

Introductory seminar  
“Mathematical Population Genetics”  
winter term 2017/18

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## 1 Hardy Weinberg Law

### 1.1 Blood types

The *ABO* blood types can (in the simplest case) be coded by the three alleles *A*, *B*, and *O* at a single locus, where *A* and *B* are dominant over *O* (genotypes *AO* and *BO* respectively have phenotypes [*A*] and [*B*]).

1. Denote with  $p_A$ ,  $p_B$  and  $p_O$  the frequencies of these alleles. Calculate, under the assumption of Hardy-Weinberg equilibrium, the relative frequencies of the blood types *A*, *B*, *AB* and *O* (phenotypes [*A*], [*B*], [*AB*], and [*O*]). Why exactly these?
2. Let  $R_A$ ,  $R_B$ ,  $R_{AB}$  and  $R_O$  denote the observed frequencies of the respective blood types within a population. From this calculate the allelic frequencies. Is it possible to do this without Hardy-Weinberg? How could one test whether the population is in Hardy-Weinberg equilibrium?

### 1.2 Dynamics of phenotype frequencies under assortative mating

Mating can be non-random with respect to some traits, respectively genes. We speak of assortative mating when similar individuals are more likely to mate with each other than expected by chance. Consider the following simple but instructive example. The two alleles  $A_1$  and  $A_2$  occur at a gene locus. Furthermore, the genotype  $A_1A_1$  has the same phenotype as  $A_1A_2$ , which is different from the phenotype of  $A_2A_2$ . Thus  $A_1$  is dominant. Let  $\rho$  denote the proportion of individuals that mate assortatively, *i.e.* only with individuals of the same phenotype. This proportion is assumed to be identical for both phenotypes. Denote the (ordered) genotype frequencies with  $P_{ij}$  and the allelic frequencies with  $p$  and  $q$ , ( $p + q = 1$ ).

1. The contribution of those individuals that mate randomly to the pool of individuals with genotype  $A_1A_1$ ,  $A_1A_2$  and  $A_2A_2$  in the next generation is then  $(1 - \rho)p^2$ ,  $(1 - \rho)2pq$  and  $(1 - \rho)q^2$  respectively (why?).

- Calculate the contribution of the assortatively mating individuals of genotype  $A_2A_2$  to the different types of offspring and also the contributions of the individuals with the dominant phenotype.
- Show that the following recursion holds:

$$P'_{11} = (1 - \rho)p^2 + \rho \frac{p^2}{1 - P_{22}}$$

### 1.3 Continued: Dynamics of phenotype frequencies under assortative mating

For details see exercise before.

- Derive also the recursion for the other two genotypes,  $P_{12}, P_{22}$ .
- Which conclusions can you draw from this set of recursions for the allelic frequencies.
- What recursion follow for the special cases of  $\rho = 0$  and  $\rho = 1$ ?

### 1.4 Dynamics of heterozygosity under assortative mating

We are considering the assortative mating case discussed in exercise 1.2,1.3. The quantity  $H = 2P_{12}$  is known as *heterozygosity*.

- Derive the recursion for  $H'$  (which can be expressed as a function of  $H$  and  $\rho$ ).
- For the case  $\rho = 1$ , derive the value of  $H(t)$  in generation  $t$  given e.g.  $p(t = 0) = \frac{1}{2}$ .
- What general conclusion can be derived for  $\lim_{t \rightarrow \infty} H(t)$  in the case of  $\rho < 1$ ?

### 1.5 Dynamics of phenotype frequencies with X-linkage

Consider a gene with two alleles  $A$  and  $a$ , located on the X chromosome (in mammals, females are XX while males are XY). The  $A$  allele is dominant (individuals with genotypes  $AA$  and  $Aa$  have phenotype  $[A]$  while individuals with genotype  $aa$  have phenotype  $[a]$ ). The frequencies of the  $A$  allele in males and females are respectively denoted  $p$  and  $q$ , initial frequencies are denoted  $p_0$  and  $q_0$ .

- Express the recursion for the allele frequencies  $p'$  and  $q'$ .
- Express the allele frequencies in males and females after  $t$  generations.
- Express the frequency of the  $[A]$  phenotype in females and in males.
- Does this dynamics change if the females:males ratio differs from 1:1 in the population?