

Total Genetic Resource Management

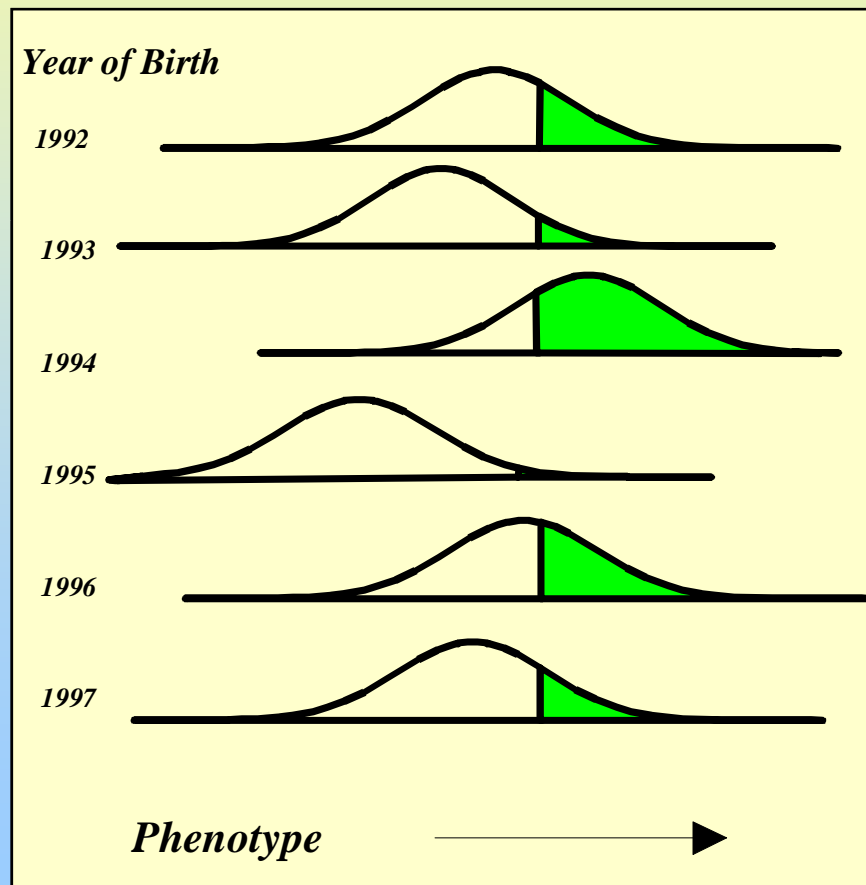
TGRM

Rules-based approach to Design

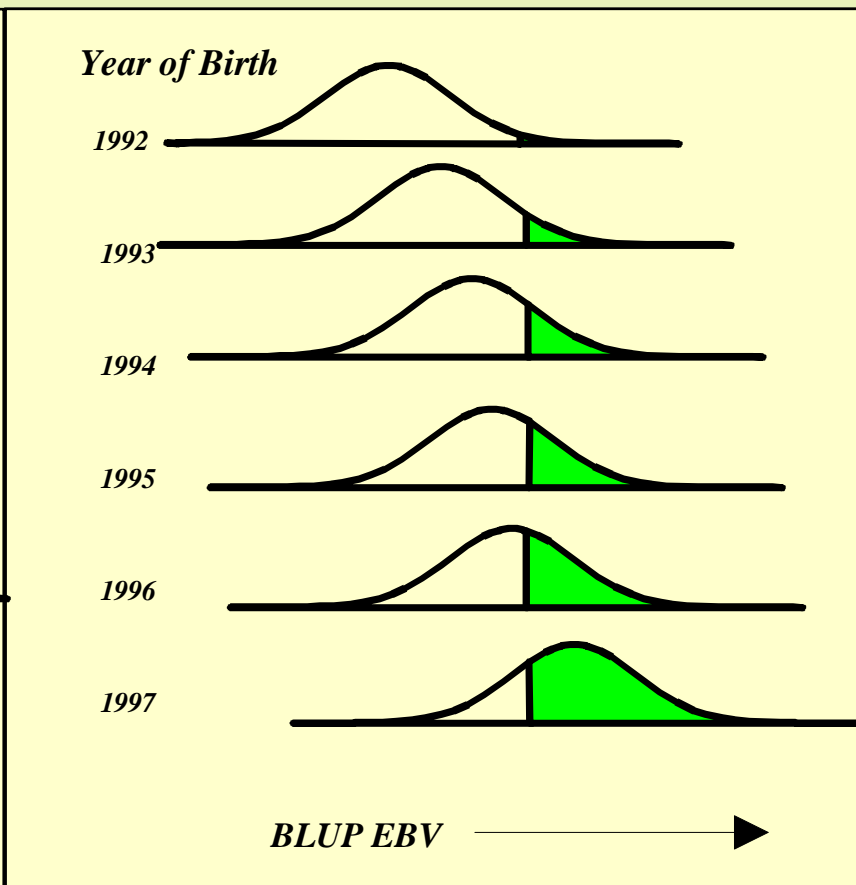
- Different methods have been developed for handling different animal breeding issues (eg. selection, inbreeding, crossing).
 - When we try to mix different methods (or mix different rules) we are likely to miss the best overall design.
- Following strict rules leads to missed opportunities. We should:
 - Fine-tune the rules as appropriate
 - Break the rules when appropriate

A genetic evaluation system helps to design breeding programs.

Select on phenotype



Select on EBV



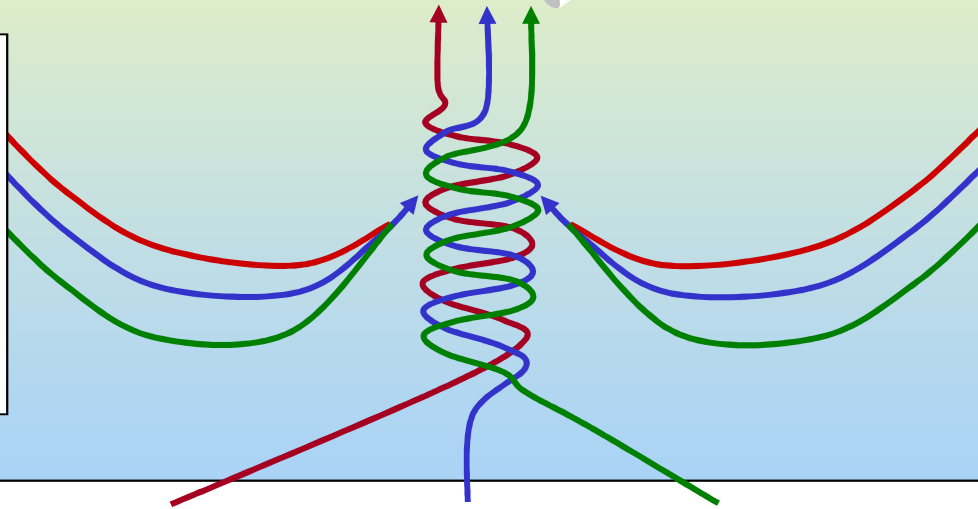
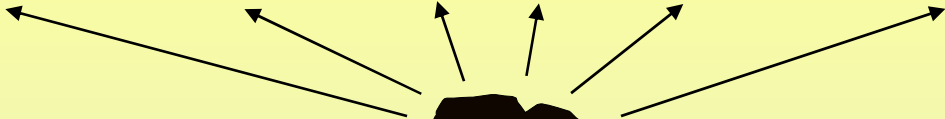
Animal breeding actions Selections
Purchases Matings Cullings etc.

