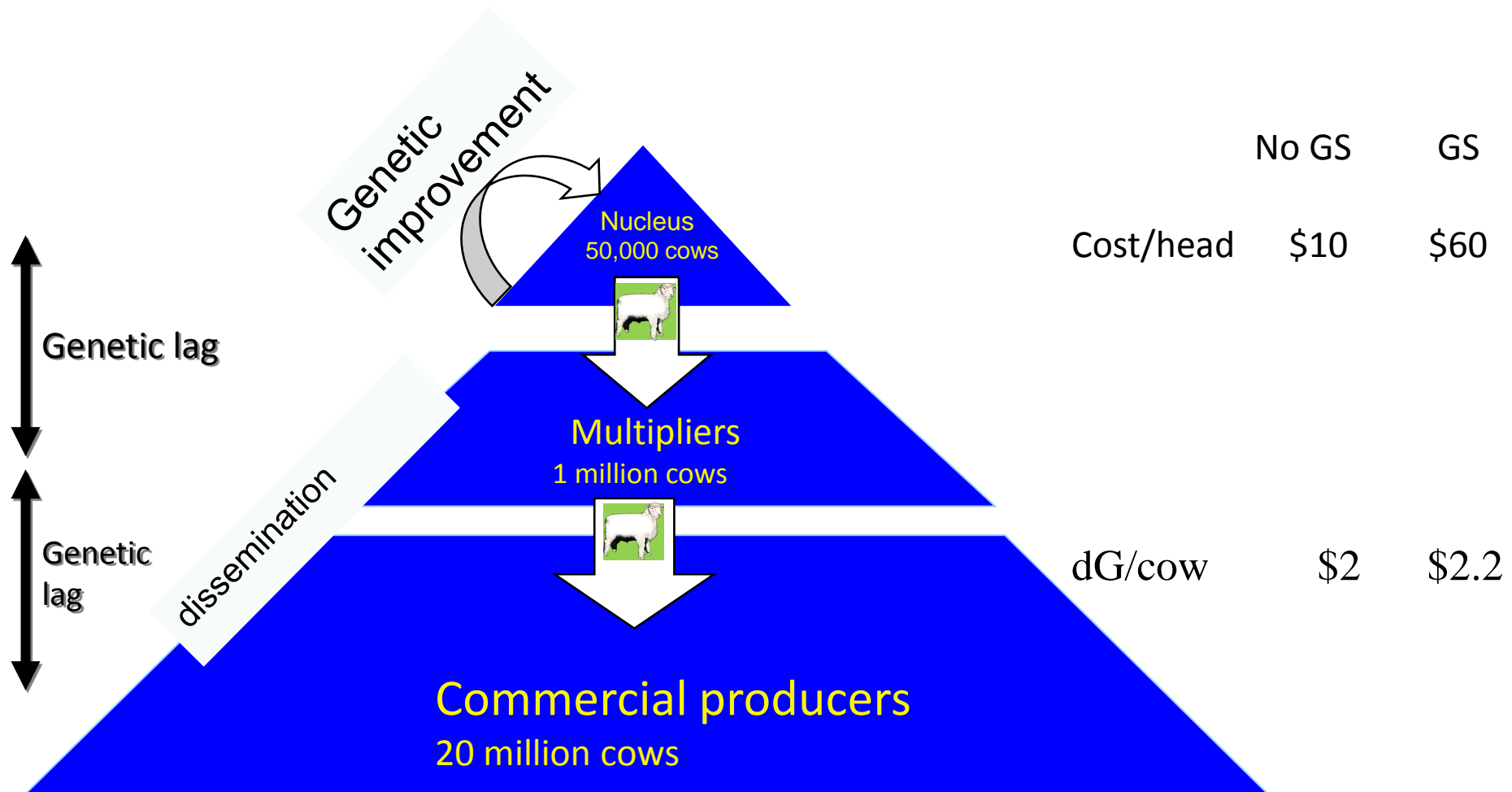
20 nucleus sires

i.e. 600 per sire

## Comparing simply dG/yr vs GFLOW

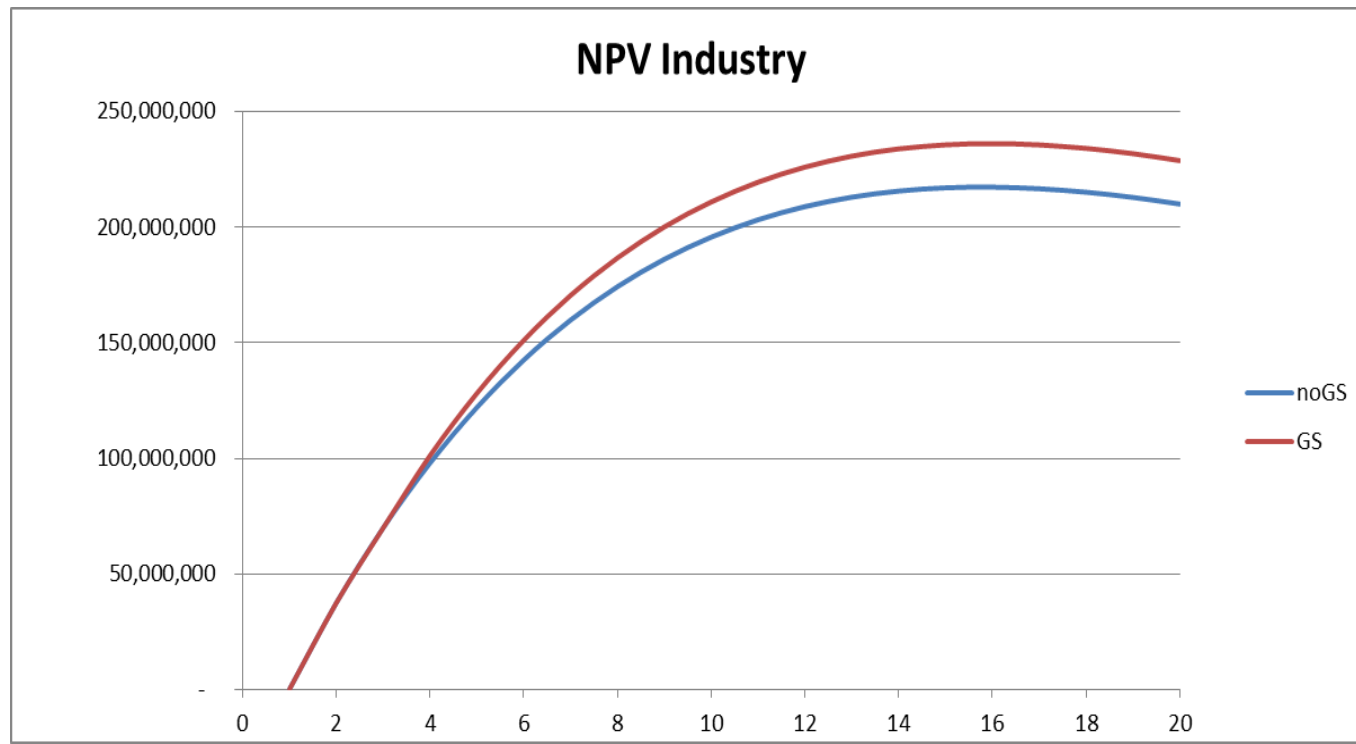


# Cost - Benefit of breeding programs



# Cost-Benefit industry wide

	<u>No GS</u>	<u>GS</u>
Cost	\$0.5 M	\$ 1.65 M
dG	\$40 M	\$ 44 M



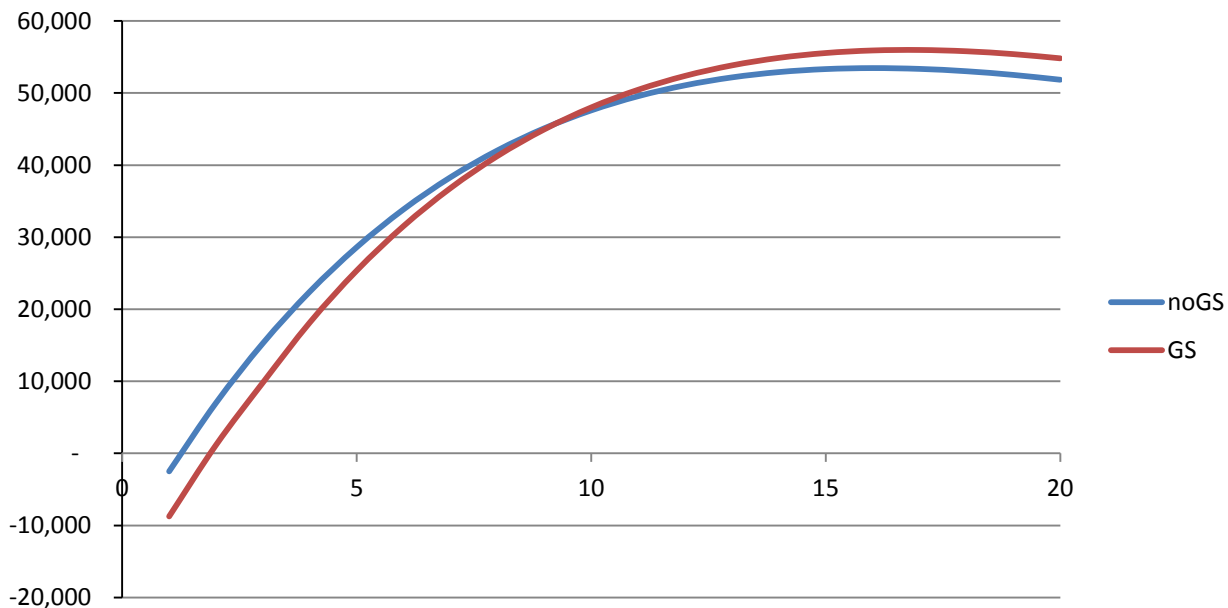
50k Nuc cows  
20M Comm

3 tier benefit

# Cost-Benefit Stud

	No GS	GS
Cost	\$ 5 k	\$17.5 k
dG	\$20 k	\$ 22 k

**NPV Stud**



500 Nuc cows  
10k Comm

2 tier benefit

# Value of selecting Stud Sires and Comm Bulls

## Value of a superior sire

= Selection Difference \* Nr.Progeny \* expression per progeny

Comm Bull	+ 1.4	100	0.55	= \$ 77
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With Genomics

+1.6			= \$ 88	+11
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Stud Sire	+ 3.0	400	1.35	= \$ 1,620
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With Genomics

+3.4			= \$ 1,836	+216
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## Cost benefit analysis

- Extra benefit  $120 * \$11 + 30 * \$216 = \$ 7,800$
- If all young stud males tested: 600
- Break even: \$13.00 per DNA test

## summary

- Can calculate additional gain on a per bull basis, assuming returns in commercial progeny
- Those figures depend on
  - Additional accuracy
  - Age structure
  - Herd parameters such as weaning rate, mating rate, prop. Sold