

une

University of
New England

Optimizing Breeding Programs

Decisions in breeding programs



Where to go?

breeding objective (which traits)

Who and what to measure?

performance, DNA test

genetic evaluation

Who to select and mate?

reproductive technol.

gains vs inbreeding

Making genetic progress is about

Selecting only the very best

Selecting accurately

$$R = \frac{i_m r_m + i_f r_f}{L_m + L_f} \sigma_A$$

Keeping generation intervals short

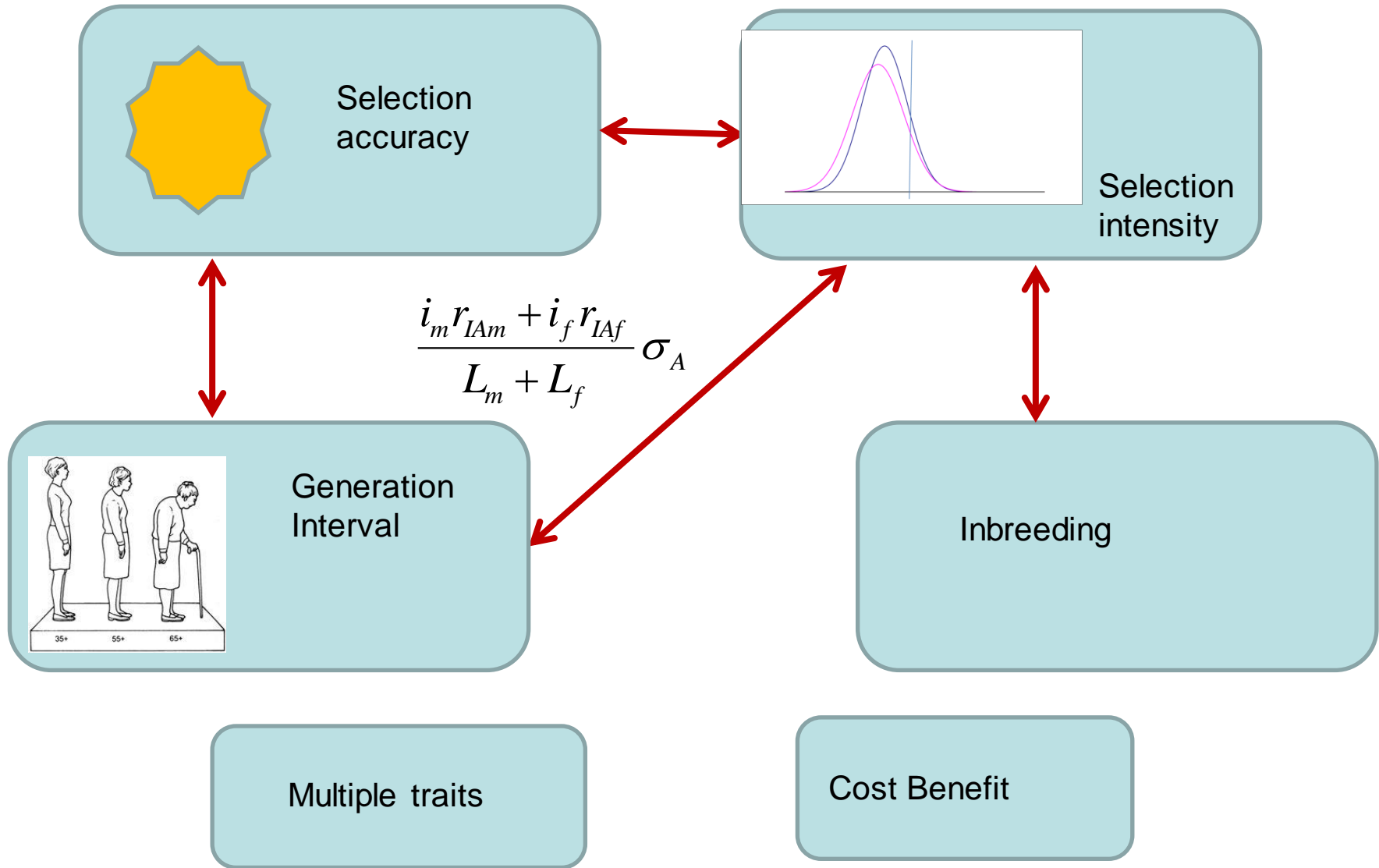
Reproductive rates affect all of the above!

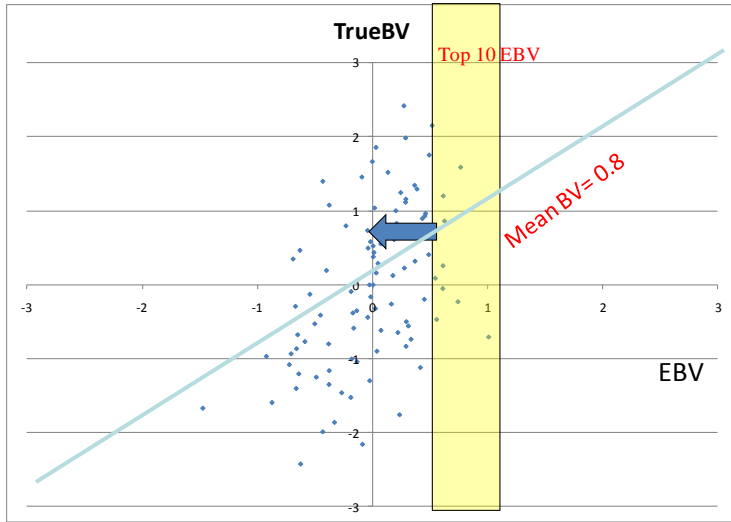
Aspects that need to be balanced:

- Selection accuracy versus generation interval
 - Short generation intervals are good for fast progress, but young breeding animals have lower EBV accuracy
- Selection accuracy versus selection intensity
 - Money available for testing (either performance or DNA) can be used to test a few animals accurately, or to test more animals with lower accuracy. For example, testing fewer young bulls but giving them more test progeny.
- Selection intensity versus generation interval
 - Selecting fewer animals for breeding each year and keeping those longer
- Selection intensity versus inbreeding
- The relative emphasis in selection for multiple traits
- Cost versus benefits

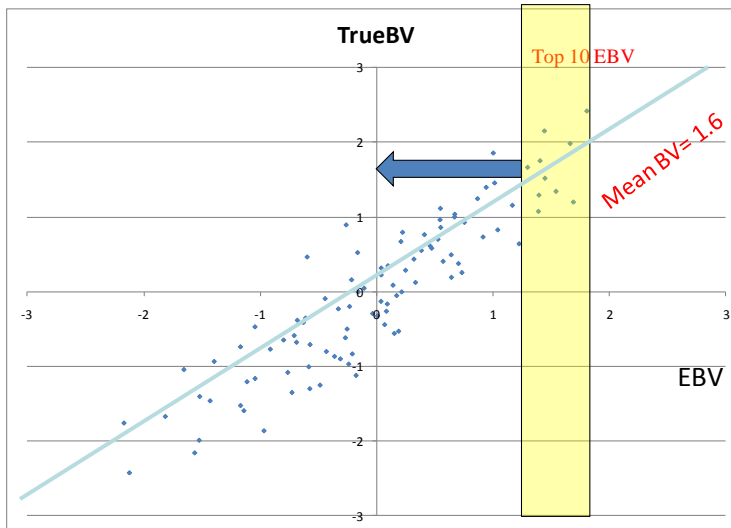
$$\frac{i_m r_{IAm} + i_f r_{IAf}}{L_m + L_f} \sigma_A$$