Experience Constraints
Attitudes **Agap** Costs

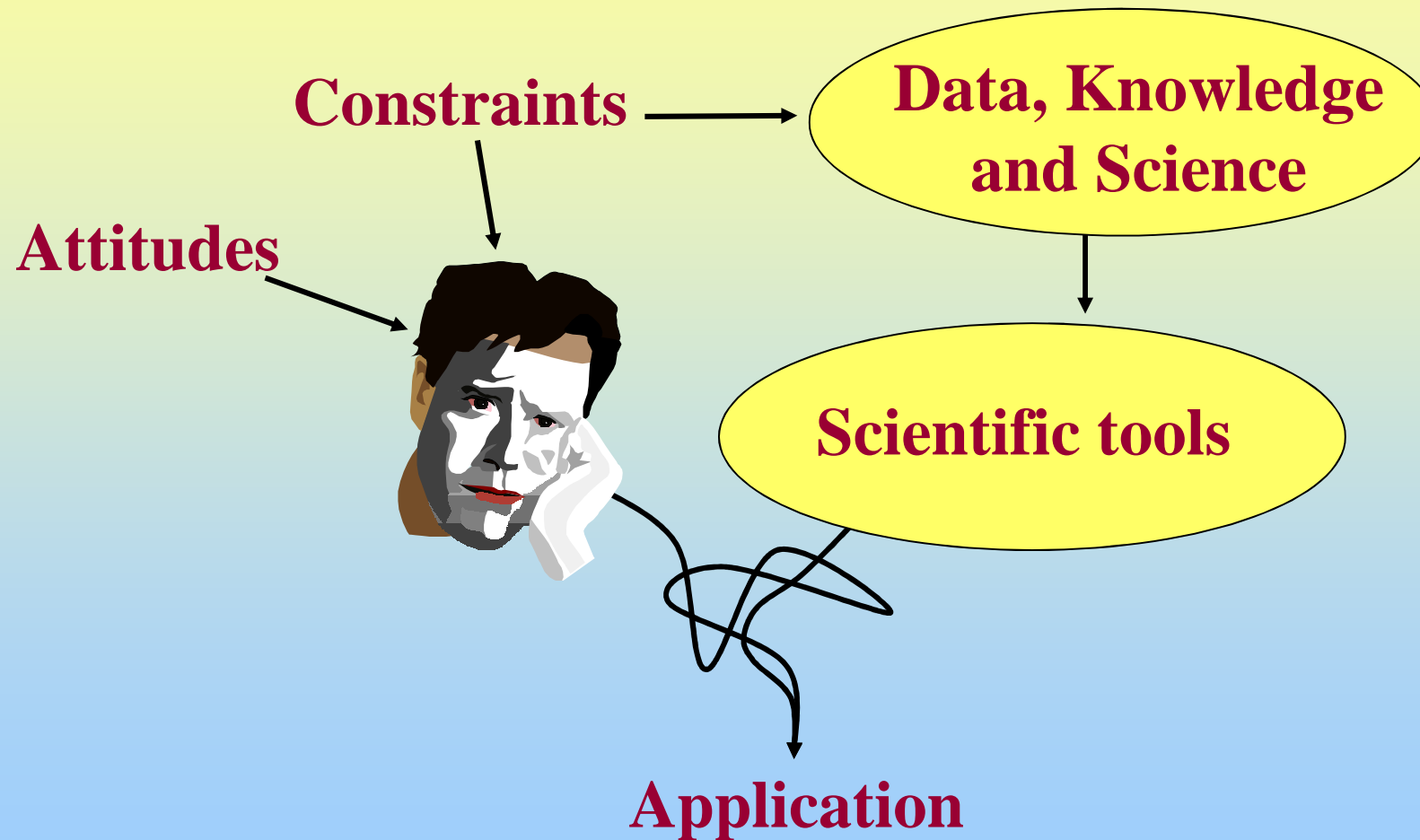
Breeding Program Design Rules

Breeding Program Design Rules

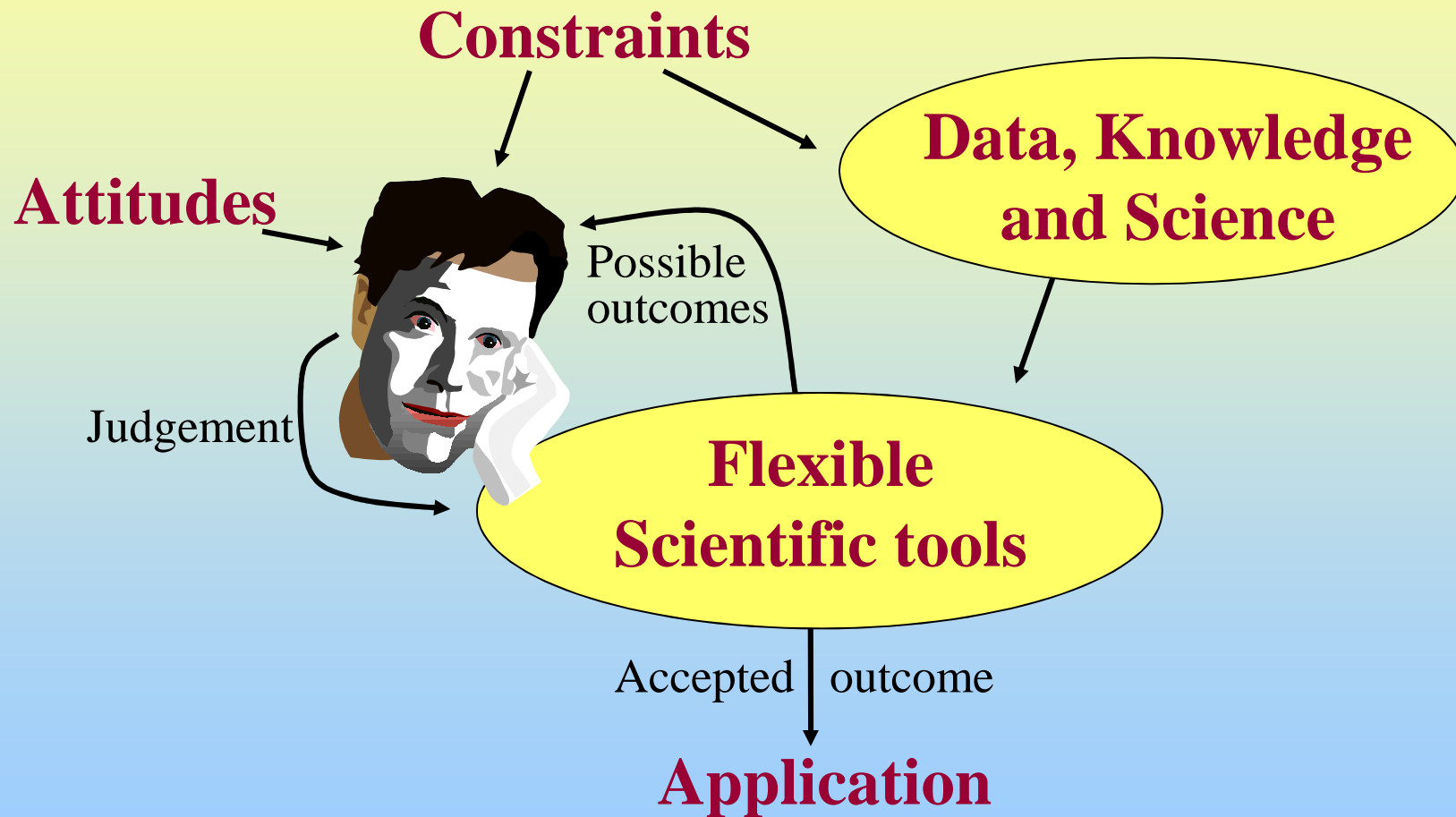
EBVs **QTLs** **Accuracies**
Pedigree Data Parameters Economics Methods Markers



