

Crossbreeding

Reasons for crossbreeding

1. Sire-Dam complementation
2. The averaging of breed effects
3. Grading up to a new breed
4. Step towards creation of synthetic/composite
5. To introduce a single gene
6. To exploit heterosis

An example of the value of selection between breeds :

- 25 *Bos Taurus* breeds - 71% of the variation in 72-week weight was due to breed effects (Thiessen et. al., 1984). Standard deviation within breeds was 33.5Kg and standard deviation between true breed means was 52.4Kg

- This gives the percentage variation that is due to breed:

$$100 \times \frac{52.4^2}{52.4^2 + 33.5^2} = 71\%$$

- Choose a breed just one standard deviation 'better' - 52.4 Kg improvement
- Same as moving from eg. the average breed (50% standing) up to the breed with 84% standing, on a 0% (worst breed) to 100% (best breed) scale.

An example of the value of selection within breeds :

Age at calving:	2	3	4	5	6
Bulls:	1	1			
Cows:	8	8	8	8	8

40 Cows x 0.8 gives 16 males + 16 females per year cohort.

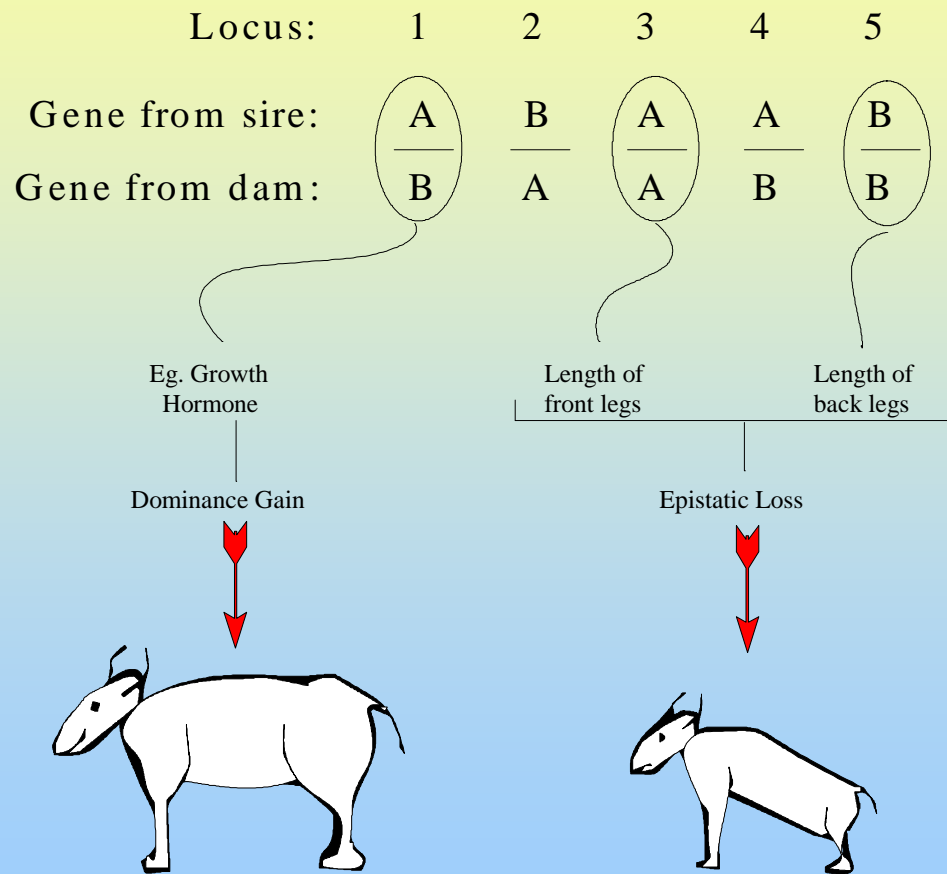
$$\begin{array}{lll}
 p_m = 1/16 & \text{giving} & i_m = 1.970 \\
 p_f = 8/16 & \text{giving} & i_f = 0.798 \\
 L_m = 2.5 & \text{and} & L_f = 4
 \end{array}$$

$$R_{yr} = \frac{i_m + i_f}{L_m + L_f} h^2 \sigma_P = \frac{1.970 + 0.798}{2.5 + 4} \times 0.4 \times 33.5 = 5.7 \text{ Kg / yr}$$

The comparison :

- Between breeds: 52.4Kg
- Within breeds: 5.7Kg/yr
- Our selection between breeds is worth 9 years 2 months and 6 days selection work with breed!
- Selection between breeds is worth considering - depending on costs and opportunities.