Aspects that need to be balanced





Accuracy = 45%

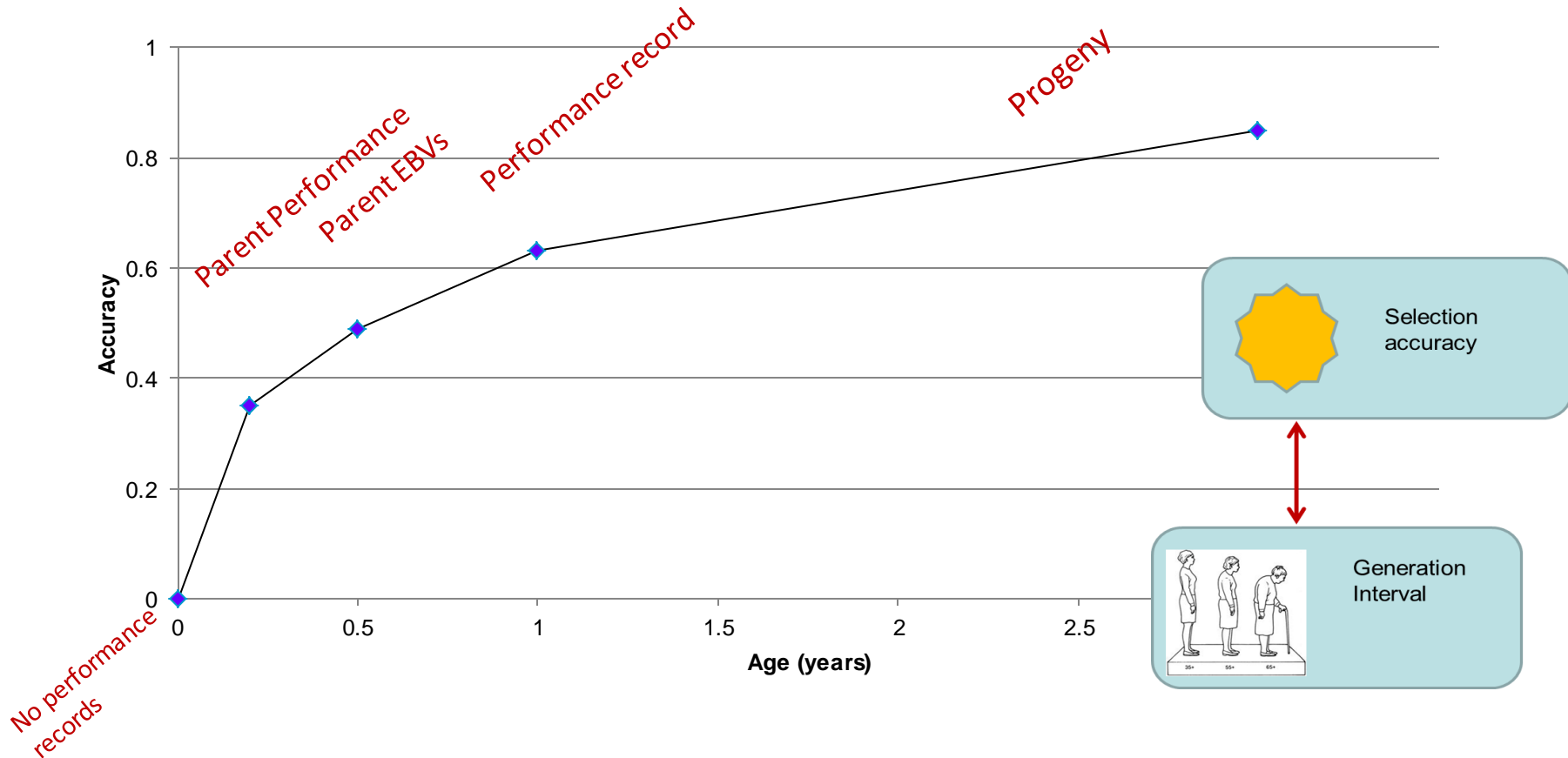


Selection accuracy

the more accuracy,
the more response

Accuracy of predicting a breeding value

- increases as an animal gets older

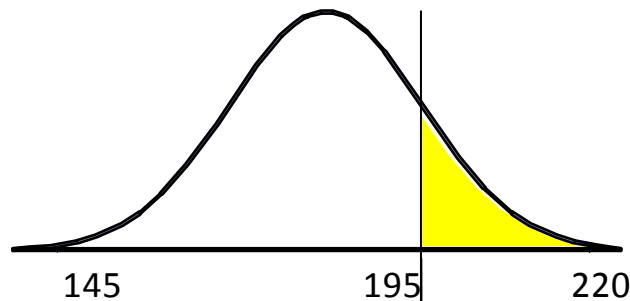


Assumed heritability = 25%

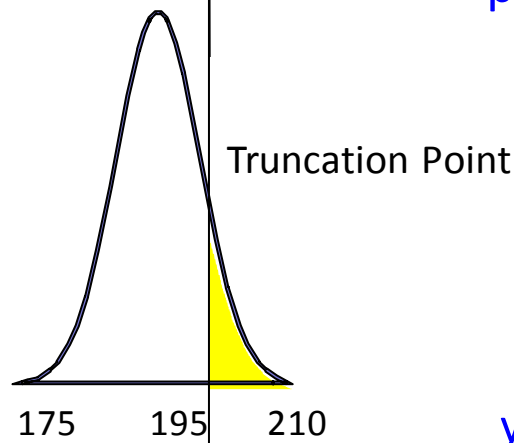
Need to balance accuracy and generation interval!

BLUP helps selecting between old and young bulls

- EBVs can be compared directly over age classes
- Selection on BLUP EBVs optimizes generation interval



proven sires

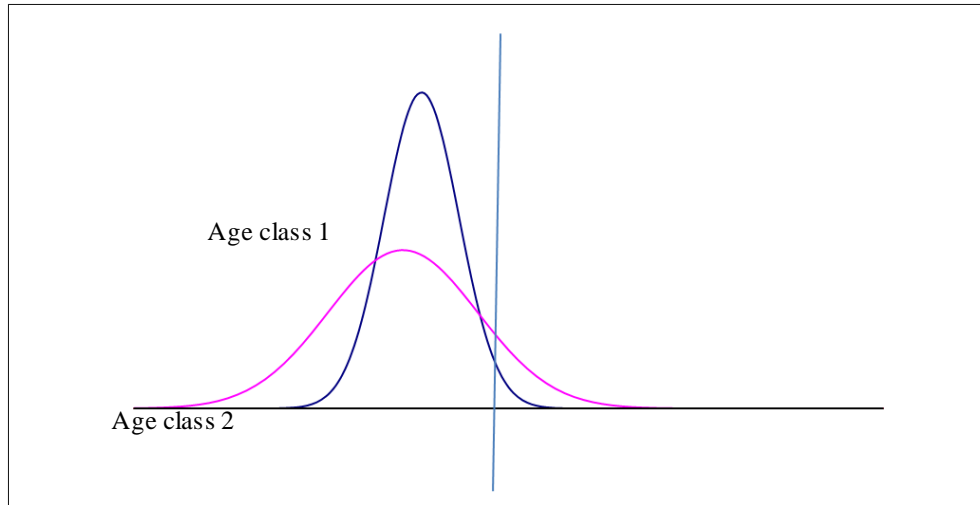


young sires



Optimizing age structure

Accuracy changes with age class !



Without genomic selection

| ageclass | N in group | mean | SD | Nr Selected |
|----------|------------|-------|-----|-------------|
| 1 | 50 | 10.20 | 0.4 | 2.7 |
| 2 | 50 | 10.00 | 0.8 | 7.3 |

