

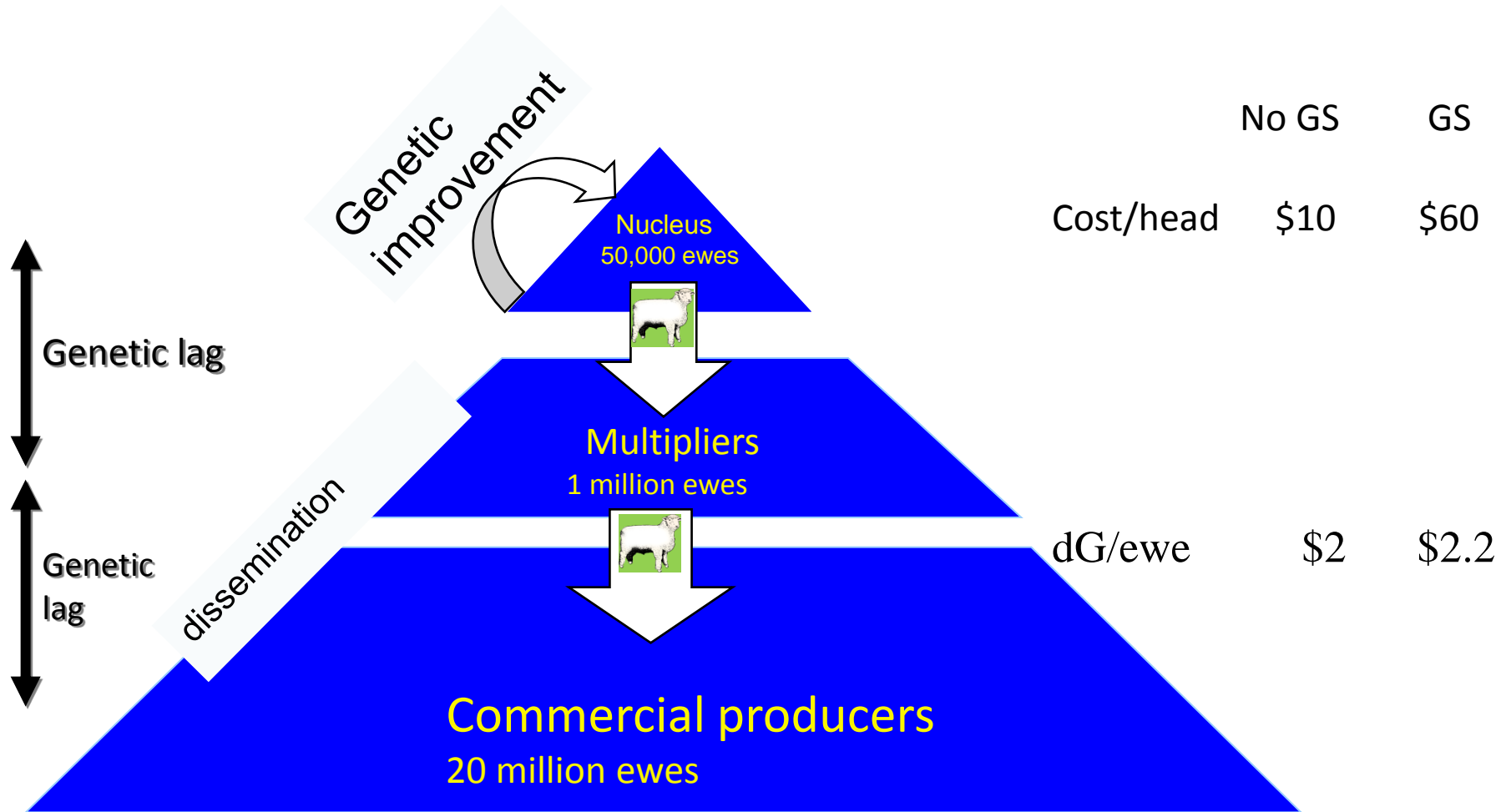
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University of  
New England