A classic decision system



A Dynamic Tactical Decision System



Key actions for Animal Breeders

- Selection

- Mating

Σ = “Mate selection”

Mating List - Netscape

File Edit View Go Communicator Help

Back Forward Reload Home Search Netscape Print Security Shop Stop

Location: http://tgrm.une.edu.au/servlet/reports_show/987823367381_0018_01/matlst0.htm What's Related

Total Genetic Resource Management
[Report List](#)

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

up/down <u>Sire</u>	up/down <u>Dam</u>	up/down <u>index</u>	up/down <u>F</u>	up/down <u>SireCoan</u>	up/down <u>DamCoan</u>	up/down <u>SortIndex</u>	up/down <u>600d-1</u>	up/down <u>600d-2</u>	up/down <u>imf%</u>	up/down <u>ema-1</u>	up/down <u>ema-2</u>	up/down <u>ced</u>	up/down <u>cem</u>	up/down <u>ce</u>
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

Document: Done

“Total Genetic Resource Management”

Covers ...

- ◆ Selection
- ◆ Crossbreeding
- ◆ Costs
- ◆ Genetic diversity
- ◆ Optimal selection
- ◆ Progeny inbreeding
- ◆ Connection between herds
- ◆ Limits on reproduction
- ◆ Marker Assisted Selection
- ◆ Use of AI, MOET, IVF
- ◆ Corrective mating
- ◆ Managing trait distribution
- ◆ Funding limits
- ◆ Quarantine barriers
- ◆ **Logistical constraints**
- ◆ etc

TGRM approach ...

A candidate solution ...

Bull→ Cow↓	1	2	3	4 ...
1	mate	mate	-	mate
2	mate	-	-	-
3	-	mate	-	-
4 ...	-	-	-	mate

[... one breed or a mixture of breeds]

Evaluate for (eg.) ...

- Progeny merit
 - EBV, inbreeding, heterosis ...
- Genetic diversity
 - including long-term inbreeding.
- Connection across herds
 - plus dispersal to comm. herds.
- Costs
 - AI and MOET, stock purchase ...
- Logistical factors
 - paddocks, stock movements ...
- Broken constraints
 - fund limits, quarantine barriers ...

Try to find a
better solution

Sum to give dollar
value of this solution

A mate selection objective function:

$$\frac{x'G}{2M} + \lambda \frac{x'Ax}{4M^2} + \phi F + \chi C - cost + \text{practical constraints} + + +$$

$x'G$ is the weighted mean EBV of selected parents

x is a vector of number of matings to be made by each candidate. We need this to be handle to handle mating.

M is the total number of matings to be made

$x'Ax/4M^2$ is the weighted mean coancestry of selected parents.

λ is a weighting factor that is typically negative, to discourage low effective population sizes.

A is the numerator relationship matrix.

ϕ is the weighing on progeny inbreeding (F).

χ is the weighing on progeny heterosis (C)

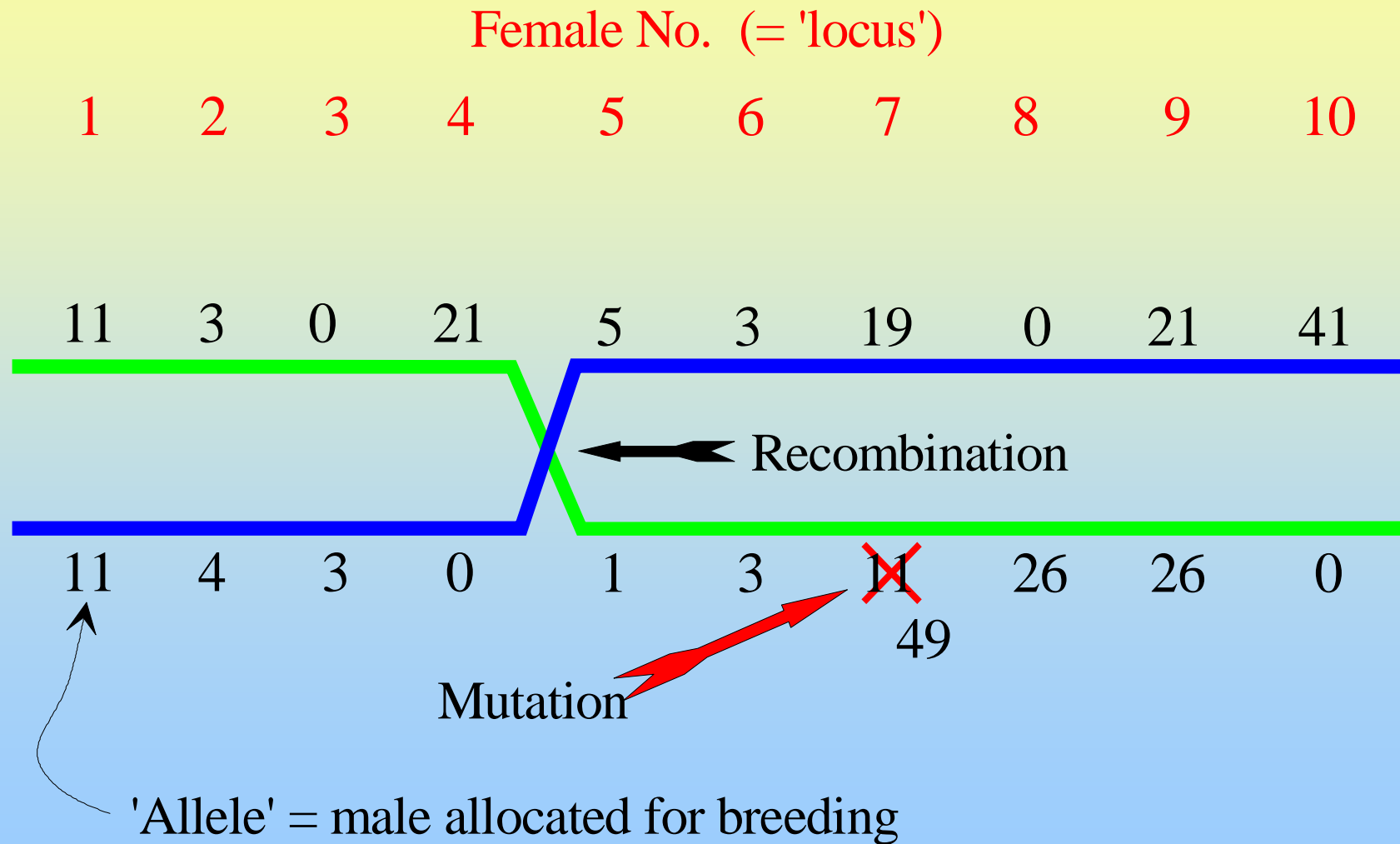
$cost$ is the cost of the mating policy implied by x .

xGxAx.xls ...