Dominance and epistasis cause heterosis



DOMINANCE - wider genetic base leads to better performance

EPISTASIS - breakdown of favourable interactions leads to loss of performance

Dominance model of heterosis

pair	one gene pair	another gene
Purebreed "A"	Genes from sire: A A A A A A A A Genes from dam : A A A A A A A A	
	Heterosis expression = 0%	
F ₁ cross "A x B"	Genes from sire: A A A A A A A A Genes from dam : B B B B B B B B	
	Heterosis expression = 100%	

Dominance model of heterosis

3 breed cross

"C x (AxB)"

Genes from sire: C C C C C C C C

Genes from dam : A B A B A B A B

Heterosis expression = 100%

Backcross

"A x (AxB)"

Genes from sire: A A A A A A A A

Genes from dam : A B A B A B A B

Heterosis expression = 50%

F₂ cross

"(AxB) x (AxB)"

Genes from sire: A A B B A A B B

Genes from dam : A B A B A B A B

Heterosis expression = 50%

	How much heterosis?
Purebreds	0
F1	100%
F2	50%
Backcross	50%

Breed-of-origin heterozygosity

\propto

allelic heterozygosity

Breed X

$p_x=0.3$

$q_x=0.7$

$p_x=0.3$.09 a	d	.21
$q_x=0.7$	d	-a	.49

$$\text{Breed X} = -0.4a + 0.42d$$

Breed Y

$p_y=0.9$

$q_y=0.1$

$p_y=0.9$.81	.09
$q_y=0.1$.09	.01

$$\text{Breed Y} = 0.8a + 0.18d$$

Parental Mean:

$$\bar{P} = 0.2a + 0.3d$$

Breed-of-origin heterozygosity

∞

allelic heterozygosity

F1 Cross

$p_x=0.3$ $q_x=0.7$

	$p_y=0.9$.27	.63
	$q_y=0.1$.03	.07

F1 Cross = $0.2a + 0.66d$

Heterosis = $.36d = d(p_x - p_y)^2$

Breed-of-origin

Heterozygosity = 1

F2 Cross

0.6 0.4

	$\frac{p_x+p_y}{2} = 0.6$.36	.24
	$\frac{q_x+q_y}{2} = 0.4$.24	.16

F2 Cross = $0.2a + 0.48d$

Heterosis = $.18d = \frac{1}{2}d(p_x - p_y)^2$

Breed-of-origin

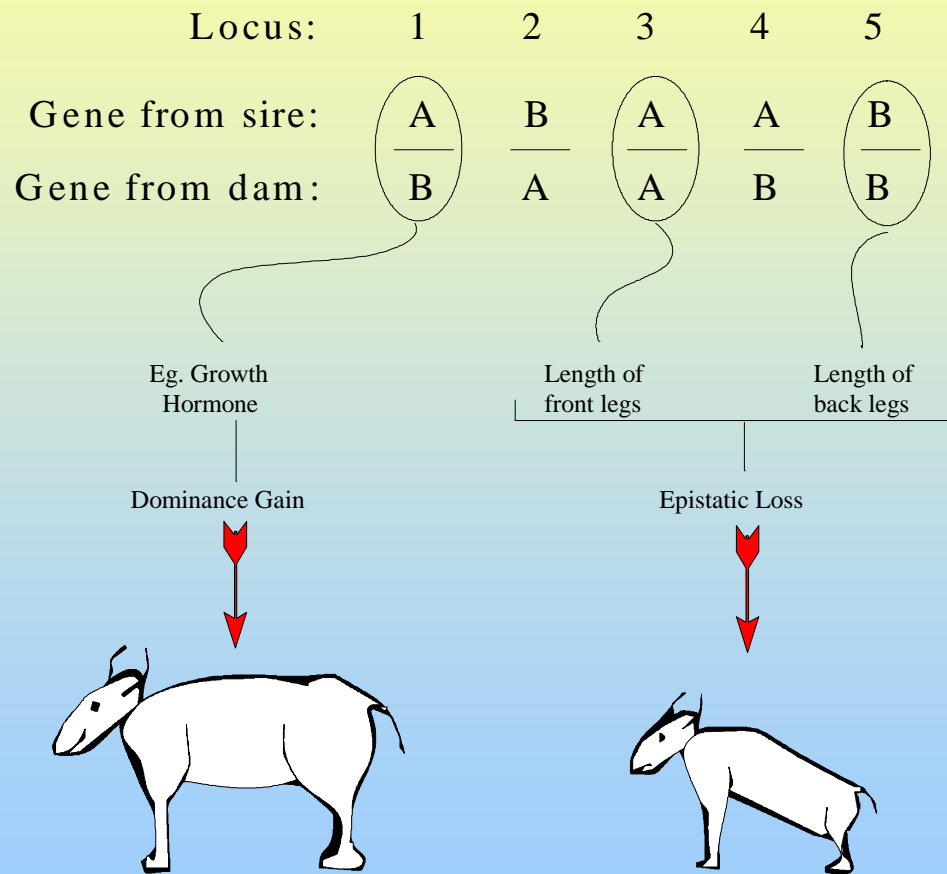
Heterozygosity = $\frac{1}{2}$

Epistasis

Material in book is

For reference only !