# Optimizing Breeding Programs

## COST-BENEFIT of genomics

# Cost - Benefit of breeding programs



# Cost-Benefit industry wide

3 tier benefit

Per annum added

This does not accumulate ←

This accumulates ←

Cost

dG Return

Cum NPV

No GS

GS

\$0.5 M

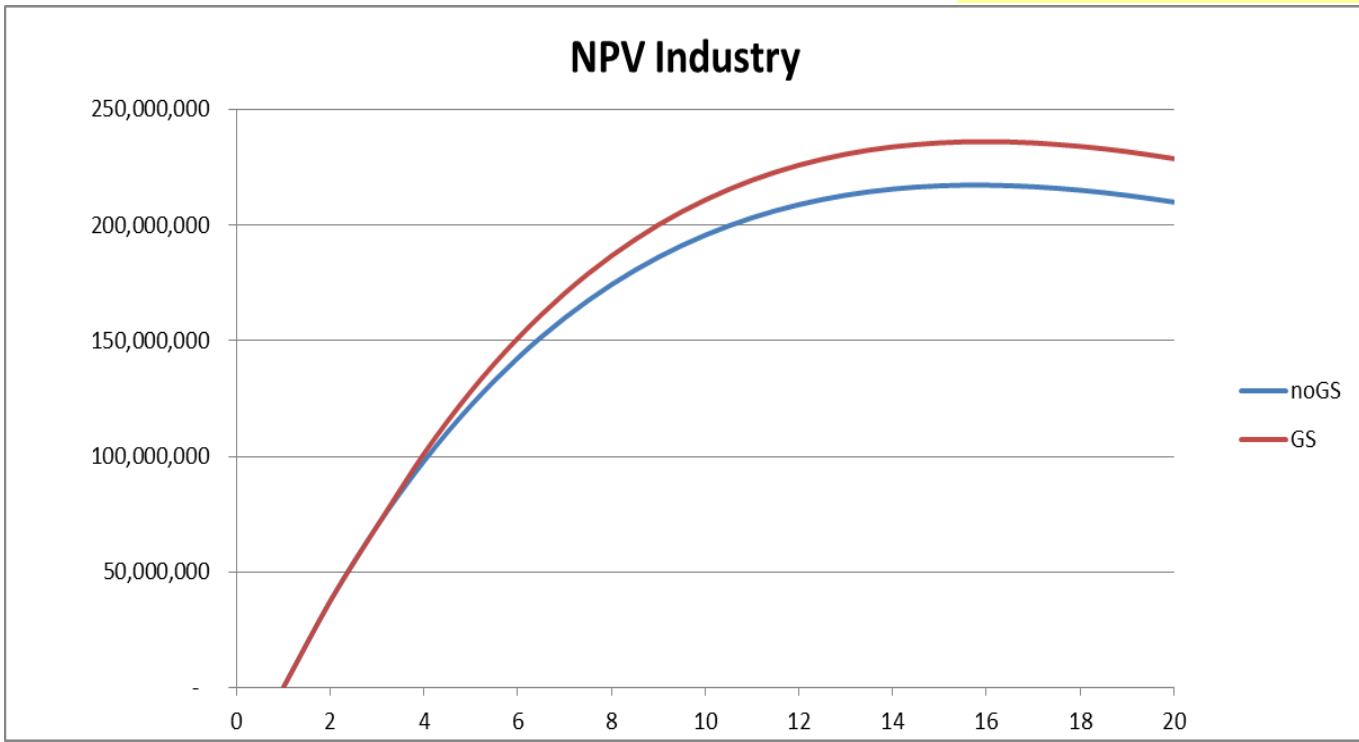
\$ 1.65 M

\$40 M

\$ 44 M

\$2.45 b

\$2.65 b



Note the much higher cost and lower RTI ratio!