Accuracy

↑

↓

With genomic selection

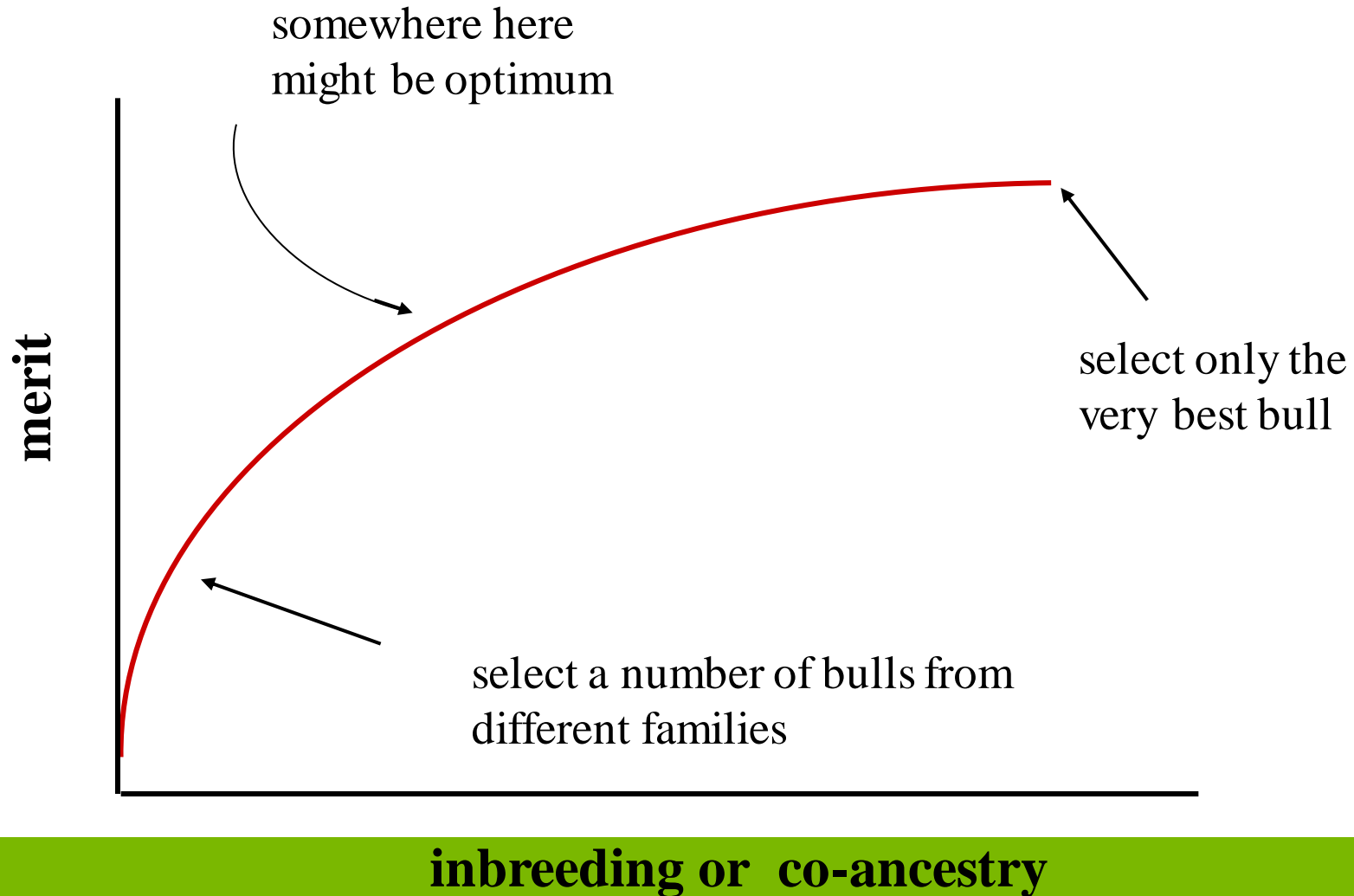
| ageclass | N in group | mean | SD | Nr Selected |
|----------|------------|-------|-----|-------------|
| 1 | 50 | 10.20 | 0.7 | 5.4 |
| 2 | 50 | 10.00 | 0.8 | 4.6 |

Best to select on EBV, irrespective of accuracy /genotyped or not / age

| | birth year | genotyped | progeny | EBV | acc |
|---------|------------|-----------|---------|------|-----|
| Kevin | 2009 | Y | 0 | +124 | 71 |
| Tony | 2005 | N | 345 | +119 | 97 |
| Bob | 2009 | N | 0 | +117 | 63 |
| John | 2008 | N | 45 | +113 | 85 |
| Paul | 2006 | N | 1087 | +112 | 99 |
| Geoff | 2009 | Y | 0 | +106 | 40 |
| Malcolm | 2007 | N | 67 | +105 | 89 |