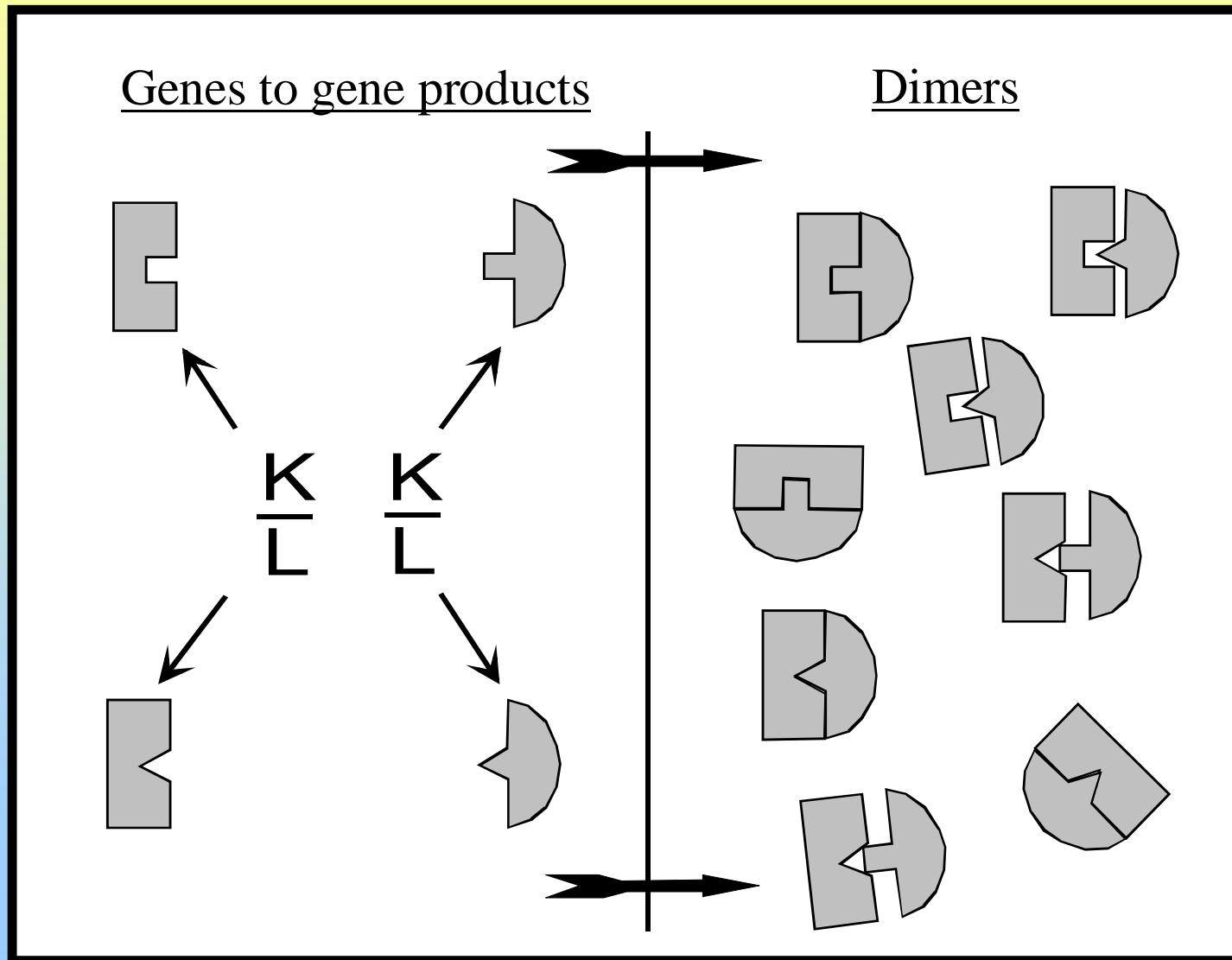
Dominance and epistasis cause heterosis



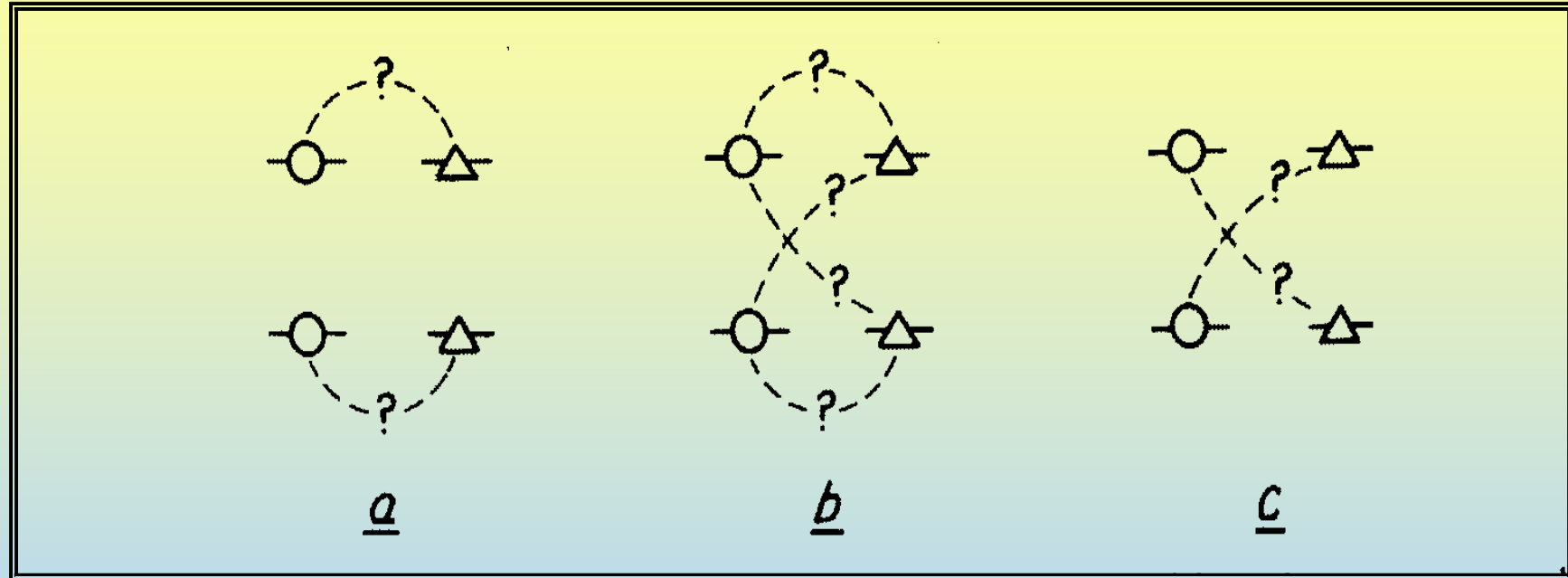
DOMINANCE - wider genetic base leads to better performance

EPISTASIS - breakdown of favourable interactions leads to loss of performance

Additive x additive model of heterosis



Where 'recombination loss' fits in



Recombination
loss, r

Additive x additive
epistasis, E_{aa}

The difference,
equal to dominance, d

$$P(E_{aa}) = [p(r) + p(d)]/2$$

Crossbreeding parameters...

- **Direct additive effects** A_{d1} , A_{d2} and A_{d3} Additive effects of purebreeds. For yearling weight, they relate to the ability to grow quickly.
- **Maternal additive effects** A_{m1} , A_{m2} and A_{m3} Additive effects of purebreeds as expressed by the dams of the crossbred individuals under consideration. They probably relate to milk production and rearing ability. Note that these effects add to zero - they describe the relative maternal performance of each pure breed.
- **Direct dominance effect** D_d The effect of heterosis in crossbred individuals, when fully expressed, as in an F1 cross.
- **Maternal dominance effect** D_m The effect of heterosis due to crossbreeding in the dam, when fully expressed, as in an F1 dam.

Estimating crossbreeding parameters...

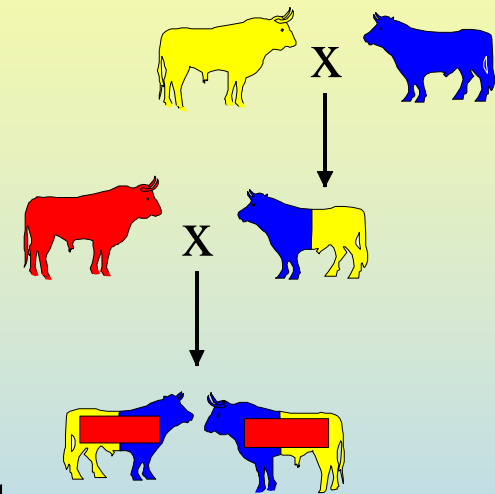