50k Nuc ewes  
20M Comm

# Cost-Benefit Stud + Direct clients

2 tier benefit

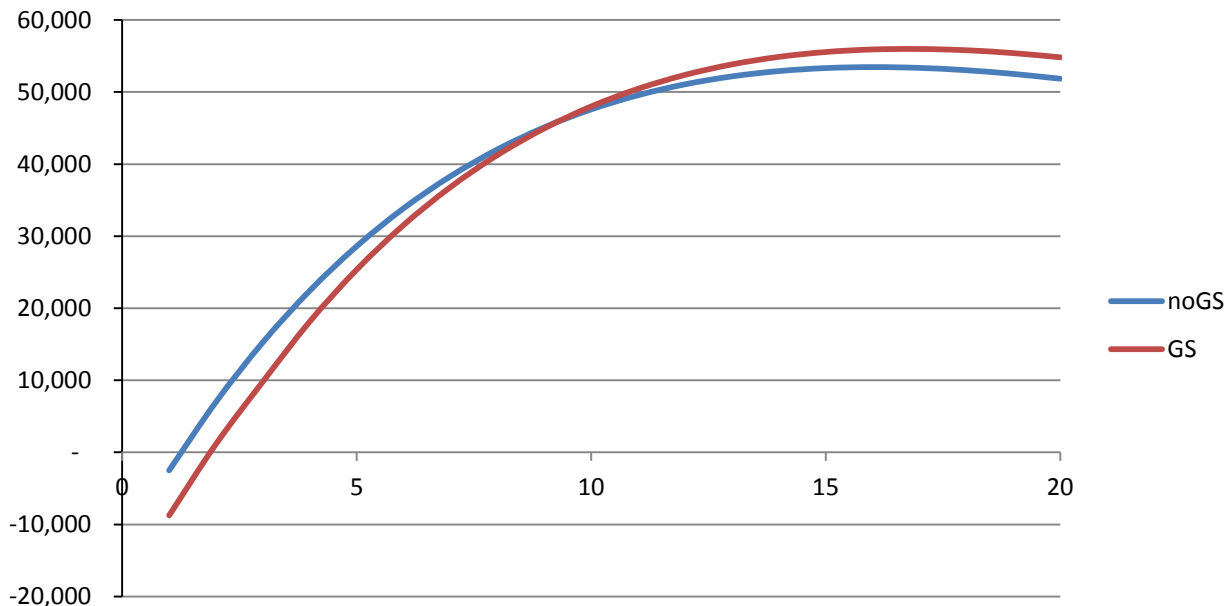
Per annum added

50-50 split of benefit  
between stud and comm

Cost  
Returns dG  
Cum NPV

No GS	GS
\$ 5 k	\$17.5 k
\$20 k	\$ 22 k
\$807 k	\$859 k

NPV Stud



500 Nuc ewes  
10k Comm

# Cost-Benefit Stud + Direct clients

2 tier benefit

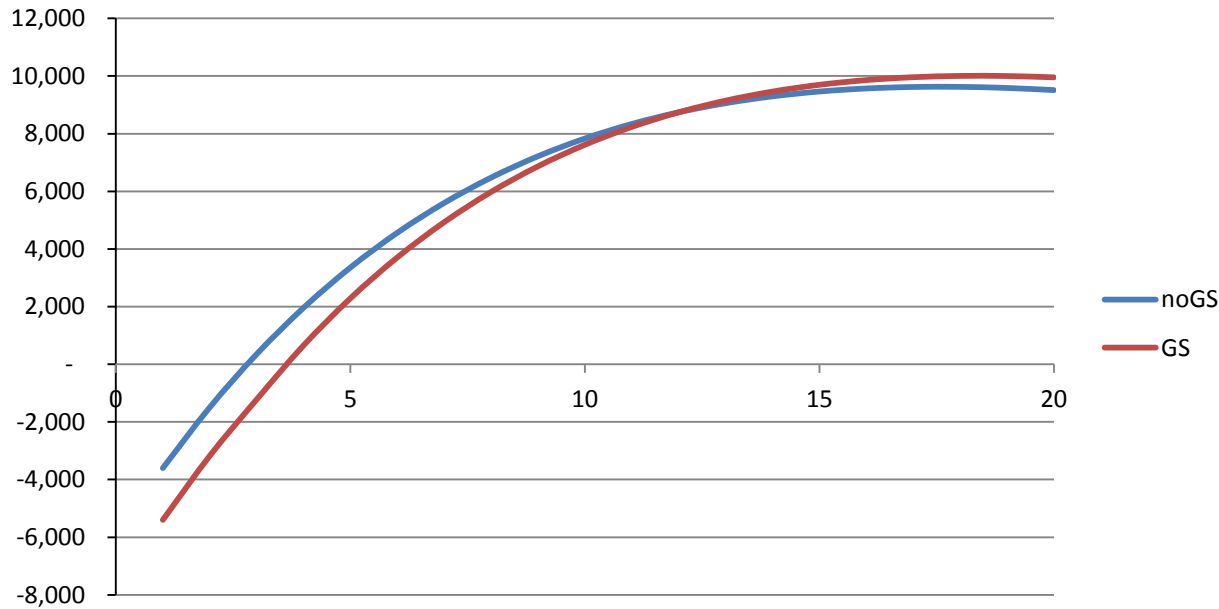
Per annum added

90 -10 split of benefit  
between stud and comm

Cost  
Returns dG  
Cum NPV

No GS	GS
\$ 5 k	\$17.5 k
\$20 k	\$ 22 k
\$125 k	\$117 k

NPV Stud



500 Nuc ewes  
10k Comm

# Value of selecting Stud Rams and Flock Rams

Value of a superior ram

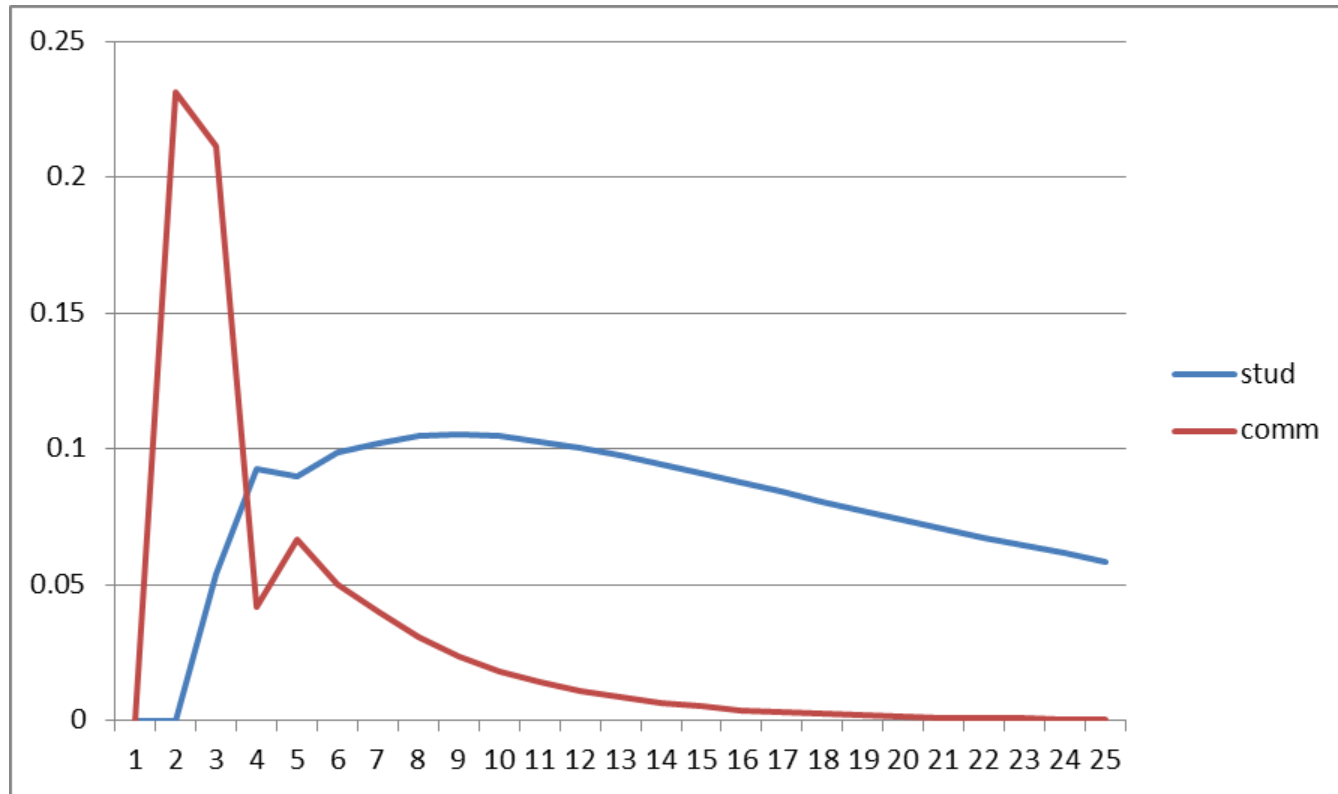
$$= \text{Selection Difference} * \text{Nr.Progeny} * \text{expression per progeny}$$

CDE

Flock Ram	+ 1.4	100	0.55	= \$ 77
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Stud Ram	+ 3.0	400	1.35	= \$ 1,620
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# (allele) frequency of one unit of superiority as expressed in commercial flock



<u>Discount rate</u>	<u>CDE flock rams</u>	<u>CDE stud rams</u>
0	0.99	3.93
0.05	0.78	1.96
0.08	0.68	1.37

# Flock structure

ck	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

100 prog/flock ram

	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	
	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	
	Nr of commercial progeny receiving genes from a stud male		600

400 prog/stud ram



## Some real data

Commerical Flock	Nr Sheep Commercial Flock	34,280	
	Comm Dams/sire	40	
	Comm Sire replacem. rate	0.33333	
	Comm Weaning rate	1.1	
	Nr new rams needed for comm flock/yr	286	
	Nr lifetime Progeny per commercial sire		132

Stud Flock	Prop. Stud.Males sold as breeding ram	40%	
	Stud weaning rate	1.28	
	Stud dams/sire	20	
	Nr stud breeding ewes	1116	
	Nr. Of stud sires	56	
	Nr of flock rams sold per year	286	
	Nr of commercial rams sold per Stud male	5.12	
	Nr of commercial progeny receiving genes from a stud male		676

# Value of selecting Stud Rams and Flock Rams

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

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

Breeding performance		Selection differential within the cohort: “The result of one round of selection”				
		SD of breeding Objective	10			
		Male Selection intensity	2			
		Female Selection intensity	0.5			
		Male Selection accuracy <i>without</i> genomics	0.44	increase		
		Male Selection accuracy <i>with</i> genomics	0.49	11%		
		Female Selection accuracy	0.4			
		Generation Interval Stud males	2			
		Generation Interval stud females	4			
	approximalely	1.36	CDE stud sires	1.36		
			CDE flock sires	0.6		
			no GS	GS		
			Sire superiority	8.8	9.8	
			Dam Superiority	2.2	2	increase
			Rate of gain/year	1.8	2.0	7%
			Break Even Cost of DNA test	41		
					value diff	
			Value of stud bull	7194	8011	817
			Value of comm bull	261	291	41

# Value of selecting Stud Rams and Flock Rams

## Value of a superior ram

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

### Stud Ram

+ 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 DNA test (\$) depends on breeding program