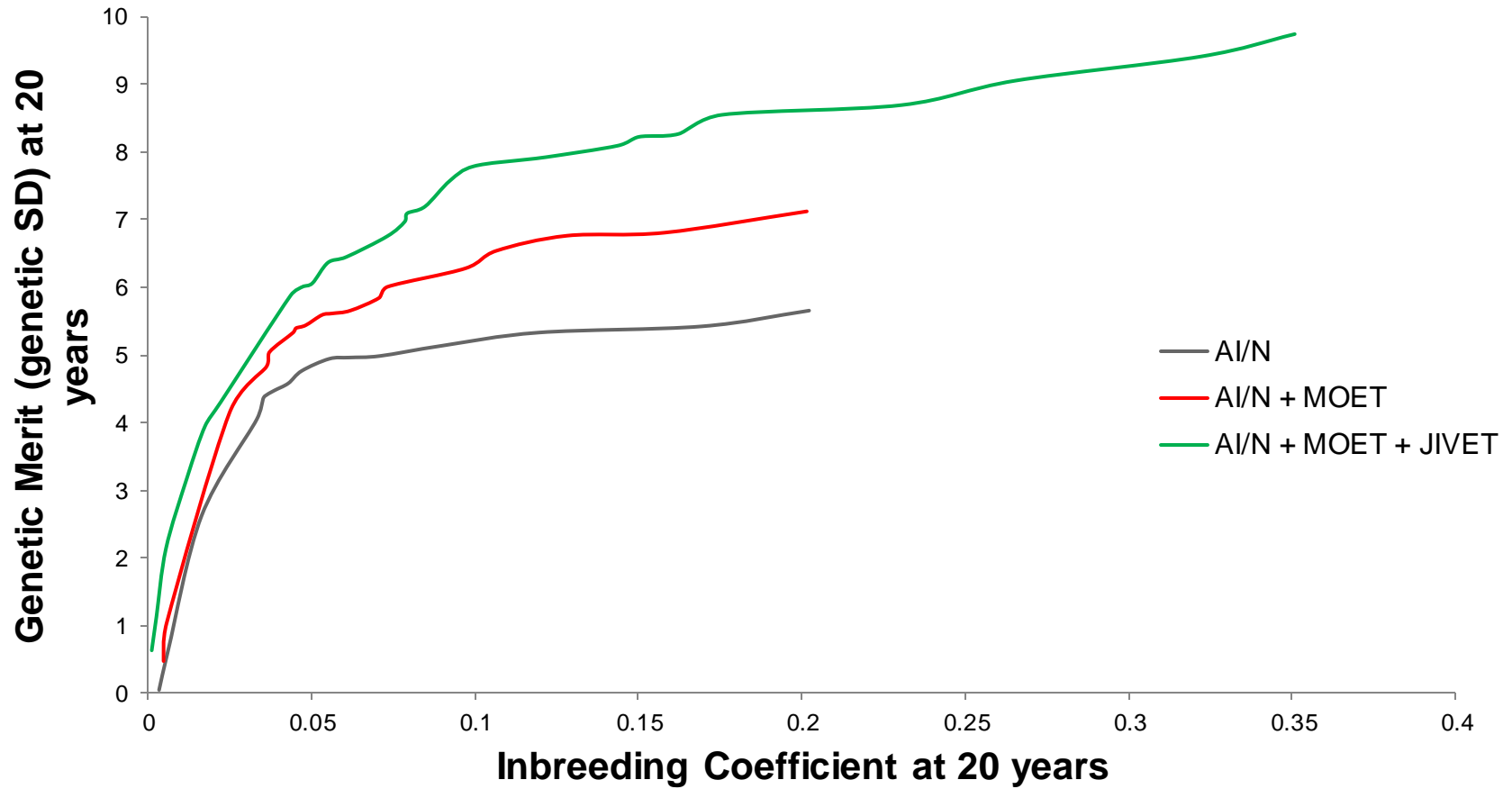
Balancing inbreeding and merit

This graph will look different for each population

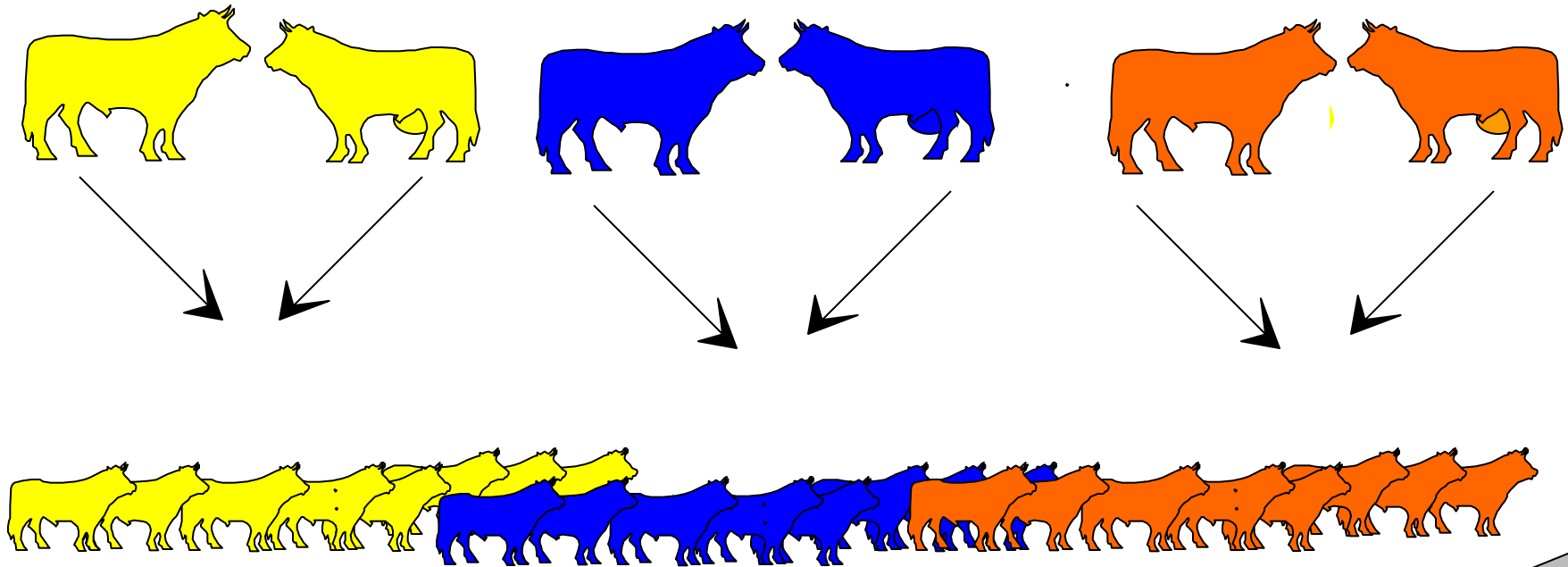


Genetic Gain vs Inbreeding After 20 Years

Tom Granleese et al., AAABG 2013



Between versus within family selection



Own information (performance or *genotype*):

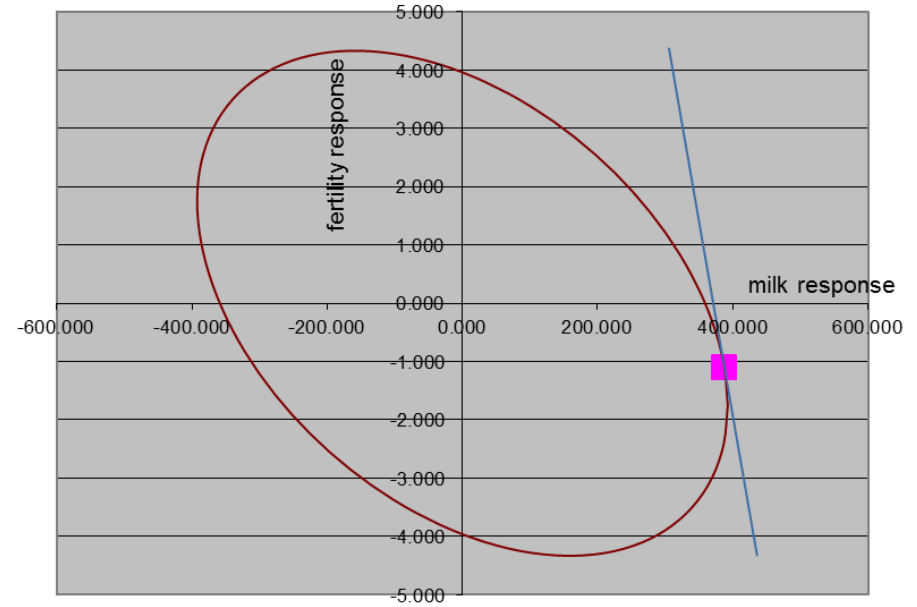
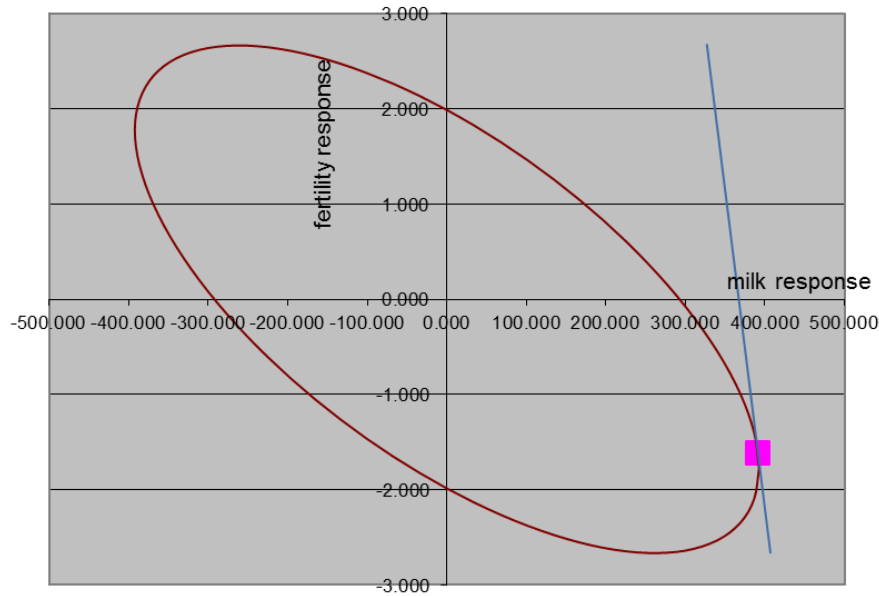
More variation within families

More within-family selection – ***less inbreeding***

Advantage of
genomic selection

Balancing Traits, weights and information

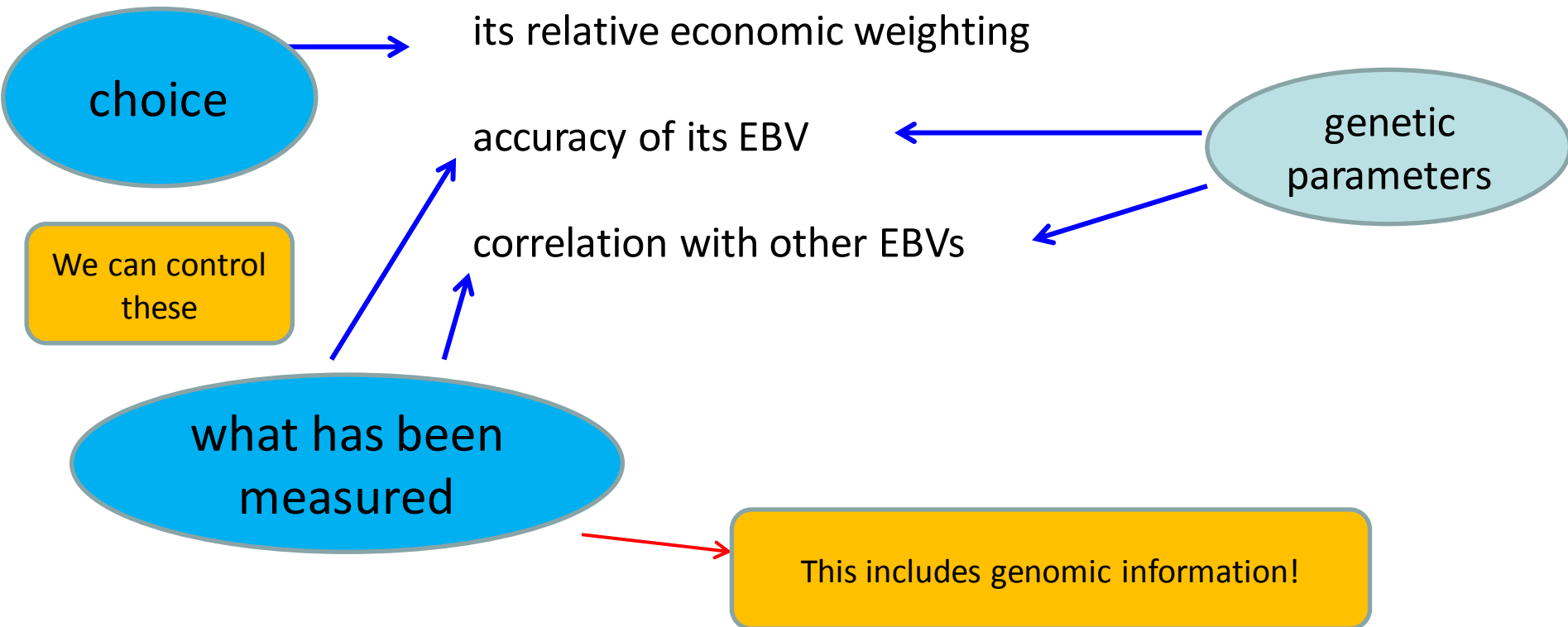
Multiple traits



Usually push the traits that have more information/higher EBV accuracy
→ Balance may change with genomic information on 'hard to measure traits'