Vector x : Number of matings

Source of animals	Animal#	$x =$ Matings	
Male candidates	1	0	} $\Sigma =$ Target number of matings
	2	21	
	3	33	
	4	0	
	5	175	
	6	0	
	7	0	
	8	0	
	
Female candidates	101	0	} $\Sigma =$ Target number of matings
	102	1	
	103	1	
	104	1	
	105	0	
	106	0	
	107	0	
	108	8	
	

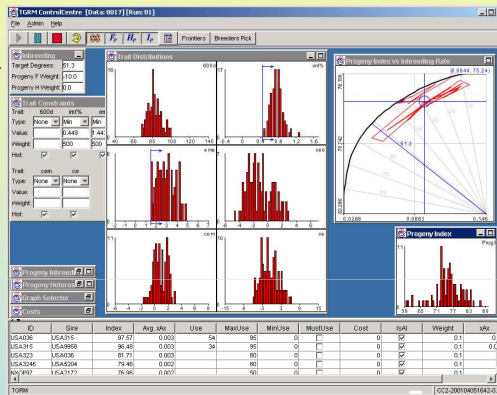
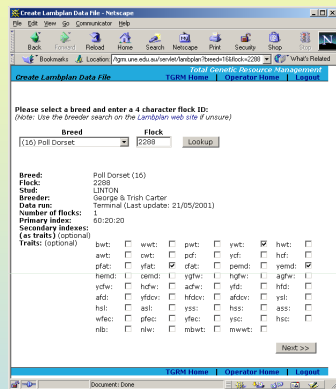
Genetic algorithm for mate selection



Opportunities
Constraints
Attitudes
Issues
Costs



Possible
outcomes



Sire	Dam	index	E	SireCan	DamCan	Sortindex	6000-1	6000-2	inf%	ema-1	ema-2	ced	cem	cc
USA3246	T229	64.68	0.0156	0.003228	0.000138	64.9366	88.00	88.00	0.50	1.15	1.25	-0.70	0.02	
USA3246	T213	70.18	0.0000	0.003228	0.000179	69.9977	81.50	81.50	0.65	1.59	1.59	2.35	0.48	
USA3246	T137	64.68	0.0225	0.003228	0.000000	62.3460	85.00	85.00	0.45	0.20	0.20	2.81	-1.08	
USA3246	T137	65.64	0.0000	0.003228	0.000000	68.4690	72.00	72.00	0.70	0.40	0.40	2.88	1.52	
USA3246	T117	68.09	0.0079	0.003228	0.000087	67.1344	76.00	76.00	0.65	1.45	1.45	2.80	0.81	
USA3246	T663	70.10	0.0029	0.003228	0.000000	75.9360	77.00	77.00	0.90	1.35	1.35	0.89	0.03	
USA3246	T657	73.06	0.0000	0.003228	0.000000	72.8840	70.00	70.00	0.90	-0.55	-0.55	2.74	0.34	
USA3246	T629	64.08	0.0225	0.003228	0.000132	61.9570	77.50	77.50	0.60	0.40	0.40	3.38	-0.97	
USA3246	T620	73.63	0.0078	0.003228	0.000000	74.6760	80.50	80.50	0.65	1.45	1.45	1.87	-0.43	
USA3246	T613	67.58	0.0000	0.003228	0.000137	67.2019	77.50	77.50	0.70	1.20	1.20	1.69	0.48	
USA3246	T608	72.18	0.0000	0.003228	0.000298	71.9982	73.50	73.50	0.75	1.05	1.05	3.04	2.21	
USA3246	T500	62.89	0.0078	0.003228	0.000144	61.7923	84.50	84.50	0.75	1.25	1.25	7.42	-0.71	
USA3246	R061	66.58	0.0000	0.003228	0.000000	64.4090	67.50	67.50	0.80	1.35	1.35	3.78	1.34	
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	
USA3246	Q004	73.62	0.0000	0.003228	0.000000	73.4460	72.00	72.00	0.80	1.30	1.30	2.50	0.36	
USA3246	R211	67.93	0.0000	0.002545	0.000093	67.9594	87.50	87.50	0.95	2.85	2.85	-1.00	-2.26	
USA315	T99	74.46	0.0000	0.002720	0.000116	73.2492	89.50	89.50	0.70	2.00	2.00	0.22	-0.21	
USA315	T270	79.83	0.0000	0.002720	0.000000	78.6263	94.00	94.00	0.80	2.60	2.60	-0.14	-1.15	
USA315	T295	73.26	0.0000	0.002720	0.000114	71.0963	85.50	85.50	0.70	0.95	0.95	-0.46	-0.01	
USA315	T243	76.79	0.0625	0.002720	0.000154	69.3261	88.00	88.00	0.75	2.95	2.95	-1.00	-2.15	
USA315	T217	72.43	0.0000	0.002720	0.000000	71.2263	82.00	82.00	0.70	0.55	0.55	0.49	0.13	

