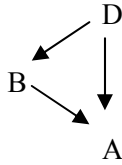


Genetic relationships and close inbreeding - solutions

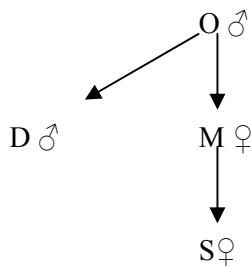
1. a)



$$F_A = \frac{1}{2} a_{BD} = \frac{1}{2} \times \left(\left(\frac{1}{2} \right)^{1+0} (1 + F_D) \right) = \frac{1}{2} \times \left(\frac{1}{2} \times (1 + 0) \right) = \frac{1}{4} = 0.25$$

$$b) \quad a_{AD} = \left(\left(\frac{1}{2} \right)^{2+0} + \left(\frac{1}{2} \right)^{1+0} \right) (1 + F_D) = \left(\left(\frac{1}{2} \right)^2 + \left(\frac{1}{2} \right)^1 \right) (1 + 0) = \frac{1}{4} + \frac{1}{2} = 0.75$$

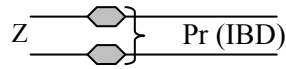
2. a)



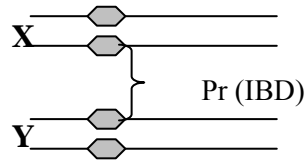
$$a_{DS} = \left(\frac{1}{2} \right)^{1+2} (1 + F_o) = \left(\frac{1}{2} \right)^3 = \frac{1}{8}$$

$$b) \quad F_{\text{progeny}} = \frac{1}{2} a_{DS} = \frac{1}{2} \times \frac{1}{8} = \frac{1}{16} = 0.625 = 6.25\%$$

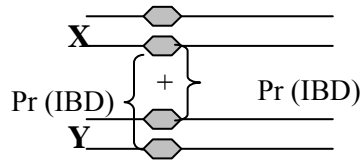
3. a) F_Z = The probability that two alleles in a locus of an individual Z are identical by descent (IBD)



- b) f_{xy} = The probability that a randomly sampled gene from individual X is identical by descent with one randomly sampled gene of the same locus from individual Y



- c) a_{xy} = The probability that a randomly sampled gene from individual X is identical by descent with any of the genes in the same locus from individual Y



- d) $F_Z = f_{xy} = \frac{1}{2} a_{xy}$

4. D is inbred: $F_D = \frac{1}{2} a_{BA} = \frac{1}{4}$

Three paths must be summarized:

$$a_{EF} = \left(\frac{1}{2}\right)^{3+2}(1+0) + \left(\frac{1}{2}\right)^{2+2}(1+0) + \left(\frac{1}{2}\right)^{1+1}\left(1+\frac{1}{4}\right)$$

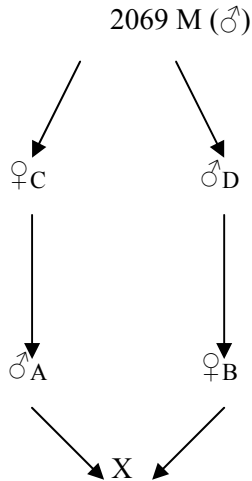


$$= \left(\frac{1}{2}\right)^5 + \left(\frac{1}{2}\right)^4 + \left(\frac{1}{2}\right)^2 \times \frac{5}{2} = \frac{1}{32} + \frac{1}{16} + \frac{5}{16} = \frac{13}{32}$$

$$F_X = \frac{1}{2} a_{EF} = \frac{1}{2} \times \frac{13}{32} = \frac{13}{64} \approx 0.20$$

5.

a)



The inbreeding coefficient of the progeny (F_X) equals half the additive relationship (a_{AB}) between parents A and B

$$a_{AB} = \left(\frac{1}{2}\right)^{m+n} (1 + F_{2069M})$$

where m and n are the number of arrows between A, B and the common ancestor (2069 M)

Thus

$$a_{AB} = \left(\frac{1}{2}\right)^{2+2} (1+0) = \frac{1}{16} = 0.0625 \text{ or } 6.25\%.$$

The inbreeding coefficient for X is

$$F_X = \frac{1}{2} \times a_{AB} = \frac{1}{2} \times \frac{1}{16} \times \frac{1}{32} = 0.03125 \text{ or } 3.125\%.$$

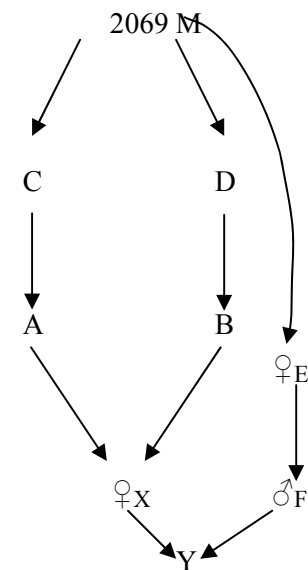
Here we assumed that the ancestor (2069 M) is not inbred.

If e.g. the inbreeding coefficient for 2069 M is 5%, then

$$a_{AB} = \left(\frac{1}{2}\right)^{2+2} (1+0.05) = 0.0625 \times 1.05 = 0.066 \text{ or } 6.6\% \text{ and}$$

$$F_X = 3.3\%$$

b)



The relationship between the parents of Y is

$$a_{XF} = \left(\left(\frac{1}{2}\right)^{3+2} + \left(\frac{1}{2}\right)^{3+2} \right) \times (1 + F_{2069M})$$

i.e. both paths X, A, C, (2069 M), E, F and X, B, D, (2069 M), E, F are summarized

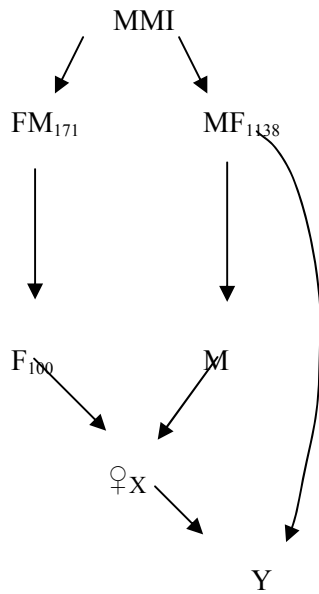
If F_{2069M} is 0 (2069 M not inbred)

$$a_{XF} = \frac{1}{32} + \frac{1}{32} = \frac{1}{16} = 0.0625 = 6.25\%$$

The inbreeding coefficient for Y is then

$$F_Y = \frac{1}{2} \times \frac{1}{16} = \frac{1}{32} = 0.03125 = 3.125\%$$

c)



The heifer X is a grand-daughter to 1138 S and furthermore related to him through 2069 M. The relationship between them is

$$a_{X-1138S} = \left(\frac{1}{2}\right)^{2+0}(1+F_{1138S}) + \left(\frac{1}{2}\right)^{3+1}(1+F_{2069M}).$$

If 1138 S and 2069 M are not inbred, i.e. F_{1138S} and $F_{2069M} = 0$ then

$$\begin{aligned} a_{X-1138S} &= \left(\frac{1}{2}\right)^{2+0} + \left(\frac{1}{2}\right)^{3+1} = \frac{1}{4} + \frac{1}{16} = \\ &= \frac{5}{16} = 0.3125 = 31.25\%. \end{aligned}$$

The inbreeding coefficient for an offspring, if the heifer X is inseminated by 1138 S would be:

$$F_Y = \frac{1}{2} a_{X-1138S} = \frac{1}{2} \times \frac{5}{16} = \frac{5}{32} = 0.15625 = 15.6\%.$$

d) 15% is definitely too high to be recommended