Importance of Trait measurement

1 The ultimate response of a trait will depend on:



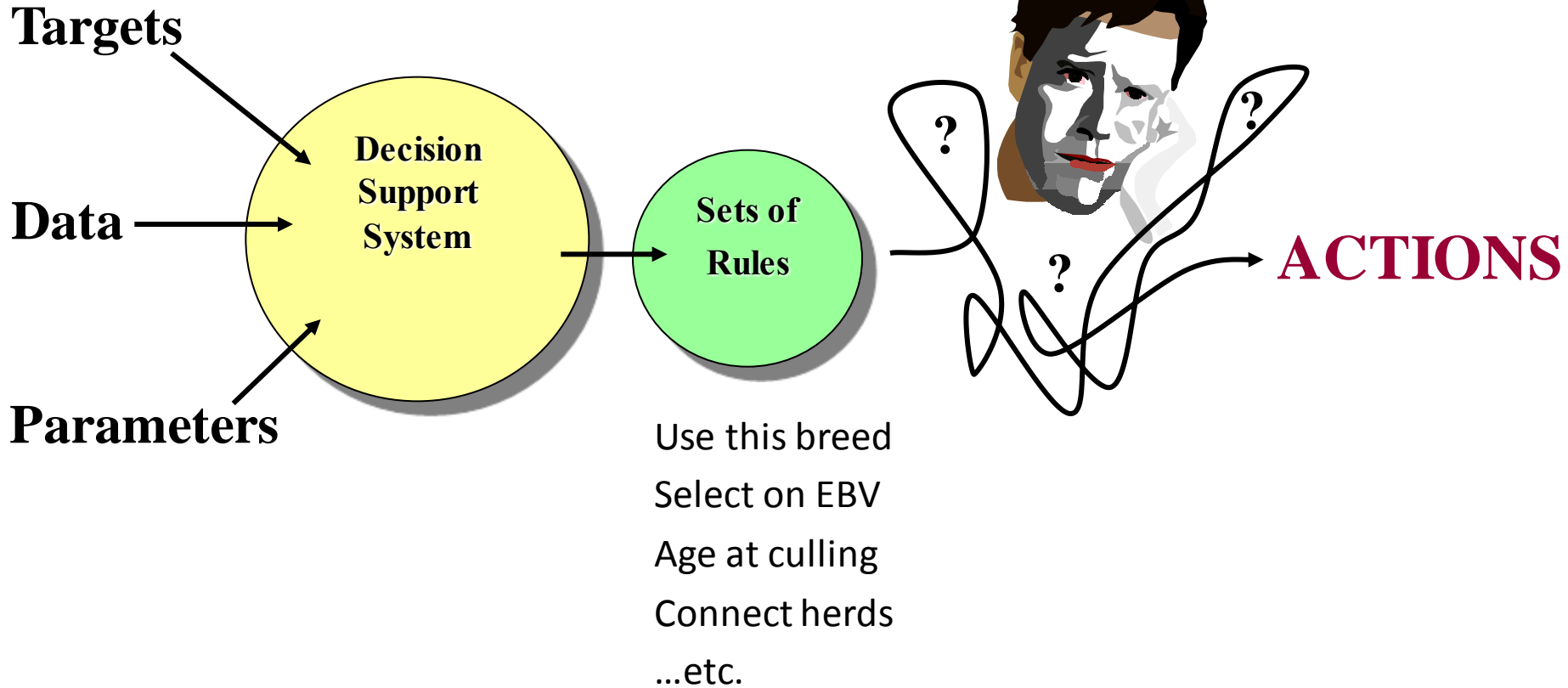
Evaluating Breeding programs

- Deterministic vs Stochastic Simulation
- Optimization strategies
- Rule based vs tactical design of breeding programs

Implementation of programs ...

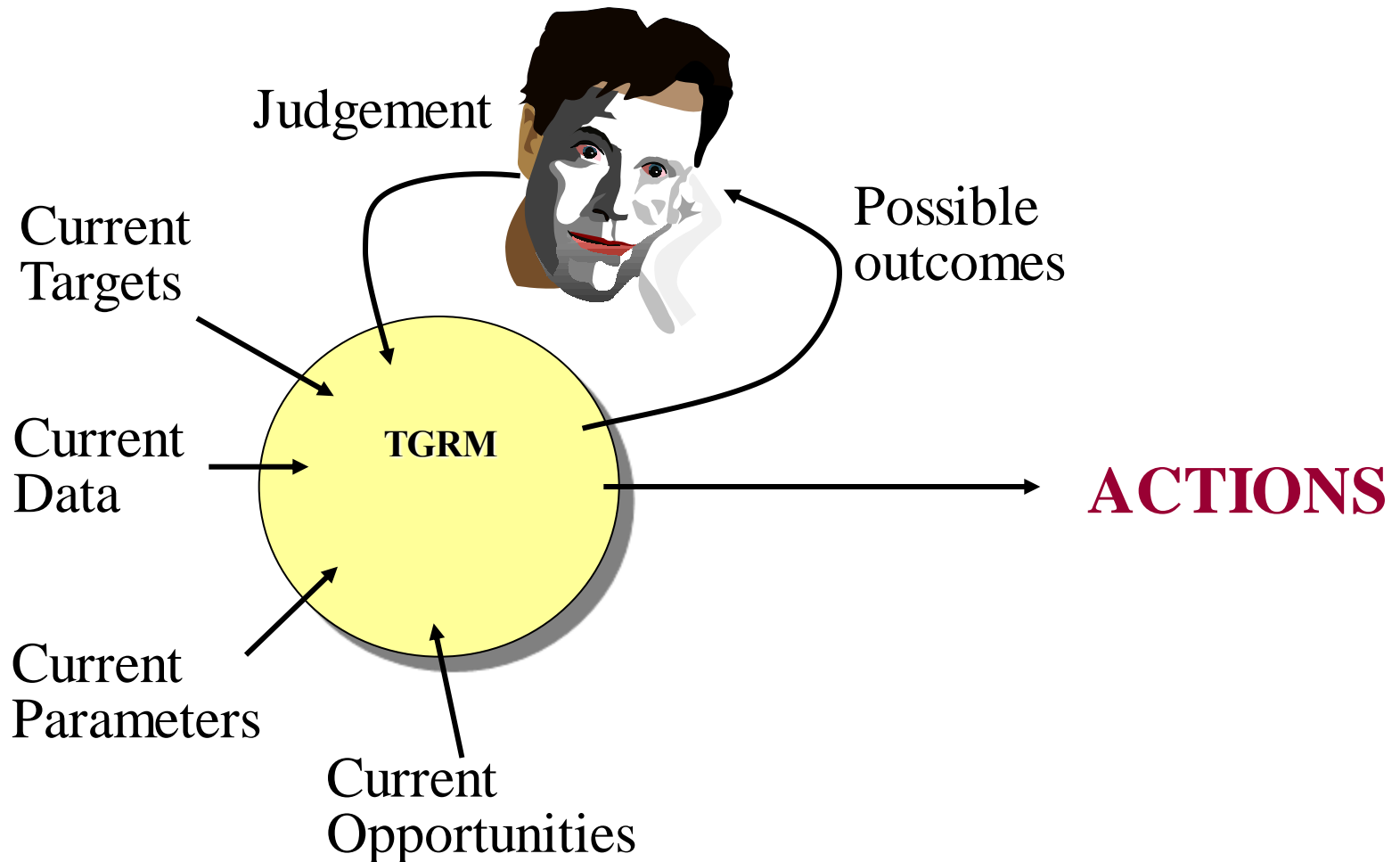
- Rules-based approach:
 - “Start joining on 1st February”
 - “Use best 10 rams mated to best 400 ewes”
 - “Set up a rotational cross”
- Tactical approach
 - Maximise impact of selection and mating, based on *prevailing* animals, markets, costs, constraints and opportunities.