Data construction

TGRM Control Center

Action : A mating list

Internet driven

Internet driven

Internet driven



Genetic evaluation service



TGRM Server
Analysis, monitoring, billing



Breeding operation

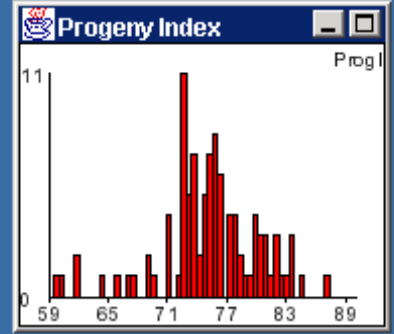
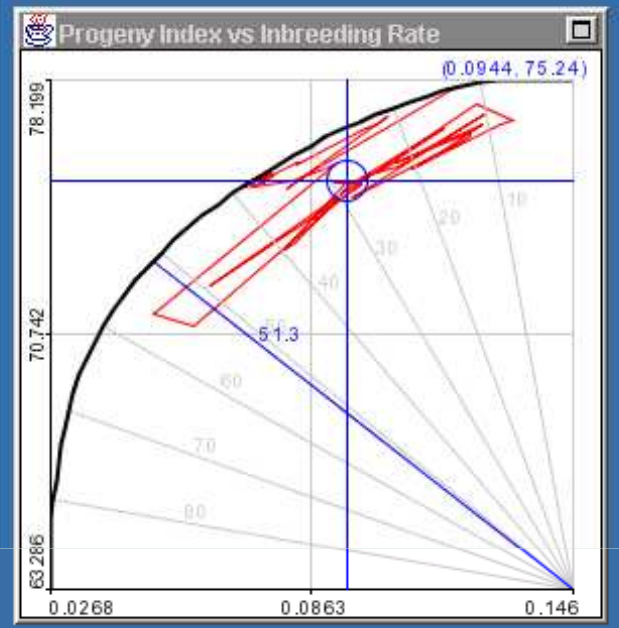
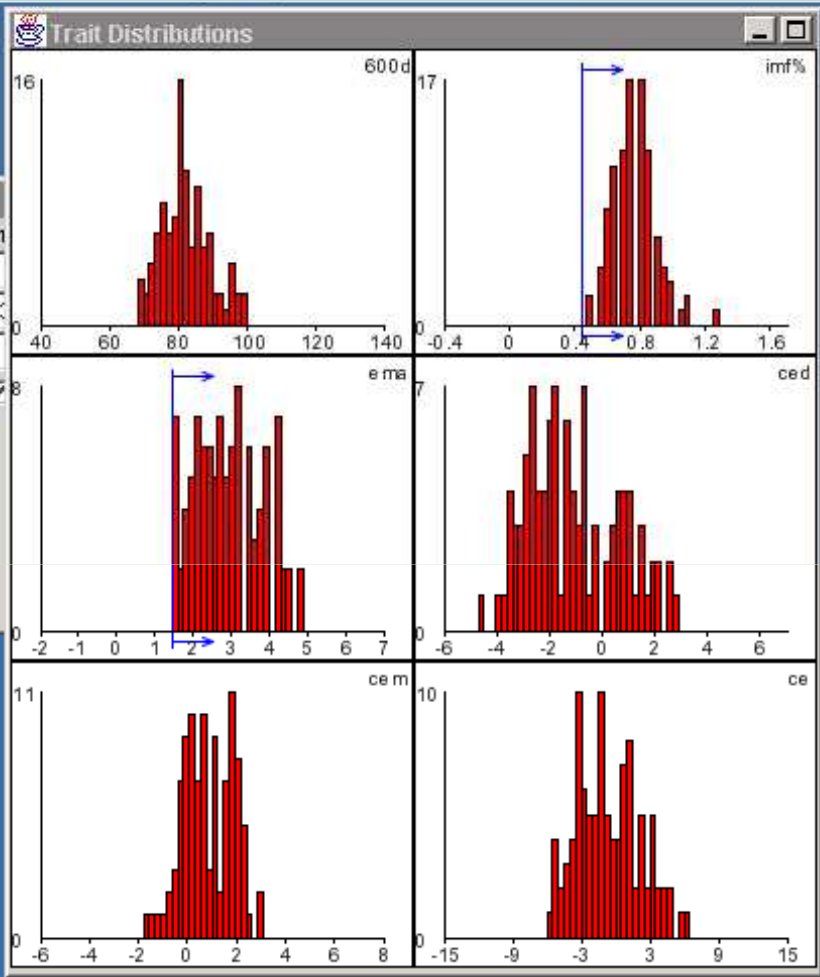
Inbreeding

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:	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>

Trait:	cem	ce
Type:	None	None
Value:		
Weight:		
Hist:	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>



Progeny Inbreed

Progeny Heteros

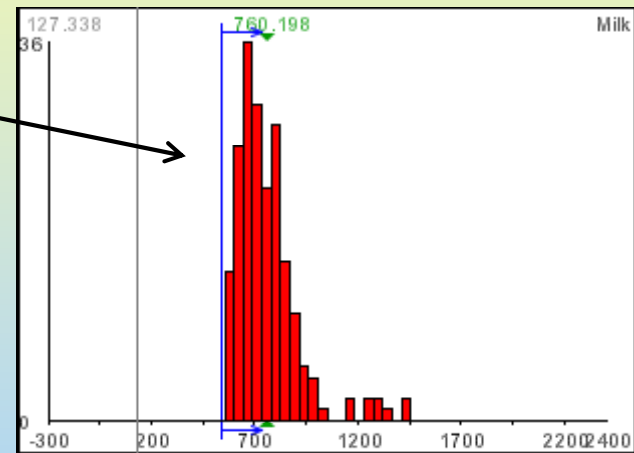
Graph Selector

Costs

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

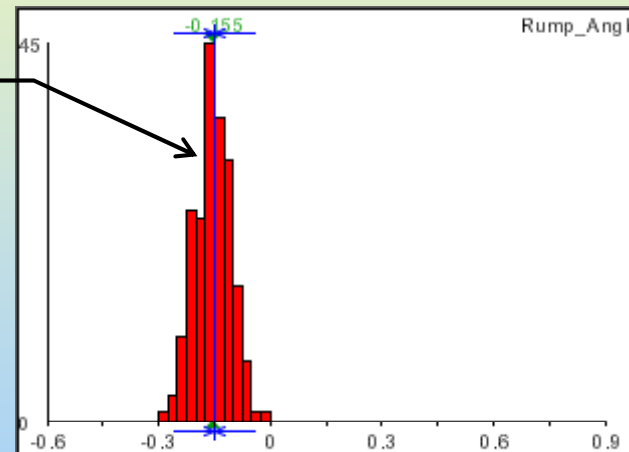
All progeny are predicted to be above the restriction of +545Kg



Predicted progeny Milk EBVs

Achieving Trait Constraints

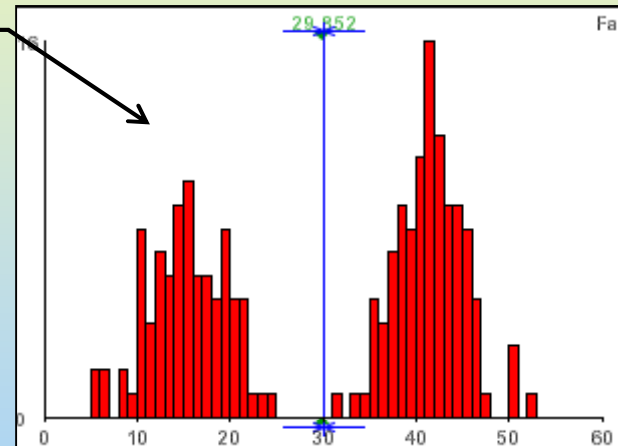
All progeny are predicted to be narrowly distributed about -0.15 degrees