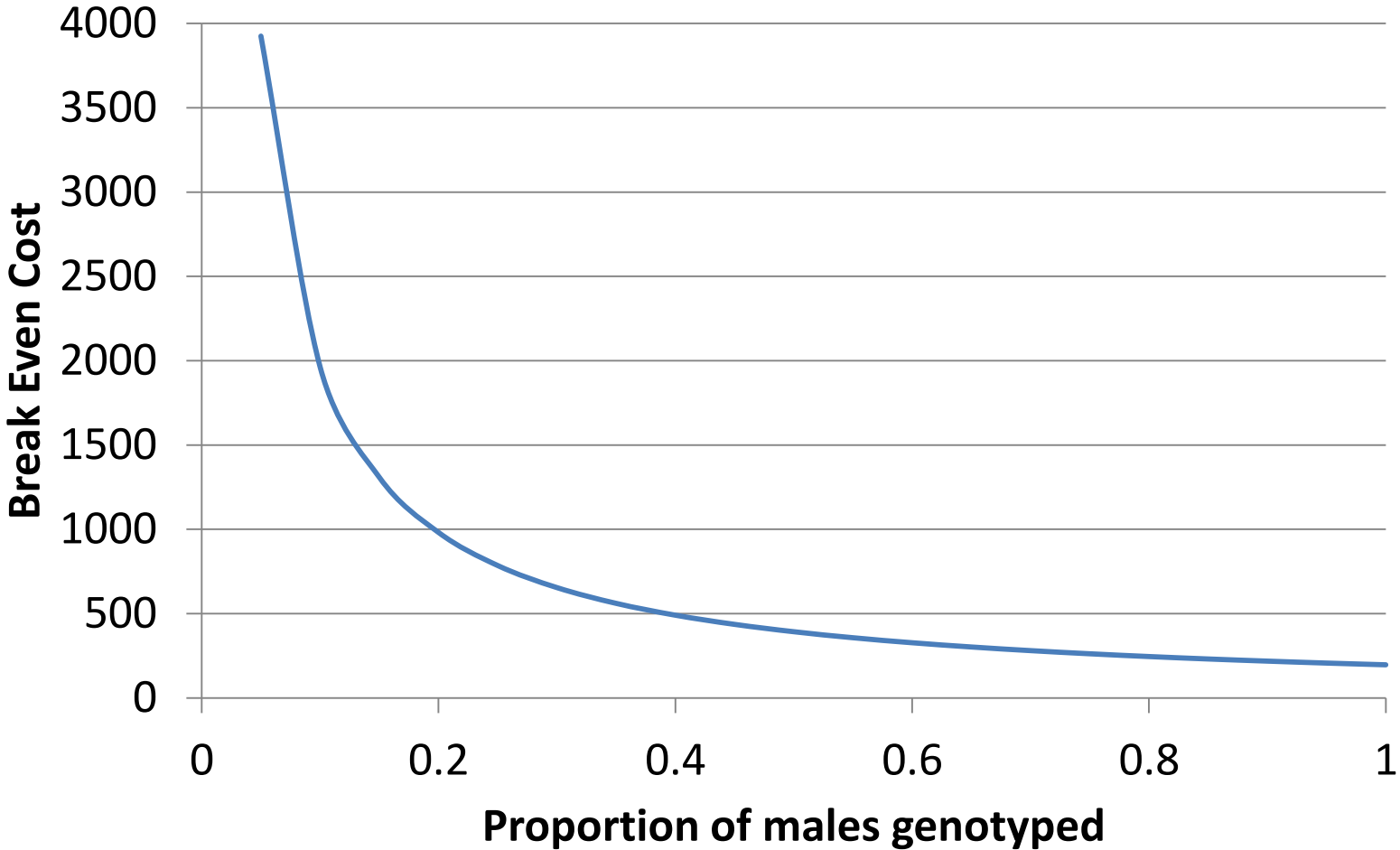
<b>Proportion tested</b>	<b>100%</b>		<b>20%</b>	
<b>Age at first progeny</b>	1yo	2yo	1yo	2yo
<b>Breakeven (\$/test)</b>	196	83	981	415

- assumes 40% males sold as rams

<b>% males born sold as rams</b>	<b>40%</b>		<b>20%</b>	
<b>Age at first progeny</b>	1yo	2yo	1yo	2yo
<b>Breakeven (\$/test)</b>	196	83	98	41

- assumes 100% of males tested

Breakeven cost and proportion genotyped (no loss assumed!)



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

# What increase in price received per ram sold do you need to cover costs?

<b>Total ram lambs weaned</b>	<b>200</b>	<b>200</b>	<b>200</b>	<b>200</b>
% DNA tested	20	100	20	100
nr tested (for use in nucleus)	40	200	40	200
Total test cost	2,000	10,000	2,000	10,000
% sold as flock rams	20	20	40	40
nr rams sold	40	40	80	80
Cost of test per ram sold	\$50	\$250	\$25	\$125

## 2 stage selection

How many rams to genotype?

All have a breeding value at stage 1      ASBV0

Only some get extra info      ASBV1

Important parameters:

ASBV0 accuracy

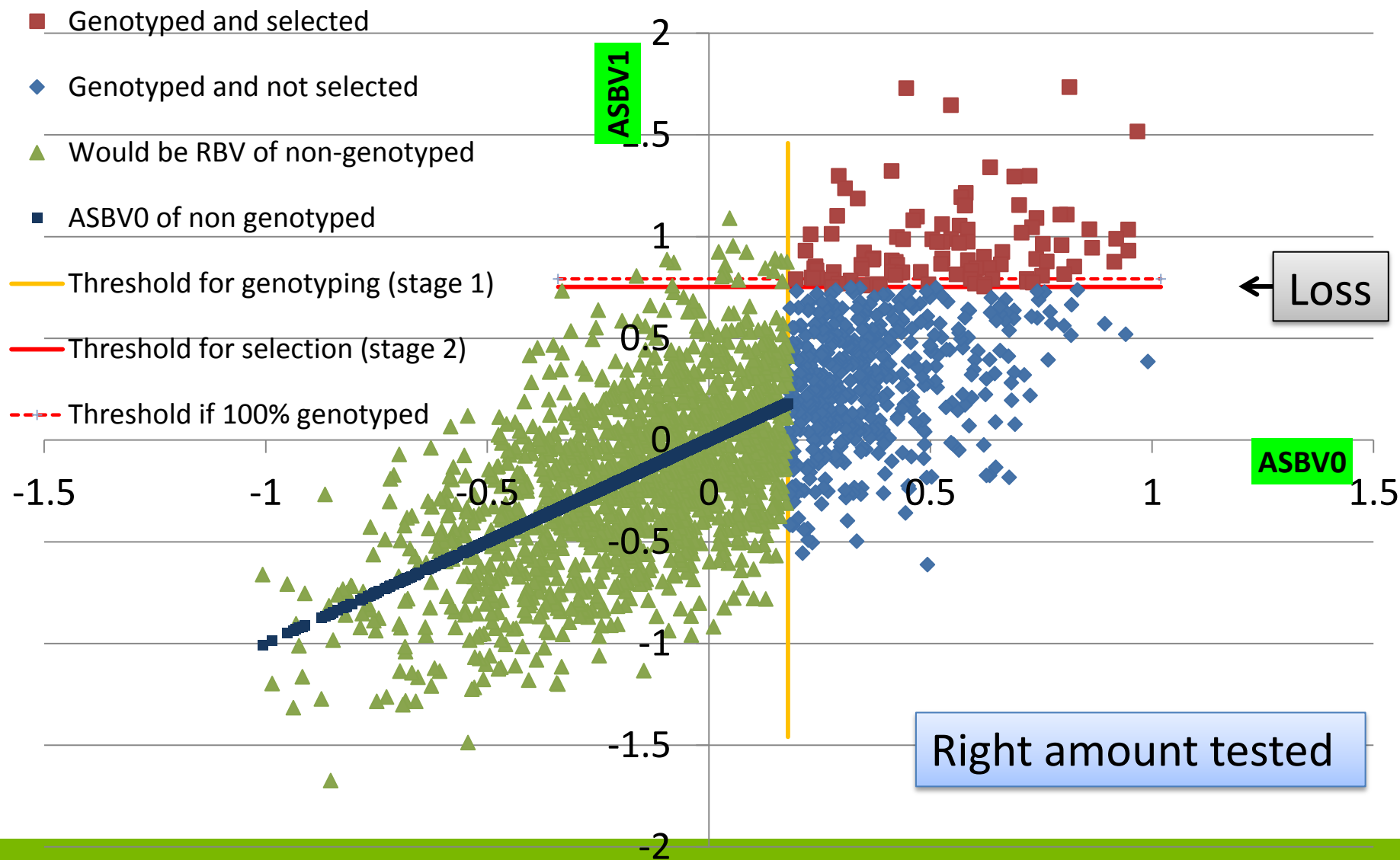


added accuracy ASBV1 → ASBV2

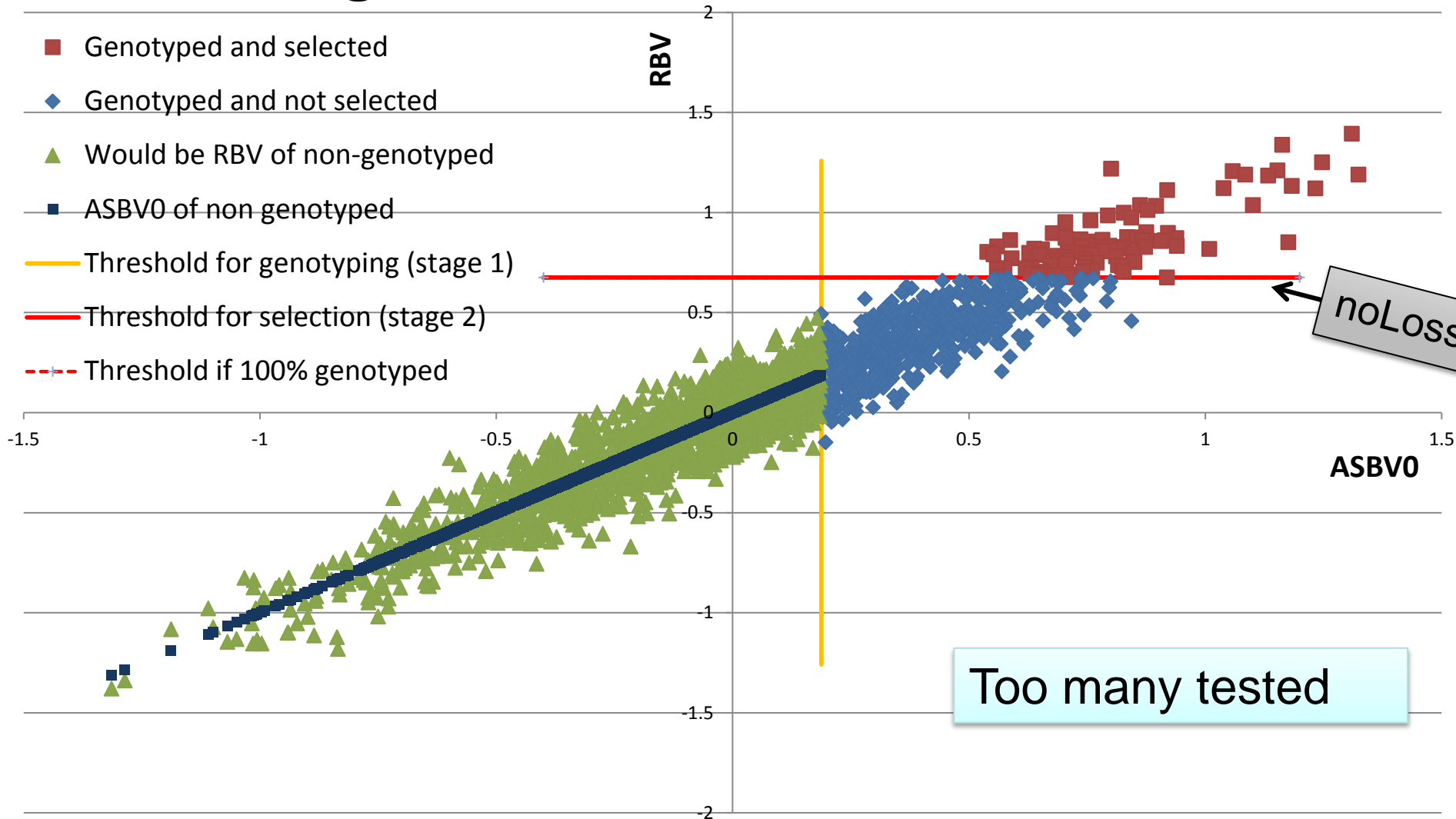
Correlation ASBV0, ASBV2



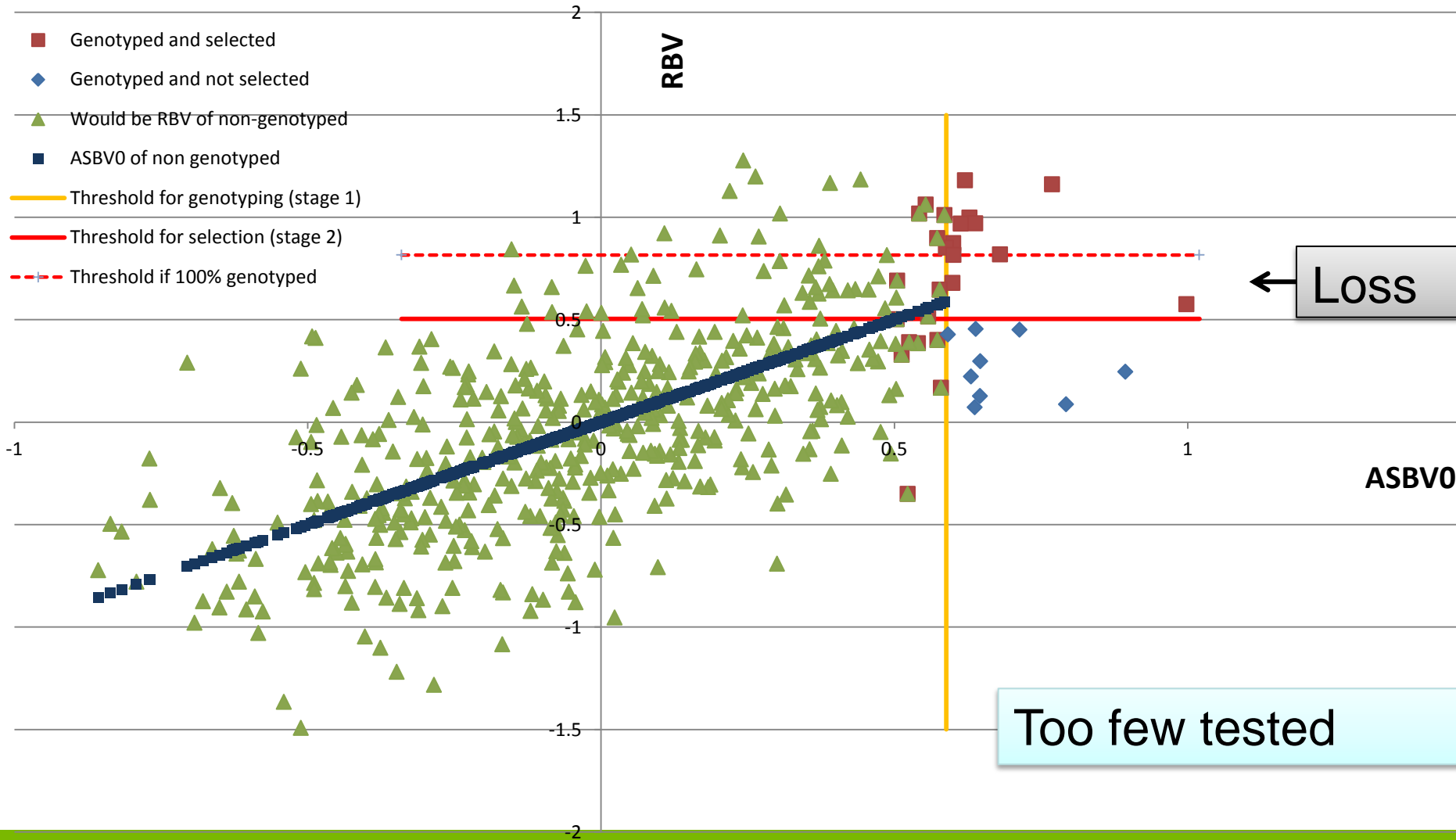