Rules-based approach to Design


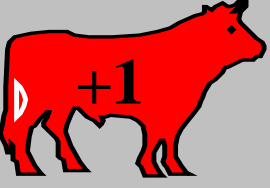
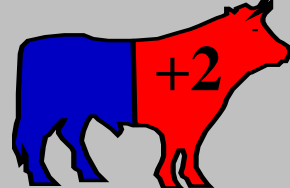
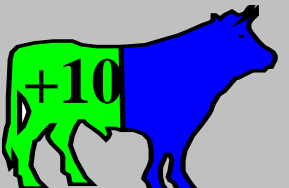
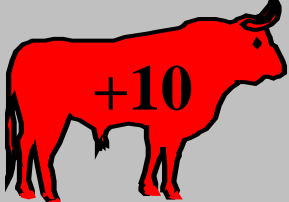

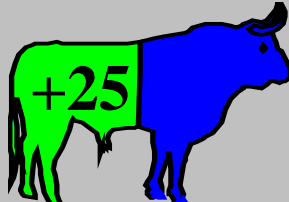

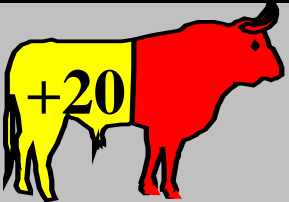



Tactical approach to Design

Action Decision Systems



Mate allocations ...

|  Cost |  0 |  10 |  10 |
|---|--|---|---|
|  0 | 311 | 312 |  |
|  3 |  | 322 | 322 |
|  2 | 309 |  | 345 |

Mate Selection Control Centre

Frontiers
Breeders Pick

Inbreeding

Target Degrees: 51.3

Progeny F Weight: -10.0

Progeny H Weight: 0.0

Trait Constraints

Trait: 600d imf% en

Type: None Min Min

Value: 0.449 1.44

Weight: 500 500

Hist:

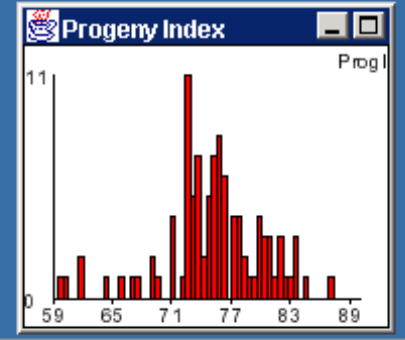
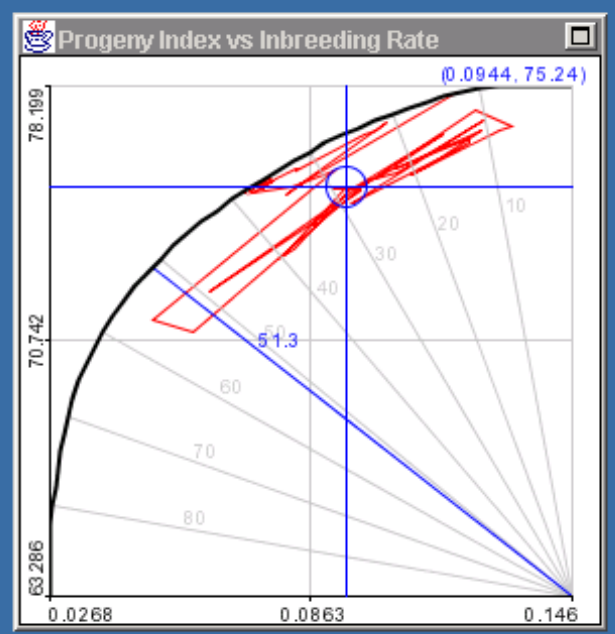
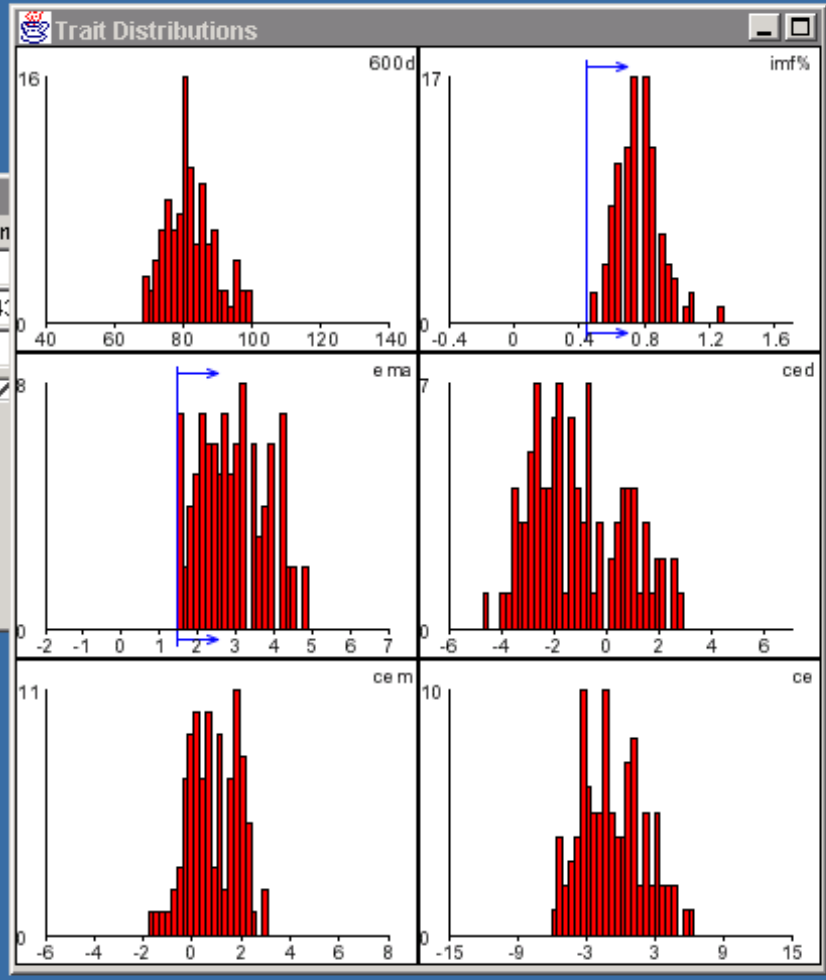
Trait: cem ce

Type: None None

Value:

Weight:

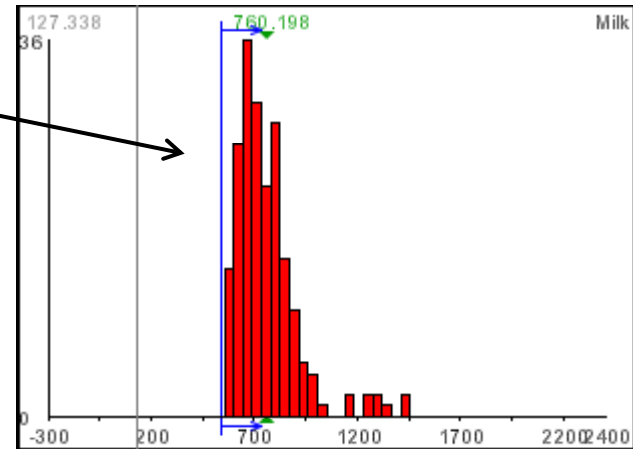
Hist:



| ID | Sire | Index | Avg. xAx | Use | MaxUse | MinUse | MustUse | Cost | IsAI | Weight | xAx |
|---------|---------|-------|----------|-----|--------|--------|--------------------------|------|-------------------------------------|--------|-----|
| USA036 | USA315 | 97.57 | 0.003 | 54 | 95 | 0 | <input type="checkbox"/> | 0 | <input checked="" type="checkbox"/> | 0.1 | 0 |
| USA315 | USA9958 | 96.48 | 0.003 | 34 | 95 | 0 | <input type="checkbox"/> | 0 | <input checked="" type="checkbox"/> | 0.1 | 0.0 |
| USA323 | USA036 | 81.71 | 0.003 | | 80 | 0 | <input type="checkbox"/> | 0 | <input checked="" type="checkbox"/> | 0.1 | |
| USA3246 | USA5204 | 79.46 | 0.002 | | 80 | 0 | <input type="checkbox"/> | 0 | <input checked="" type="checkbox"/> | 0.1 | |
| NXOP97 | USA2172 | 76.96 | 0.002 | | 50 | 0 | <input type="checkbox"/> | 0 | <input checked="" type="checkbox"/> | 0.1 | |