Cross Table (Yearling weight)															
3 Breeds		New Row		Reset Table		Breed / Trait		Update		Defaults		Close		Calculate	
Effects:		<input checked="" type="checkbox"/> Ad1 Ad2 Ad3			<input checked="" type="checkbox"/> Am1 Am2 Am3			<input checked="" type="checkbox"/> Dd		<input checked="" type="checkbox"/> Dm		Merit			
Value (Kg)		300	280	260	-6	-1	7	20	10						
1	1 x 1	1	0	0	1	0	0	0	0	294	<input checked="" type="checkbox"/>				
2	2 x 2	0	1	0	0	1	0	0	0	279	<input checked="" type="checkbox"/>				
3	3 x 3	0	0	1	0	0	1	0	0	267	<input checked="" type="checkbox"/>				
4	1 x 2	0.5	0.5	0	0	1	0	1	0	309	<input checked="" type="checkbox"/>				
5	1 x 23	0.5	0.25	0.25	0	0.5	0.5	1	1	318	<input checked="" type="checkbox"/>				
6	1 x 12	0.75	0.25	0	0.5	0.5	0	0.5	1	311.5	<input checked="" type="checkbox"/>				
7	2 Br Bal Comp	.5	.5	0	.5	.5	0	.5	.5	301.5	<input checked="" type="checkbox"/>				
8	3 Br Bal Comp	.3333	.3333	.3333	.3333	.3333	.3333	.667	.667	299.982	<input type="checkbox"/>				
9	2 Br Opt Comp	.63	.37	0	.63	.37	0	.47	.47	302.55	<input type="checkbox"/>				
10	3 Br Opt Comp	.57	.31	.12	.57	.31	.12	.56	.56	302.91	<input type="checkbox"/>				
11	2 Br Rotation	.5	.5	0	.5	.5	0	.667	.667	306.51	<input type="checkbox"/>				
12	3 Br Rotation	.3333	.3333	.3333	.3333	.3333	.3333	.86	.86	305.772	<input type="checkbox"/>				