**Predicted progeny
Rump Angle EBVs**

Achieving Trait Constraints

Targeting two different objectives
in one cycle of matings



Predicted progeny Fat EBVs

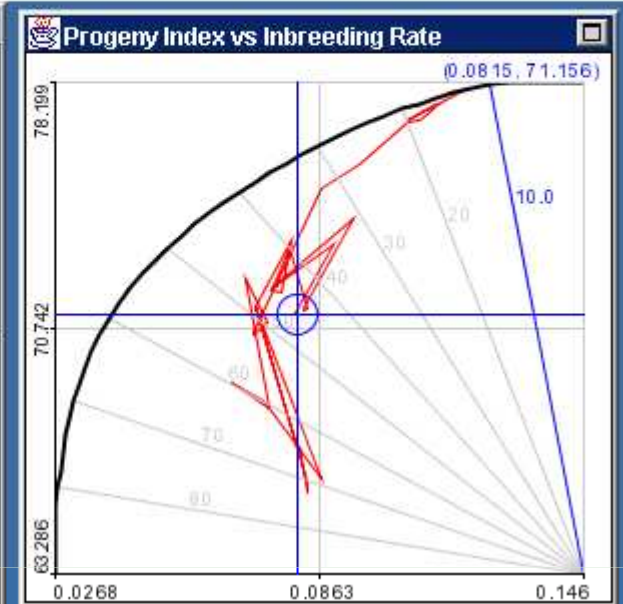
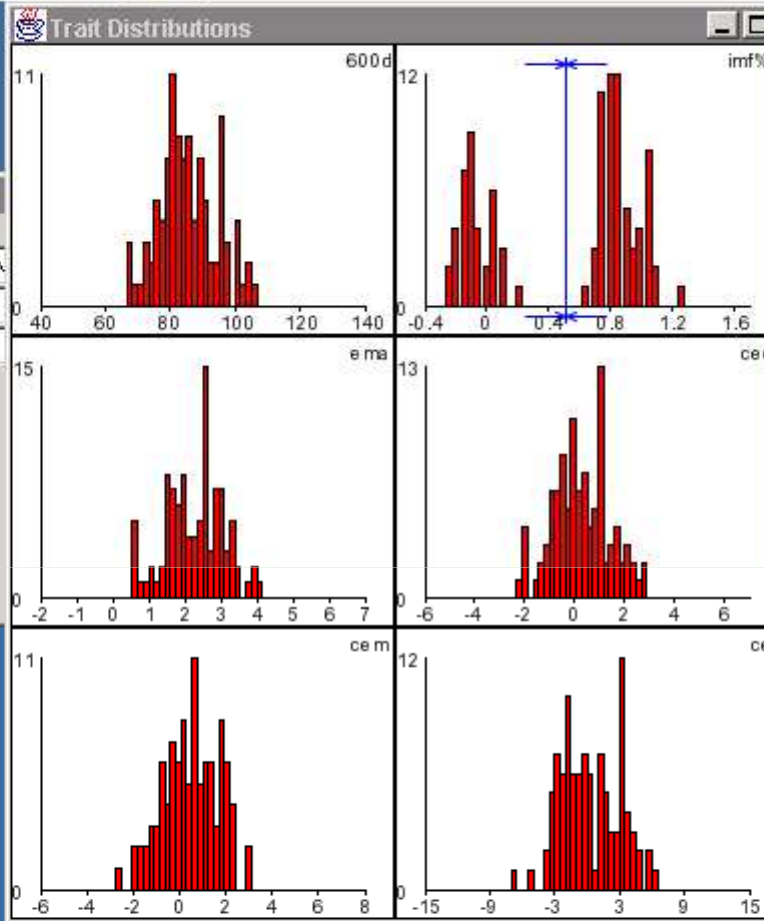
Inbreeding

Target Degrees: 10.0
 Progeny F Weight: -10.0
 Progeny H Weight: 0.0

Trait Constraints

Trait:	600d	imf%	ema
Type:	None	Opt	None
Value:	54.015	0.515	
Weight:	50	-500	
Hist:	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>

Trait:	cem	ce
Type:	None	None
Value:		
Weight:		
Hist:	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>

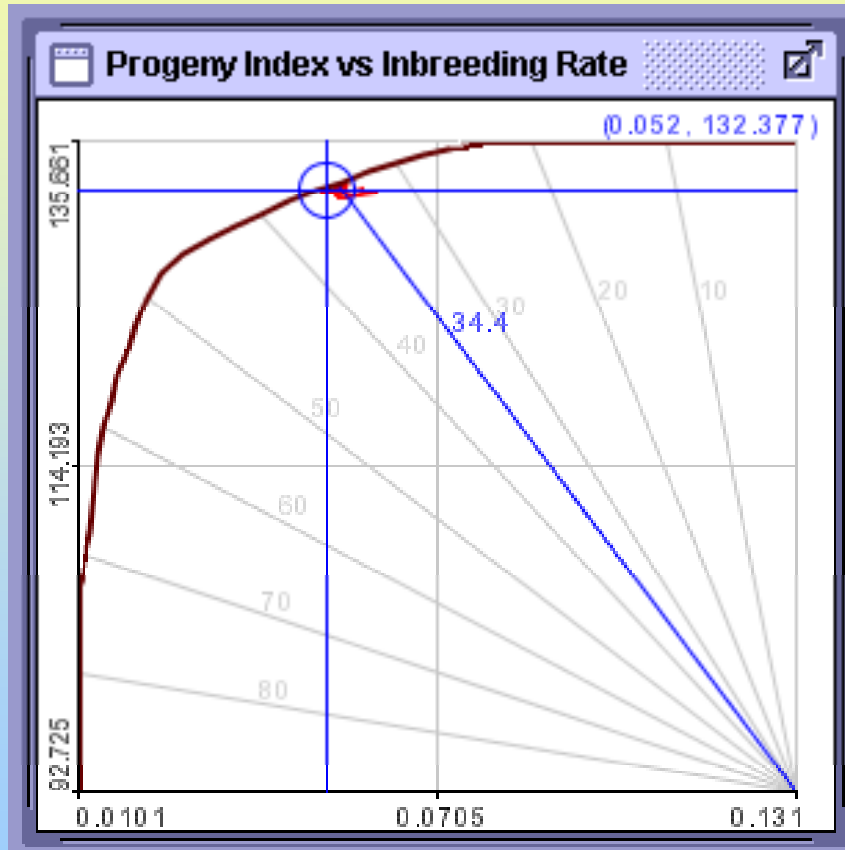


ID	Sire	Index	Avg. xAx	Use	MaxUse	MinUse	MustUse	Cost	IsAI	Weight	xAx
USA315	USA9958	96.48	0.003	62	95	0	<input type="checkbox"/>	0	<input checked="" type="checkbox"/>	0.1	0.051
USA473	0	68.09	0.001	38	80	0	<input type="checkbox"/>	0	<input checked="" type="checkbox"/>	0.1	0.018
USA036	USA315	97.57	0.003		95	0	<input type="checkbox"/>	0	<input checked="" type="checkbox"/>	0.1	
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	