# 30% genotyped, 5% selected, correlation ASBV to RBV of 0.7



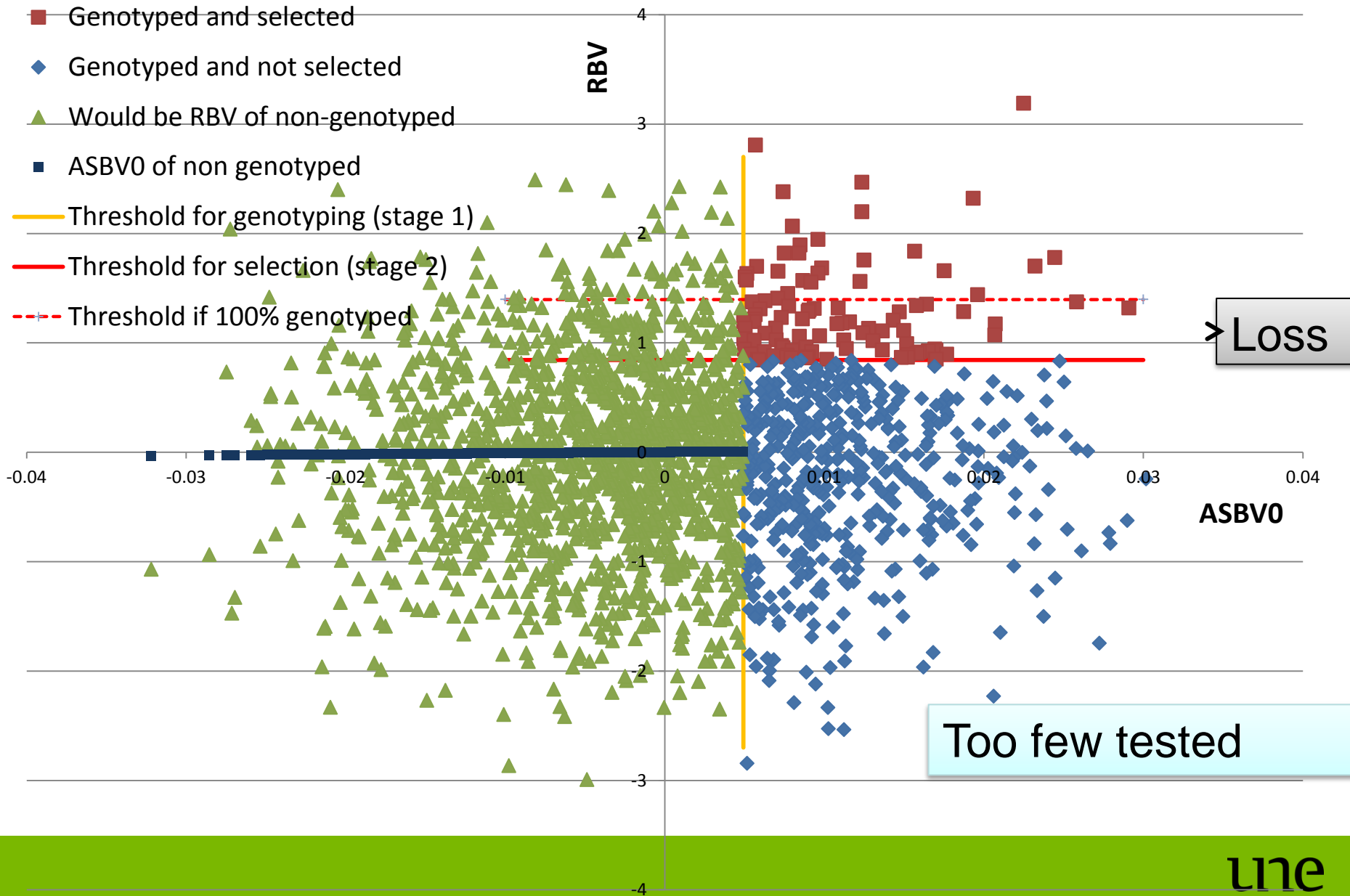
# 30% genotyped, 5% selected with very high correlation ASBV to RBV