Crossbreeding:

More 'structure' gives more merit ...

In general ...

The shorter the breed pedigree
back to purebred parents:

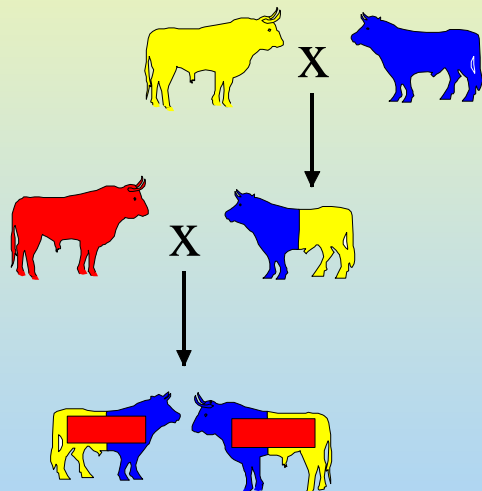


- the more heterosis can be expressed.
- the more sire-dam complementarity can be expressed

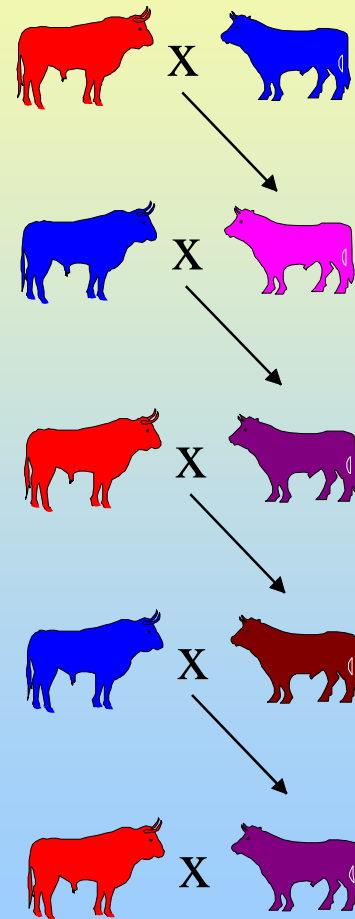
BUT: The more expensive the operation is to run

Loss of heterosis and complementarity ...

3-Breed Cross

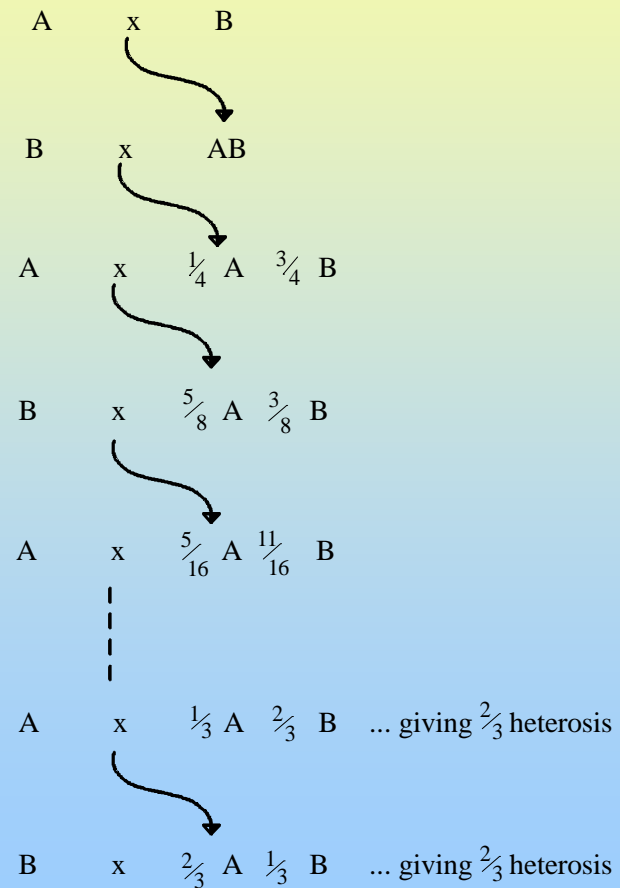
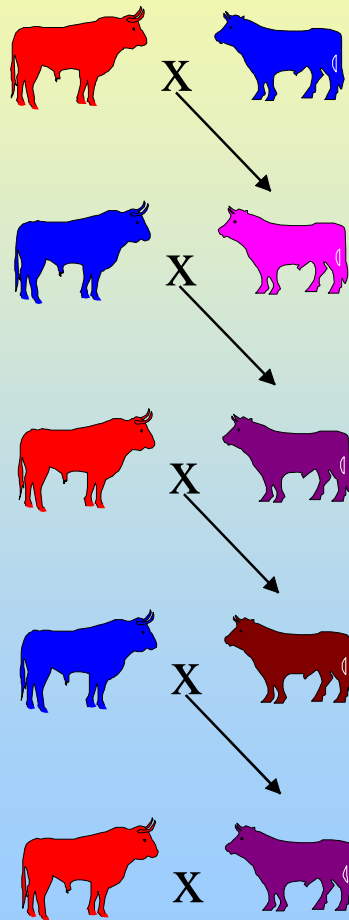


