

Optimizing Breeding Programs

COST-BENEFIT of genomics



Cost - Benefit of breeding programs



Cost-Benefit industry wide



Cost-Benefit Stud + Direct clients



Cost-Benefit Stud + Direct clients



Value of selecting Stud Rams and Flock Rams



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



Discount rate	CDE flock rams	CDE stud rams
0	0.99	3.93
0.05	0.78	1.96
0.08	0.68	1.37

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		
	Nr. Of stud sires	20	
	Nr of comm bulls sold per year	80	
	Proportion of males DNA tested	100%	
Nr. Stud born Male DNA tested/yr		400	
Nr of commercial bulls sold per Stud male		4	
Nr of DNA tested young male per stud bull		20	
Ir of commercial progeny receiving genes from a stud male			600

100 prog/comm bull

400 prog/stud sire

Value of selecting Stud Rams

Value of a superior sire = Selection Difference * Nr.Progeny * expression per progeny

Stud sire

	+ 8.8	400	1.36	= \$7,194
With				
Genomics	+ 9.8	400	1.36	= \$ 8,011

+817

Cost benefit analysis

• Extra benefit per stud sire

\$817

- Extra Cost If all young stud males tested: 20 young males/stud sire
- Break even: \$817 / 20 = \$41 per DNA test

Breakeven cost and proportion genotyped (no loss assumed!)



1yo male, 2yo female, Fine10% + SS, 40% males sold as rams

2 stage selection

How many bulls to genotype?

All have a breeding value at stage 1 EBV Only some get extra info from GBV GEBV

Important parameters:

EBV accuracy

Correlation EBV GEBV

added accuracy GBV \rightarrow GEBV

% gain compared with 100% genotyping EBV 0.34, GBV 0.39, GEBV = 0.50, **r** = **0.7**



At high(ish) correlation between EBV and GEBV only need to genotype ~20%

% gain compared with 100% genotyping EBV 0.10, GBV = 0.39, GEBV 0.40, **r** = **0.25**



Proportion genotyped

At low(er) correlation between ASBV and ASBV1 need to genotype more

summary

- Can calculate additional gain on a per ram basis, assuming returns in commercial progeny
- Those figures depend on
 - Additional accuracy
 - Age structure
 - Flock parameters such as weaning rate, mating rate, proportion sold
 - Can have strategies to save costs, e.g. test top 20%