# Very low proportion tested



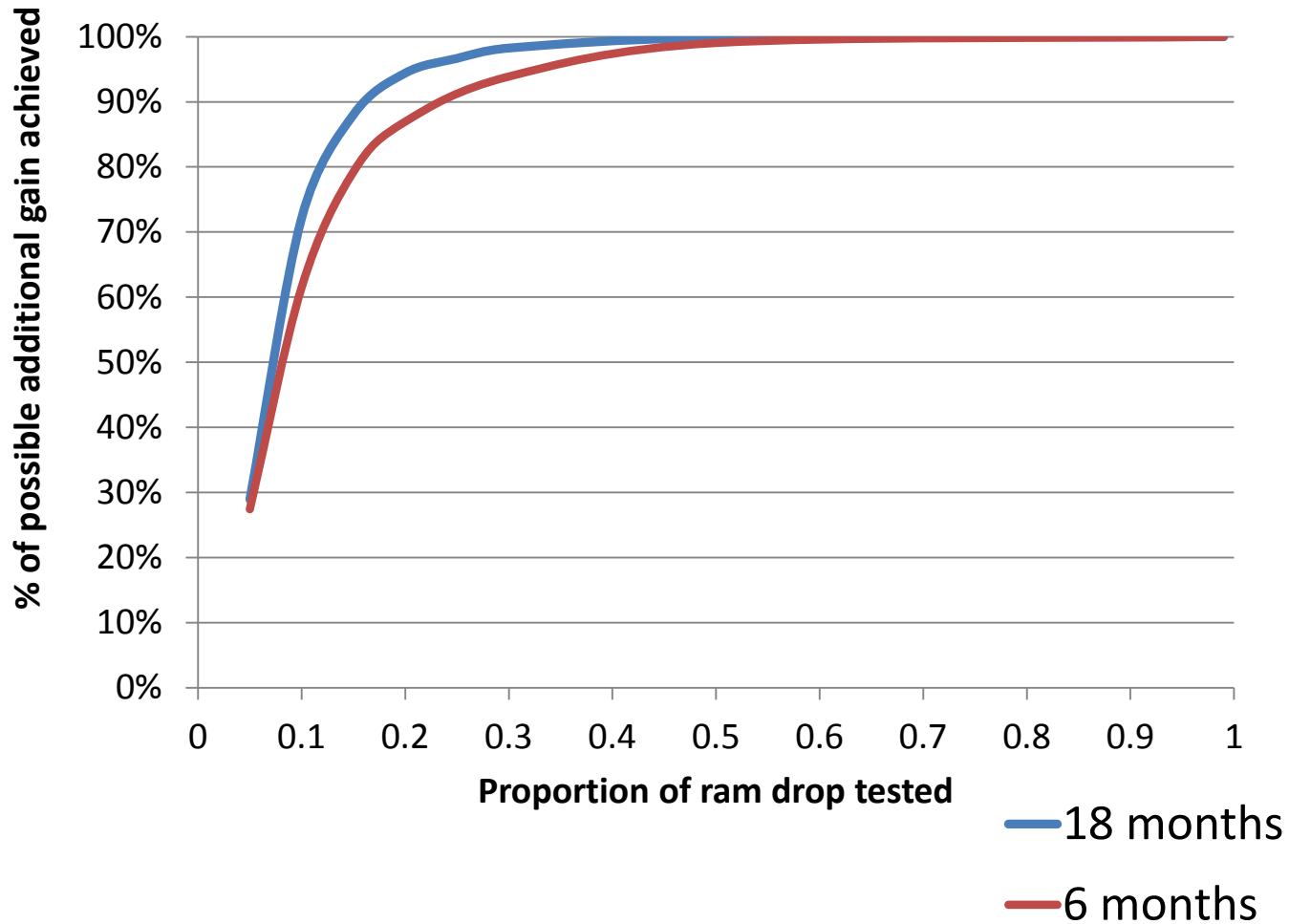
$$r_{\text{ASBV0}} = 0.01, r_{\text{GBV}} = 0.90, r = 0.01$$