Achieving Trait Constraints

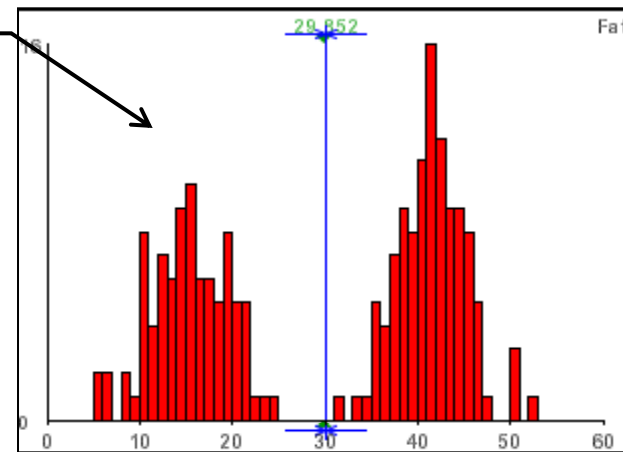
All progeny are predicted to be above the restriction of +545Kg



Predicted progeny Milk EBVs

Achieving Trait Constraints

Targeting two different objectives in one cycle of matings

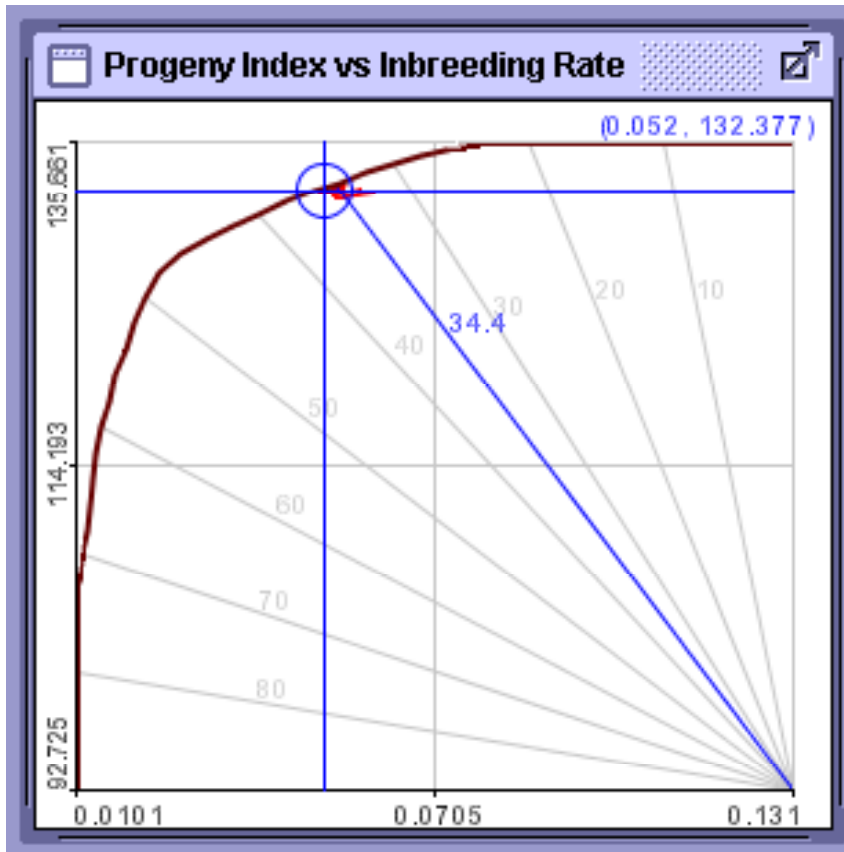


Predicted progeny Fat EBVs

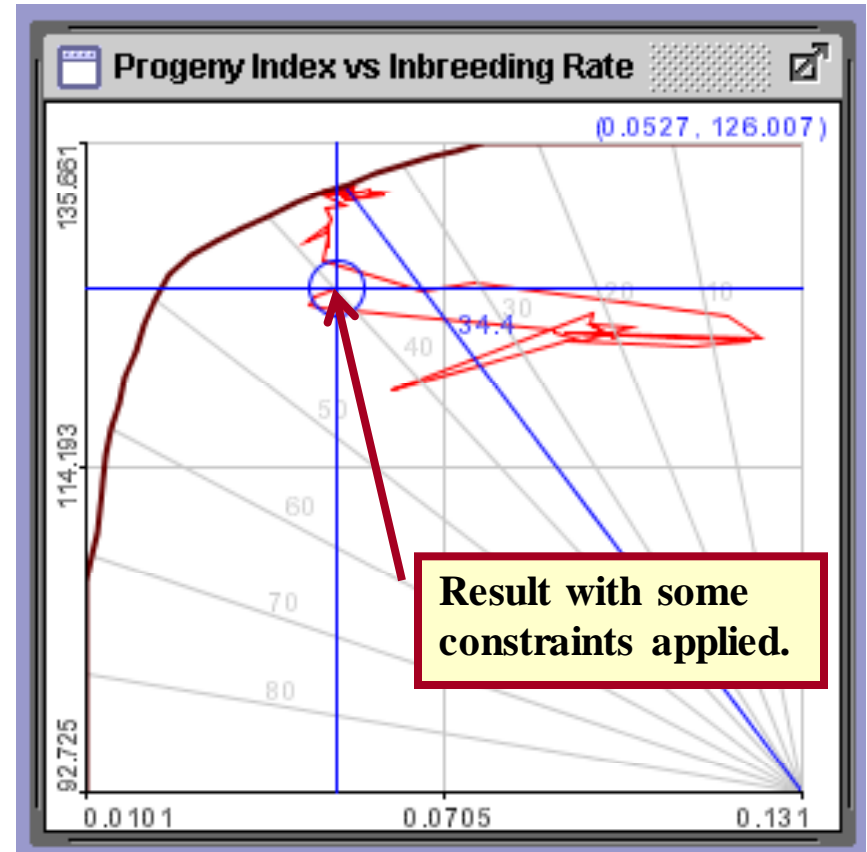
Imposing constraints

(eg. Sire use, QTL outcome, trait distributions)

Predicted progeny index



Mean parental coancestry



Mean parental coancestry



Mating List (by Sire)

Total Genetic Resource Management

[Report List](#)
[Report Summary](#) | [Sires Summary](#) | [Mating List](#) | [Mating List \(CSV\)](#) | [Mating List \(TXT\)](#)