Rotational Cross



Loss of heterosis and complementarity ...

Rotational Cross



Which crossing system to adopt?

PUREBREED	when no cross is better.
F ₁ CROSS	when direct heterosis is important.
3 BREED CROSS	when both direct and maternal heterosis are important.
4 BREED CROSS	when paternal heterosis is important as well.
BACKCROSS	when only 2 good parental breeds are available and/or when direct heterosis is not important.
ROTATIONAL CROSSES	when females are too expensive to either buy in or to produce in the same enterprise.
OPEN OR CLOSED COMPOSITE	when both males and females are too expensive. A few initial well judged importations establish the synthetic (or 'composite'), and it can then either be closed (which helps to establish a breed 'type'), or left open to occasional well judged importations.

Patterns of use of crossbreeding


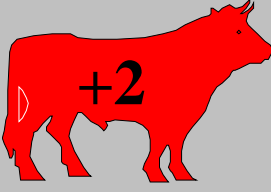
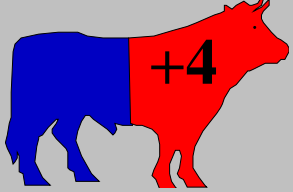
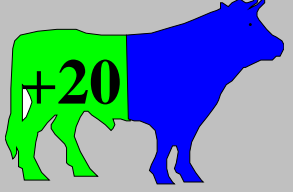
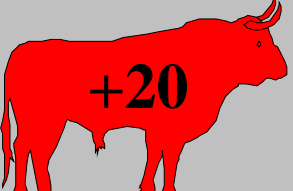
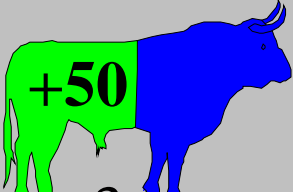
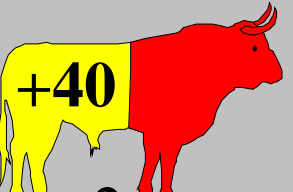
Industry	Fecundity	Typical crossing systems
Poultry	highest ↓ lowest	4-breedcrosses
Pigs		3-breed crosses;back crosses
Meat sheep		3-breedcrosses
Wool Sheep		purebred*
Dairy		purebred*
Temperate Beef		rotations;composites
Tropical Beef		composites

*Wool sheep and dairy industries are exceptions due to availability of an outstanding pure breed in each.

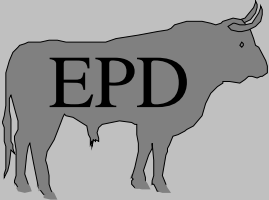
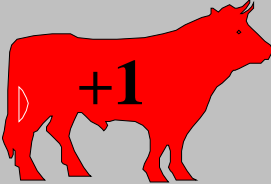
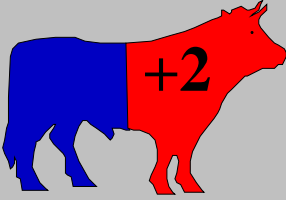
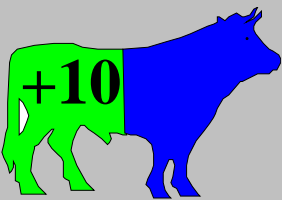
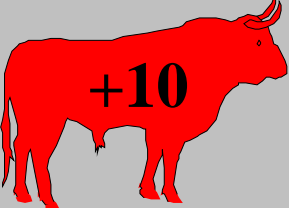

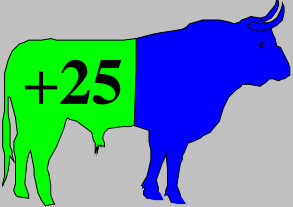

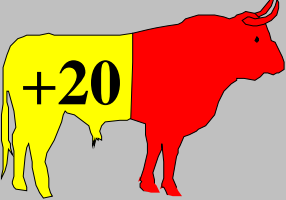

Exploiting both between and within breed variation

- Designing crossbreeding systems based on breed means
- Actually selecting animals based on within breed EBVs
 - E.g. mate merinos with the best BL ram
 - Use the best Angus bulls for rotational crossing
 - Etc.
- Could also use across breed EBV (not always) but that does not exploit heterosis

'Automatic design' through mate selection.

