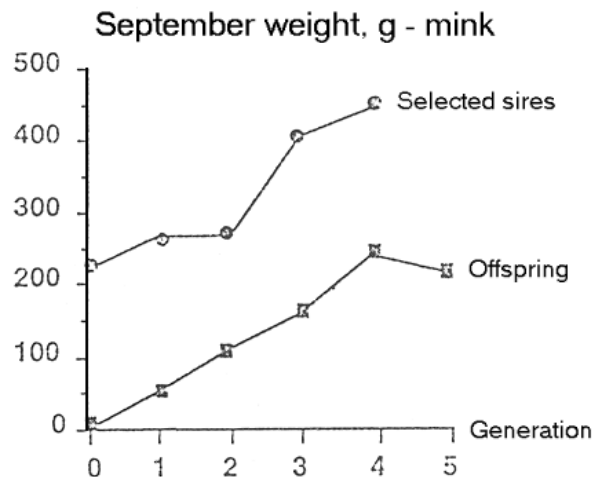


6.4 Estimating the heritability and common environmental effect

Selection experiment for estimation of heritability.

In Figure 6.5 is shown a selection experiment for September weight in mink, carried out by G. Lagerquist, the Swedish Agric. Univ. published in J. Am. Sci., 1993, 71:3261-72. Only the males were selected with S equal to around 225 gram in each generation - the upper curve in the figure. The lower curve in the figure shows the changes for each generation, R , that is equal to around 50 gram per generation. The heritability is estimated by $R = h^2 S/2$, as only the males were selected: and inserted, $50 = h^2 * 225/2$ which correspond to h^2 equal to 0.44.



$R = h^2 * S$ can also be interpreted as the regression of offspring on the average of the two parents, where the regression coefficient b correspond to h^2 .

Heritability estimation based on calculated correlation. The use of a selection experiment for estimating the heritability takes long time, at least several generations. Now will be presented methods for estimation of the heritability based on data from one generation only.

The heritability (h^2) can be estimated as the calculated correlation (r or t) between related individuals divided by the coefficient of relationship (a). This is correct if inheritance (genes) is the only cause of similarity between the involved individuals. The formulas are as follows:

$$h^2 = r/a, \text{ or } r = a * h^2.$$

The calculation of the correlation can be done according to the classical method (see Figure 6.4). When considering several individuals in a group, as for instance half sibs, calculations are done by means of analysis of variance and estimation of the interclass correlation.

Estimation of the heritability can be based on full and half sib material as can be found in all multipare species. The heritability measures what degree of similarity exists between related animals sharing identical genes.

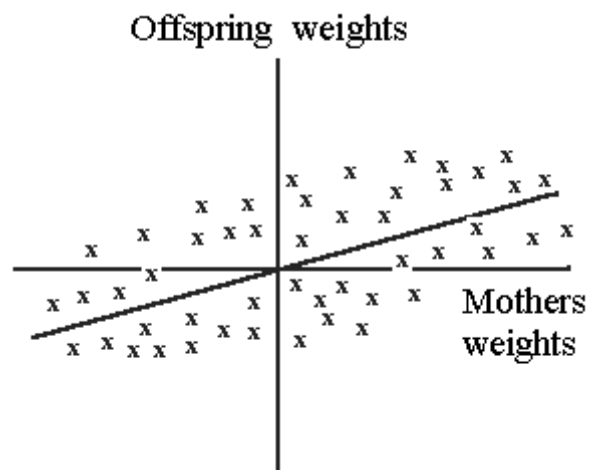
If environmental factors are causing the similarity, this can be estimated as shown in the formula below. The common environmental factors are particularly important in the litter's first months, when the mother and its maternal abilities cause the special environment. In half sibs, having a common father, the common genes are usually the only cause of similarity.

The general formula for interpretation of correlation between related individuals is shown below, and as symbol of this correlation t is used, this is also the symbol of the interclass correlation.

$$t = a * h^2 + c^2$$

Figure 6.4.

An example of a two-dimensional distribution for calculation of correlation, typically a trait of a mother and one of its offspring.



Now the heritability can be calculated on the basis of half sib correlation, c^2 being 0. Then c^2 for full sibs can be calculated based on the full sib correlation.

Example: Weight of mink aged 8 weeks, sample size: 508 animals in 107 litters with 37 sires.

In a statistical analysis of the weight of 508 mink puppies was recorded.

Half sib correlation $t = 0.03$ and

Full sib correlation $t = 0.41$.

These numbers put into the genetic model results in:

Half sib correlation $t = (1/4)h^2 + 0 = 0.03$ corresponding to $h^2 = 0.12$.

Full sib correlation $t = (1/2)h^2 + c^2 = (1/2)*0.12 + c^2 = 0.41$ corresponding to $c^2 = 0.35$. For calculation of c^2 for full sibs the heritability calculated for half sibs is applied.

The material for calculation of heritability of height of humans was shown in chapter 1. As mentioned there, the heritability was calculated as the regression of offspring on basis of the average of the parents. For height of humans it was around 0.6.

Mother-offspring correlation was found to be 0.4 while the correlation between father and offspring was only 0.2. This lower correlation between father and offspring might be caused by the fact that these do not share the same significant common environment as the mother and offspring. The 0.2 correlation between father and offspring gives a heritability of 0.4 for height of humans. This means that the mother-offspring correlation is highly influenced by their common environment.