| up/down Sire | up/down Dam | up/down index | up/down F | up/down SireCoan | up/down DamCoan | up/down SortIndex | up/down 600d-1 | up/down 600d-2 | up/down imf% | up/down ema-1 | up/down ema-2 | up/down ced | up/down cem | up/down ce |
|---------------------------------|--------------------------------|----------------------------------|------------------------------|-------------------------------------|------------------------------------|--------------------------------------|-----------------------------------|-----------------------------------|---------------------------------|----------------------------------|----------------------------------|--------------------------------|--------------------------------|-------------------------------|
| USA3246 | T229 | 66.68 | 0.0156 | 0.003228 | 0.000138 | 64.9366 | 88.00 | 88.00 | 0.50 | 0.15 | 0.15 | 1.25 | -0.70 | 0.02 |
| USA3246 | T213 | 70.18 | 0.0000 | 0.003228 | 0.000175 | 69.9997 | 81.50 | 81.50 | 0.65 | 1.55 | 1.55 | 2.35 | 0.48 | 3.48 |
| USA3246 | T157 | 64.68 | 0.0235 | 0.003228 | 0.000000 | 62.1640 | 85.00 | 85.00 | 0.45 | 0.20 | 0.20 | 2.81 | -1.08 | 0.83 |
| USA3246 | T137 | 65.64 | 0.0000 | 0.003228 | 0.000000 | 65.4690 | 72.00 | 72.00 | 0.70 | 0.40 | 0.40 | 2.88 | 1.52 | 6.09 |
| USA3246 | T117 | 68.09 | 0.0078 | 0.003228 | 0.000087 | 67.1344 | 76.00 | 76.00 | 0.65 | 1.45 | 1.45 | 2.80 | 0.81 | 4.59 |
| USA3246 | T063 | 76.10 | 0.0039 | 0.003228 | 0.000000 | 75.5340 | 77.00 | 77.00 | 0.90 | 1.35 | 1.35 | -0.09 | 0.03 | 0.14 |
| USA3246 | T057 | 73.06 | 0.0000 | 0.003228 | 0.000000 | 72.8840 | 70.00 | 70.00 | 0.90 | -0.55 | -0.55 | 2.74 | 0.34 | 3.60 |
| USA3246 | T029 | 64.08 | 0.0235 | 0.003228 | 0.000132 | 61.5570 | 77.50 | 77.50 | 0.60 | 0.40 | 0.40 | 3.38 | -0.97 | 1.62 |
| USA3246 | T020 | 75.63 | 0.0078 | 0.003228 | 0.000000 | 74.6740 | 90.50 | 90.50 | 0.65 | 1.45 | 1.45 | 1.87 | -0.43 | 1.17 |
| USA3246 | T013 | 67.38 | 0.0000 | 0.003228 | 0.000133 | 67.2019 | 77.50 | 77.50 | 0.70 | 1.20 | 1.20 | 1.69 | 0.48 | 2.82 |
| USA3246 | T008 | 72.18 | 0.0000 | 0.003228 | 0.000298 | 71.9982 | 73.50 | 73.50 | 0.75 | 1.05 | 1.05 | 3.04 | 2.21 | 7.62 |
| USA3246 | S305 | 63.88 | 0.0078 | 0.003228 | 0.000141 | 62.9215 | 81.50 | 81.50 | 0.55 | 1.35 | 1.35 | -0.42 | -0.71 | -1.67 |
| USA3246 | R001 | 66.58 | 0.0000 | 0.003228 | 0.000000 | 66.4090 | 67.50 | 67.50 | 0.80 | 1.35 | 1.35 | 3.75 | 1.14 | 6.20 |
| USA3246 | Q075 | 62.24 | 0.0000 | 0.003228 | 0.009841 | 61.5426 | 84.00 | 84.00 | 0.55 | 0.35 | 0.35 | 2.09 | 0.25 | 2.76 |
| USA3246 | Q001 | 73.62 | 0.0000 | 0.003228 | 0.000000 | 73.4490 | 73.00 | 73.00 | 0.80 | 1.30 | 1.30 | 2.50 | 0.36 | 3.40 |
| USA323 | R211 | 67.93 | 0.0000 | 0.000296 | 0.000093 | 67.9094 | 87.50 | 87.50 | 0.55 | 2.85 | 2.85 | -3.00 | -2.26 | -8.13 |
| USA315 | T99 | 74.46 | 0.0000 | 0.022720 | 0.000116 | 73.2452 | 89.50 | 89.50 | 0.70 | 2.00 | 2.00 | 0.22 | -0.21 | -1.76 |
| USA315 | T270 | 79.83 | 0.0000 | 0.022720 | 0.000000 | 78.6263 | 94.00 | 94.00 | 0.80 | 2.60 | 2.60 | -0.14 | 0.27 | -1.15 |
| USA315 | T259 | 72.26 | 0.0000 | 0.022720 | 0.000114 | 71.0503 | 85.50 | 85.50 | 0.70 | 0.55 | 0.55 | -0.46 | -0.01 | -2.04 |
| USA315 | T243 | 76.79 | 0.0625 | 0.022720 | 0.000154 | 69.3281 | 88.00 | 88.00 | 0.75 | 2.95 | 2.95 | -1.00 | 0.20 | -2.15 |
| USA315 | T217 | 72.43 | 0.0000 | 0.022720 | 0.000000 | 71.2263 | 82.00 | 82.00 | 0.70 | 0.55 | 0.55 | 0.49 | 0.13 | -0.80 |



Genetic Gain vs Inbreeding while using female reproductive technologies

Tom Granleese, 2015

Reproductive technologies

Natural

MOET

JIVET



Mature Ewe

Mature Ewe

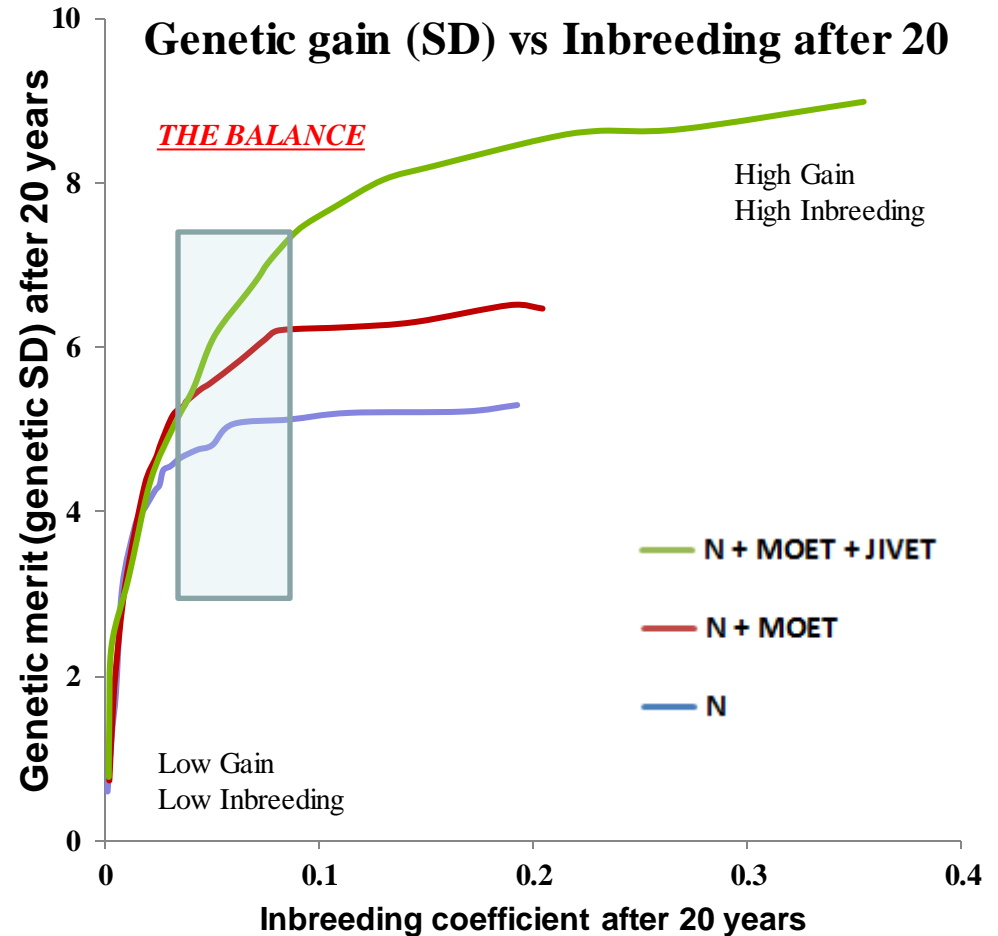
Juvenile Ewe



1 Lamb

Many Lambs

Loads of Lambs



Proportion of females assigned technologies

at 1% ΔdF per gen

| | AI/N | MOET | JIVET | Total Females* | Males Used* | Females per male |
|--------------------|------|------|-------------|----------------|-------------|------------------|
| Early Trait | | | | | | |
| With GS | 0.29 | 0.28 | 0.43 | 85 | 19 | 4.5 |
| NO GS | 0.34 | 0.36 | 0.30 | 88 | 20 | 4.4 |
| Late Trait | | | | | | |
| With GS | 0.31 | 0.26 | 0.43 | 88 | 14 | 6.3 |
| NO GS | 0.34 | 0.35 | 0.31 | 89 | 15 | 6.0 |
| Dairy | | | | | | |
| With GS | 0.38 | 0.28 | 0.34 | 218 | 39 | 5.6 |
| NO GS | 0.47 | 0.35 | 0.18 | 237 | 41 | 5.8 |

GS SHIFTS PROPORTION to JIVET

Compensate female lack of diversity with more sire diversity



Optimizing use of repro technologies

| Proportion Captured by breeder | AI | MOET | JIVET | Dams Used | G/yr (\$) | L |
|--------------------------------|------|------|-------|-----------|-----------|------|
| 0.06 | 0.95 | 0.00 | 0.05 | 261 | \$2.26 | 1.87 |
| 0.32 | 0.77 | 0.04 | 0.19 | 221 | \$2.82 | 1.46 |
| 0.64 | 0.36 | 0.10 | 0.54 | 136 | \$3.96 | 1.21 |



If breeder captures more benefit she/he can afford to invest more and make more gain