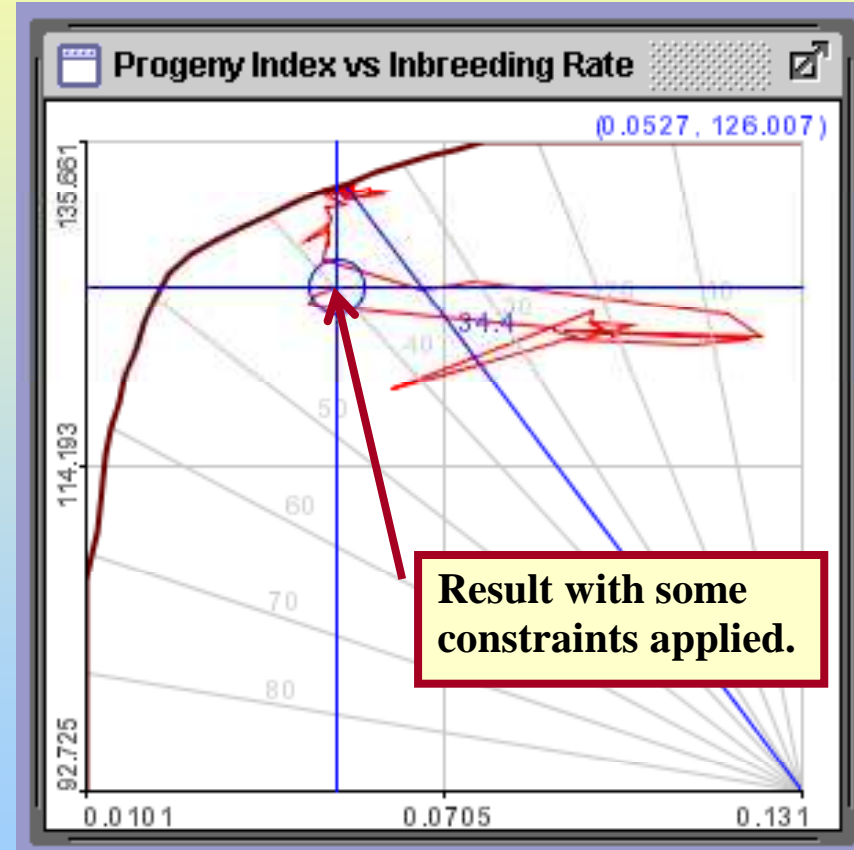
Imposing constraints

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

Predicted progeny index

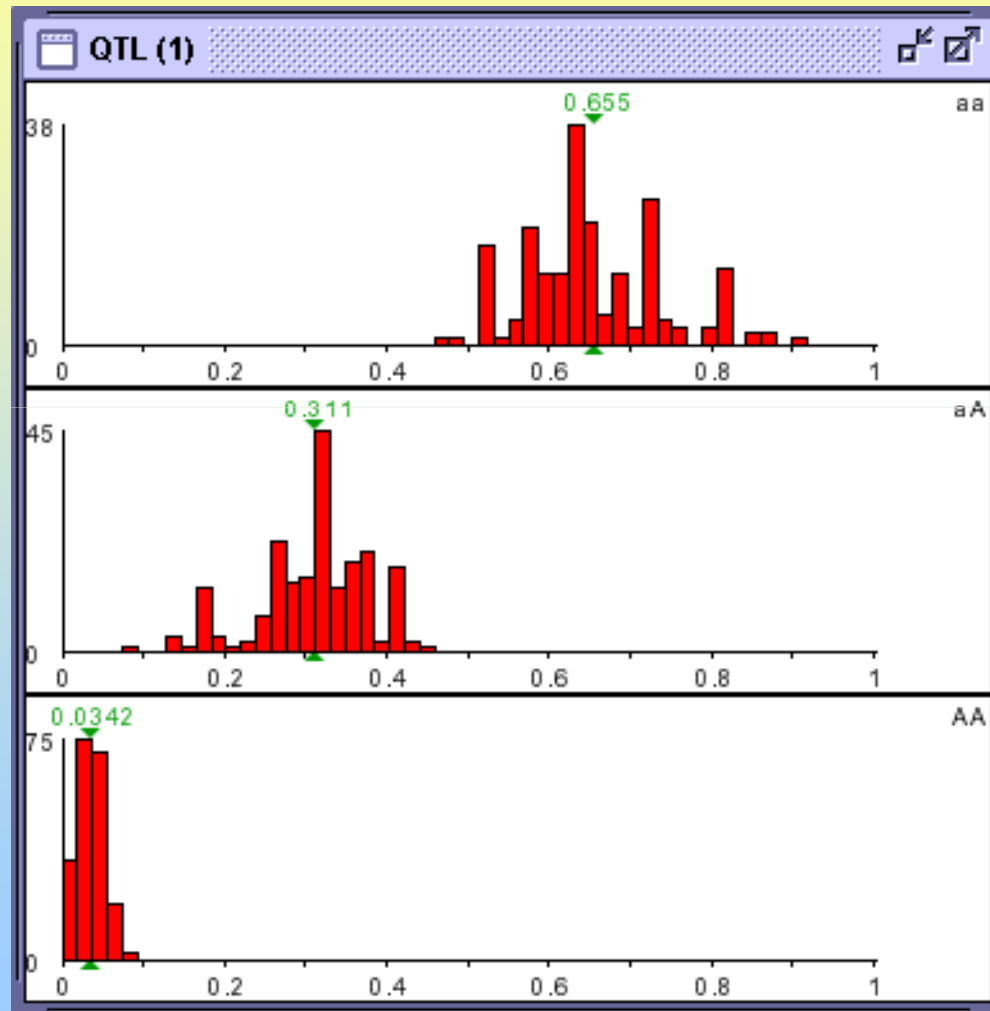


Mean parental coancestry



Mean parental coancestry

MAS: Targetting QTL/gene outcomes



Mating List - Netscape

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Bookmarks Location: http://tgrm.une.edu.au/servlet/reports_show/987823367381_0018_01/matlst0.htm What's Related

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Mating List (by Sire) [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

Document: Done

Use of sires across flocks

Sire	EBV	Sire of Sire	Uses	Flock				
				5001	5002	5003	5004	5008
2350011996961011	148.51	1700171993930114	125	28	48	0	13	36
2350011996961017	143.62	1700171993930114	44	0	0	0	0	44
2350021996962003	146.45	1700171993930114	89	0	24	0	0	65
2350021996962048	138.64	1600341994940171	33	0	0	0	0	33
2350031994943086	131.72	0	71	0	8	40	23	0
2350031995953035	130.58	0	57	0	22	0	35	0
2350031996963053	151.18	1700171993930114	113	0	19	54	0	40
2350031996963054	144.70	1700171993930114	38	0	0	0	0	38
2350031996963057	140.70	1600341994940171	90	0	10	19	0	61
2350031996963071	146.24	1700171993930114	83	0	45	18	20	0
2350031997973113	131.09	1618901992920099	38	0	0	38	0	0
2350031997973201	142.59	2350031996963059	33	0	0	33	0	0
2350031997973256	153.40	2350031996963057	37	0	0	37	0	0
2350041996964066	134.92	2350021994942107	73	11	0	62	0	0
2350041996964183	133.67	2350021994942107	70	0	20	28	6	16
2350081996960176	133.76	1600341994940171	33	0	0	33	0	0
TOTALS:			1027	39	196	362	97	333