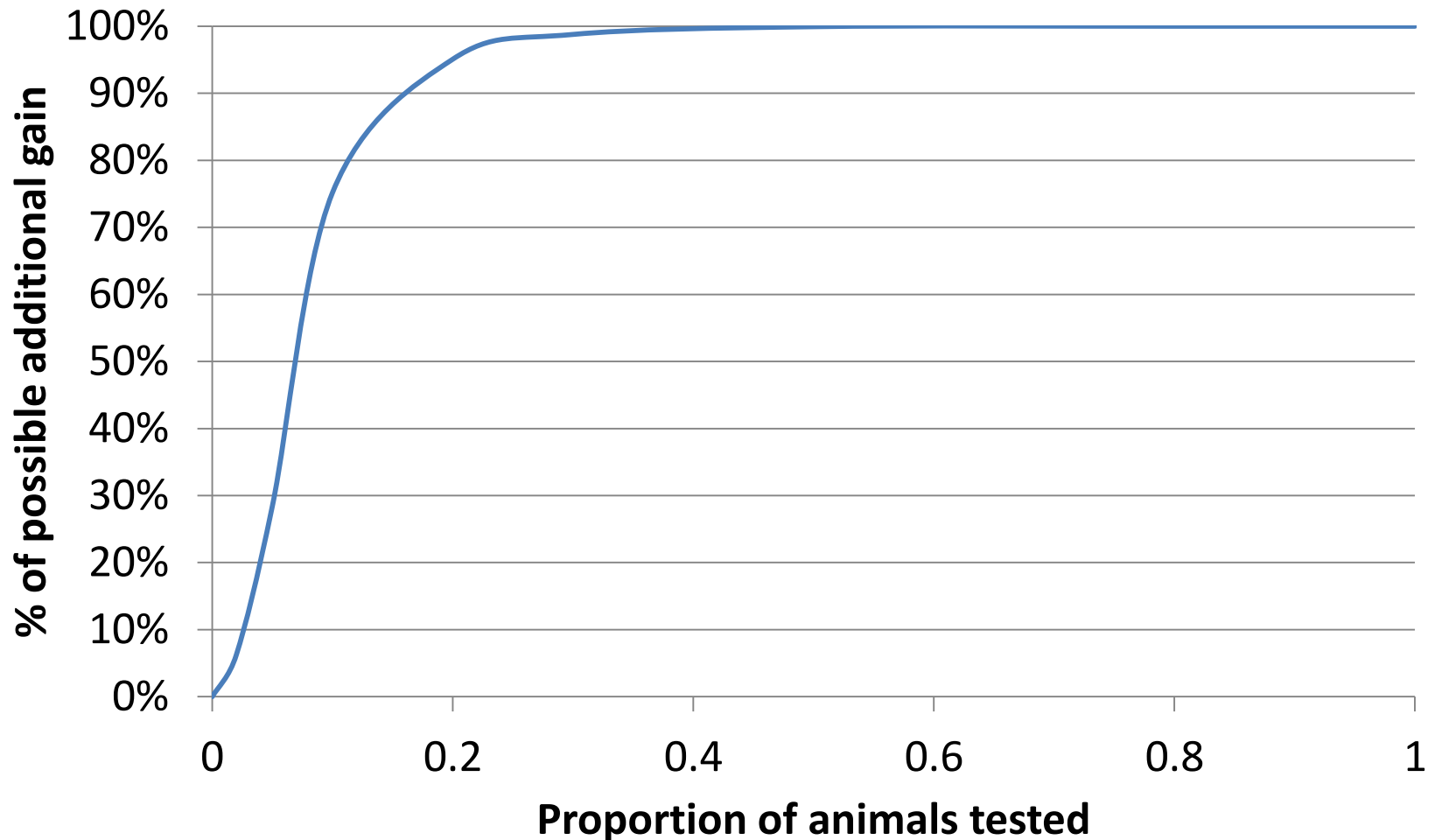
# Low ASBV acc% & high GBV

ASBV0	0.20
GBV	0.50
ASBV1	0.52
correlation ASBV0-ASBV1	0.38
prop genotyped	0.3
prop selected final	0.05
Selection Differential	0.96
SelDiff 100% genotyping	1.08
SelDiff 0% genotyping	0.40
% of possible additional gain	82%

# Testing 20% of drop gives most of benefit

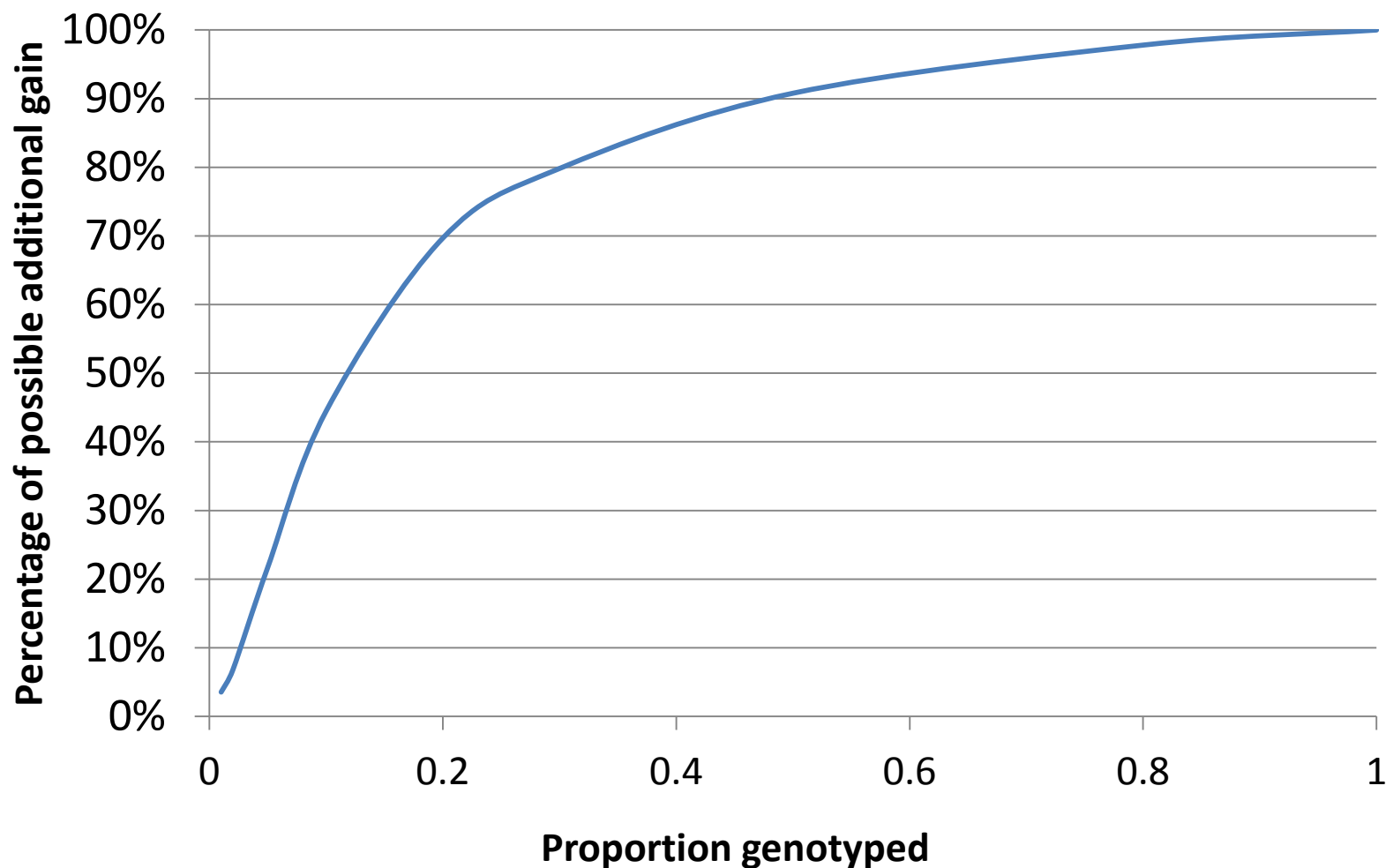


% gain compared with 100% genotyping  
ASBV0 0.34, GBV 0.39, ASBV1 = 0.50, **r = 0.7**



At high(ish) correlation between ASBV and RBV only need to genotype ~20%

% gain compared with 100% genotyping  
ASBV 0.10, GBV = 0.39, ASBV1 0.40, **r = 0.25**



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%