 EBV Cost	 +2 0	 +4 10	 +20 10
 +20 0	Crossing: 300 EBV: 11 Cost: 0 Total: 311	Crossing: 310 EBV: 12 Cost: -10 Total: 312	Crossing: 325 EBV: 20 Cost: -10 Total: 335
 +50 3	Crossing: 318 EBV: 26 Cost: -3 Total: 341	Crossing: 308 EBV: 27 Cost: -13 Total: 322	Crossing: 300 EBV: 35 Cost: -13 Total: 322
 +40 2	Crossing: 290 EBV: 21 Cost: -2 Total: 309	Crossing: 320 EBV: 22 Cost: -12 Total: 330	Crossing: 327 EBV: 30 Cost: -12 Total: 345

Mate allocations ...

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

Linear Programming approach

Jansen and Wilton (1985)

Bull→ Cow↓	1	2	3	Dummy bull
1	311	312	335	999
2	341	322	322	999
3	309	330	345	999
Dummy Cow	999	999	999	999

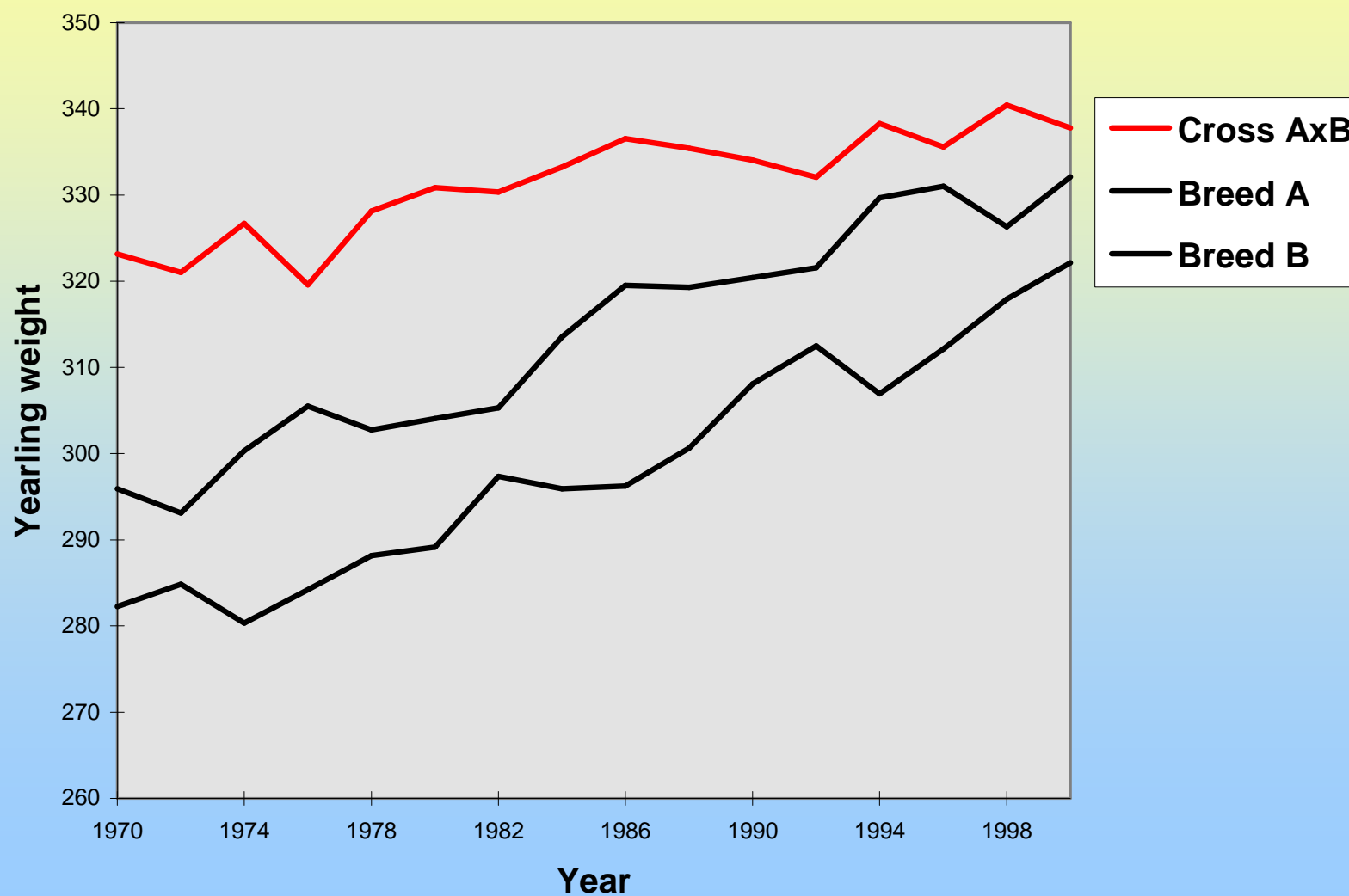
Maximise sum of merit of chosen cells, with constraints such as:

- Max. one mating per cell
- Max. one mating per female
- Max. d matings per male

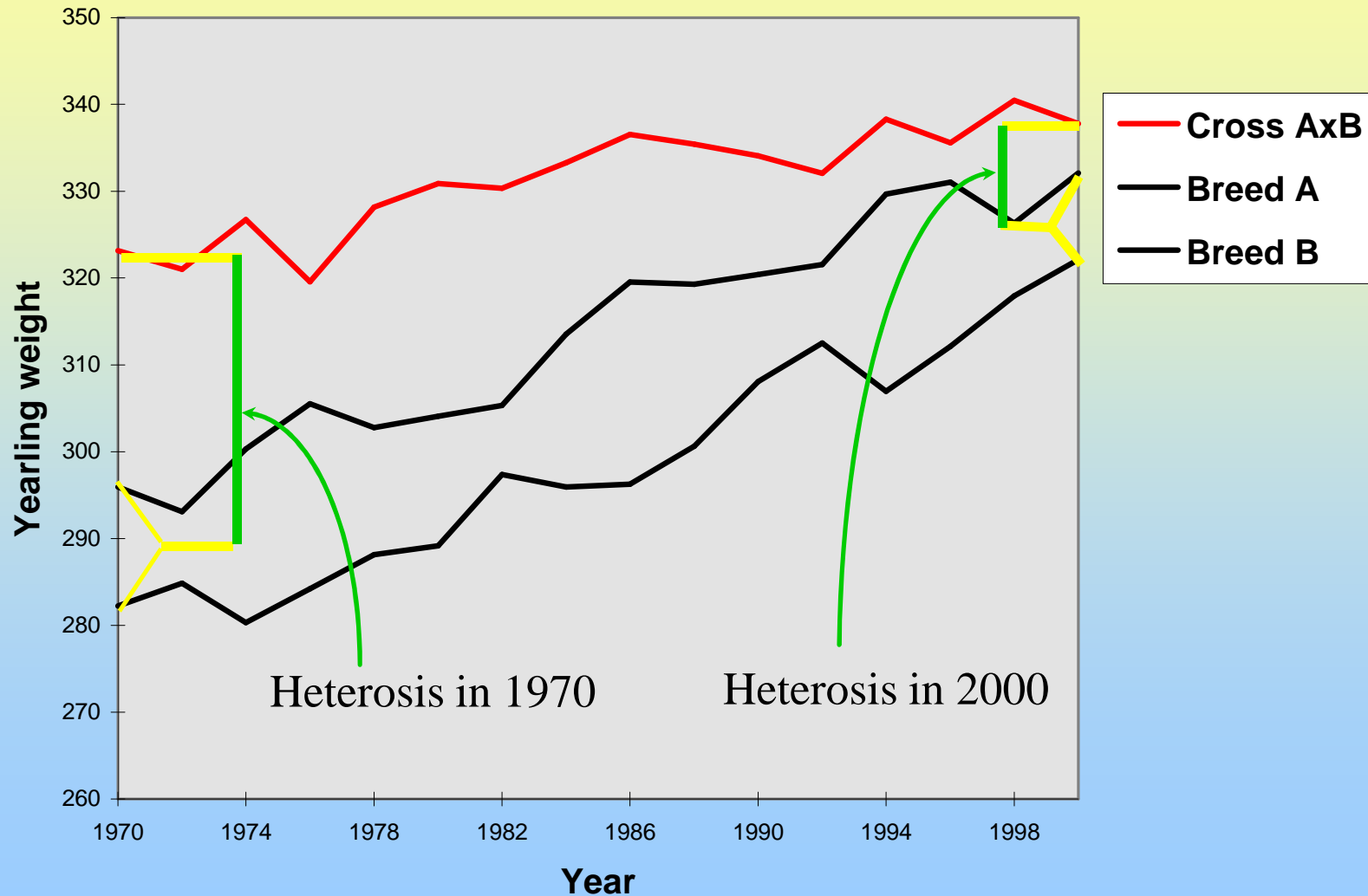
Does bull ranking depend on mate breed?

EPD's	Angus Bull No. 1	Angus Bull No. 2	Angus Bull No. 3
Angus cows	+++	++	+
Hereford cows	+	++	+++
Brahman cows	-	++	+

If ‘Pure-Cross correlation’ were less than 100% ...



If “Pure-Cross correlation” were less than 100% ...



Conclusion about crossbreeding

- Design optimal structure based on breed parameters (averages)
- Operationally: select and mate based on EBVs and predicted heterosis
- Non-additive effects not important for selection within lines or breeds