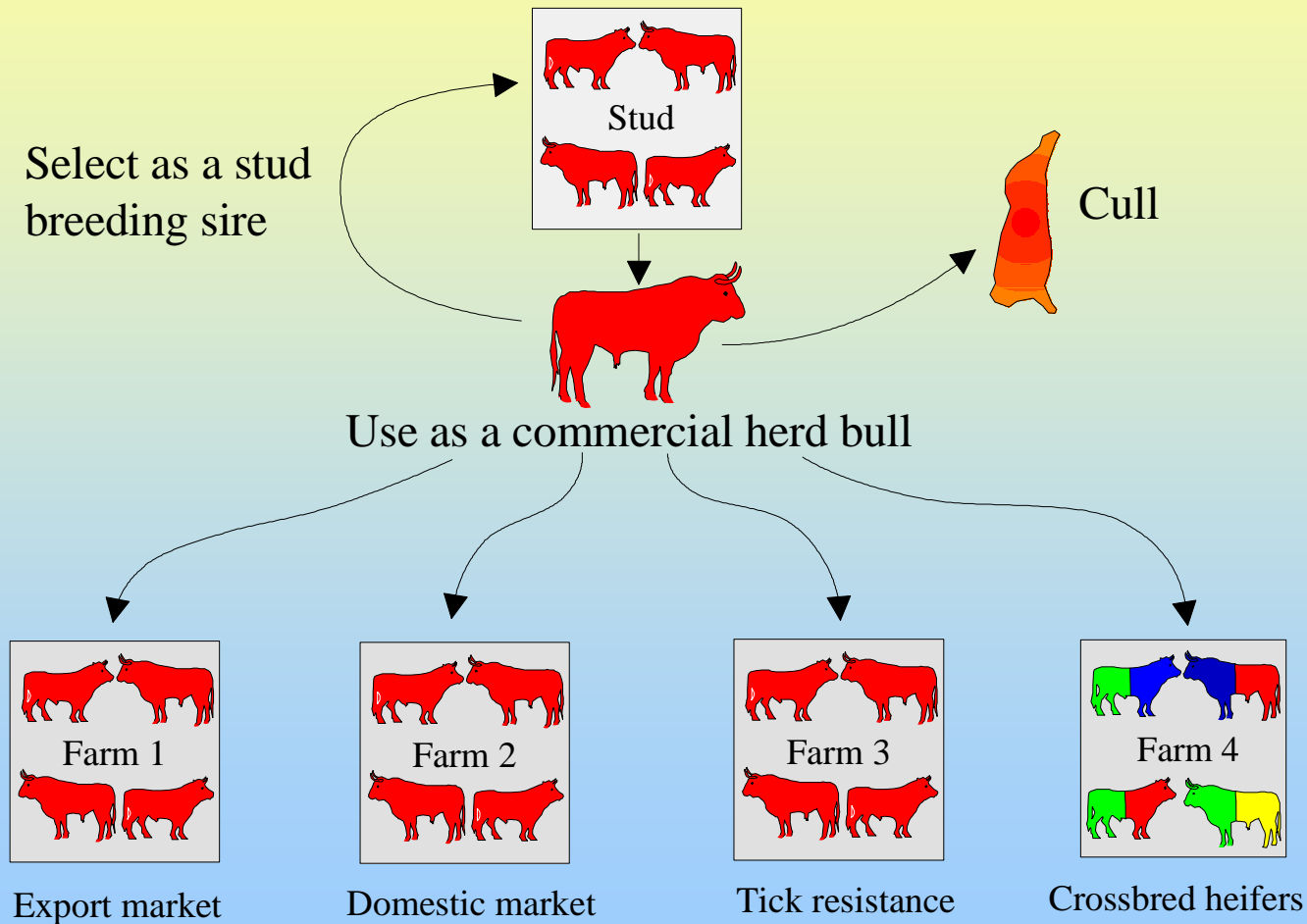
Examples of Costs that can be accommodated

- Reproductive costs.
 - AI, MOET, JIVET
 - find breakeven costs
- Seedstock purchase costs
 - rationalise the balance between price and value
- Seedstock maintenance costs
 - optimise herd size
- Limits on funds available

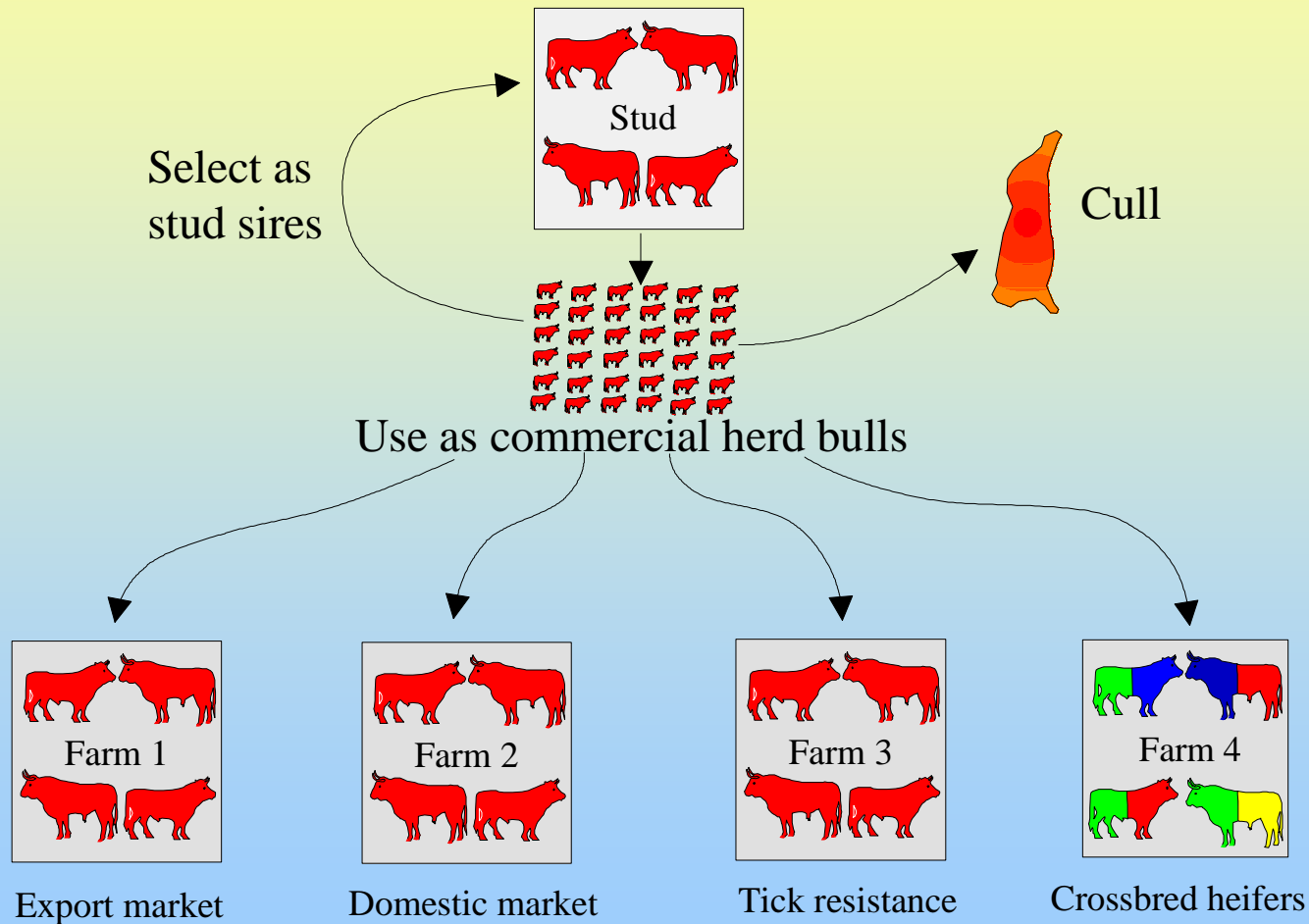
Examples of other factors that can be controlled.

- **Mating logistics**
 - number and size of mating paddocks
 - “don’t migrate young rams”
 - “put young ewes only to old rams”
- **Opportunities and constraints**
 - “Lets consider using this New Zealand sire”
 - “I want to use all my 50 doses of this ram’s semen”
 - “Use this link sire for at least 25 matings”
 - “Don’t use any one bull for less than 25 matings”
- **Desired outcomes**
 - narrow variation in birthweight, fat, etc
 - corrective matings

The fate of a stud-born bull ...



The fate of many stud-born bulls ...



Breeding program

