

Genetic variation in populations (allele and genotype frequencies), HWE

Modul no.: Animal Genetics

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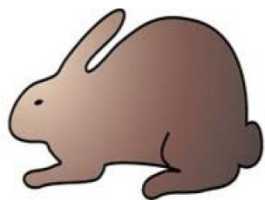
Fakulty of Agrobiolology, Food and Natural Resources



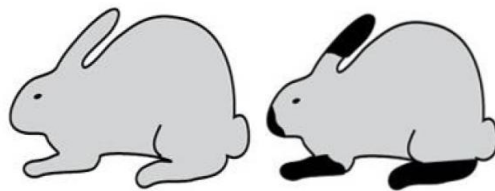
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At the level of individuals

Phenotype



wild



chinchilla



himalayan



albino

$$c^+ > c^{ch} > c^h > c$$

Genotype: one gene – one locus


 $c^+c^+, c^+c,$
 c^+c^{ch}, c^+c^h
 $c^{ch}c, c^{ch}c^{ch}, \underline{c^{ch}c^h}$
 c^hc^h, cc^h
 cc

*Note: Even when more than two alleles exists, any given individual can have no more than two alleles (one for the mother and one from the father).

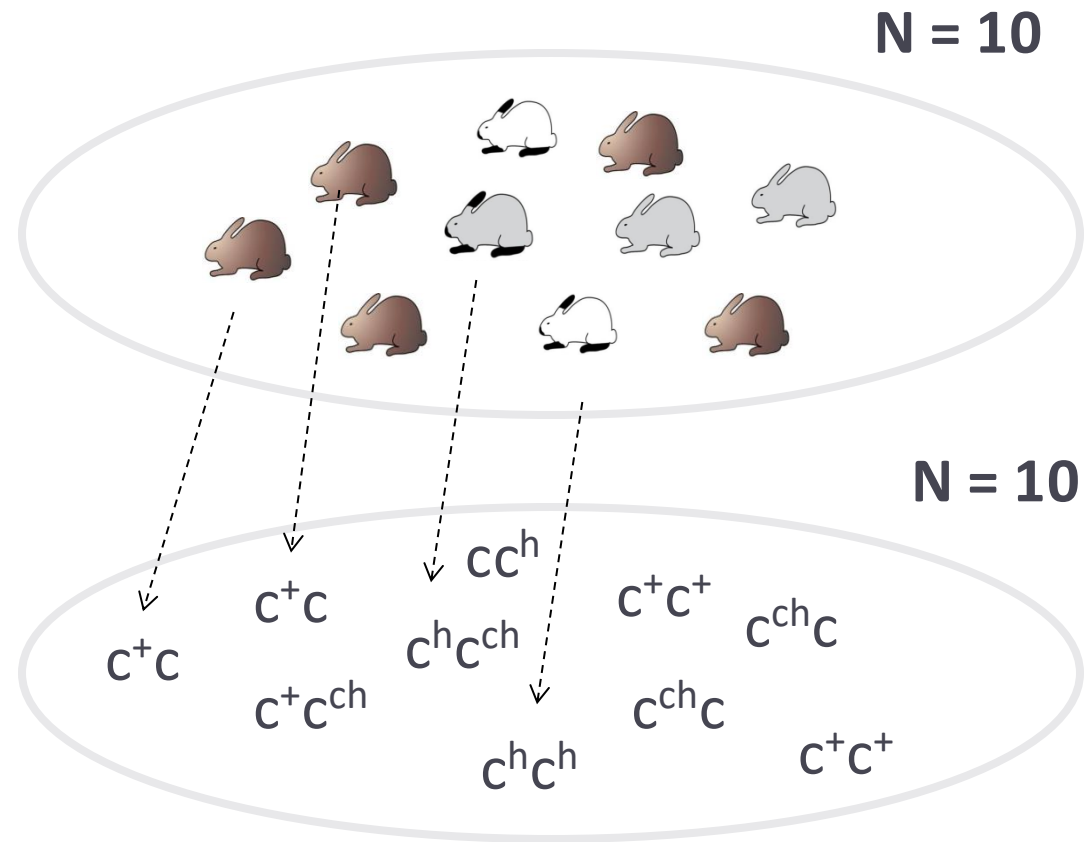
Population level

Alely

c^+ (wild), c (albino),
 c^{ch} (chinchila), c^h (himalayan)

10 rabbits \rightarrow 20 allele:

7 c^+ (A_1), 5 c (A_2),
 4 c^{ch} , (A_3) 4 c^h (A_4)



frequency of all. c^+ (A_1) = $p = ? = 7/20 = 0,35$

frequency of all. c (A_2) = $r = ? = 5/20 = 0,25$

frequency of all. c^{ch} (A_3) = $w = ? = 4/20 = 0,20$

frequency of all. c^h (A_4) = $v = ? = 4/20 = 0,20$

frequency of all. A_2

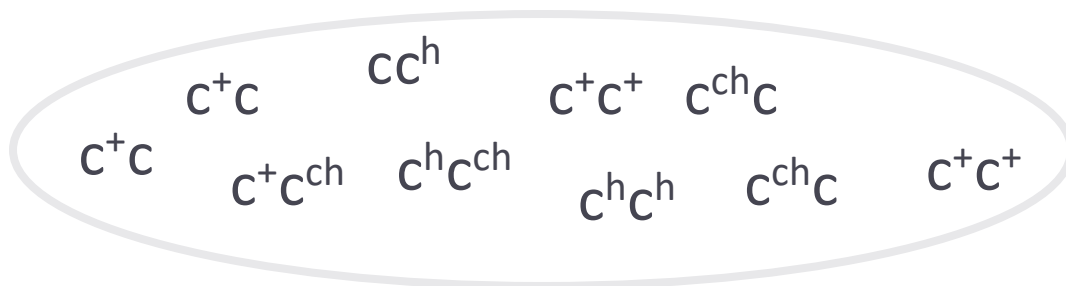
$q = r + w + v$
 $= ? = 13/20 = 0,65$

*Note: $p + q = 1$ (0.35 + 0.65)



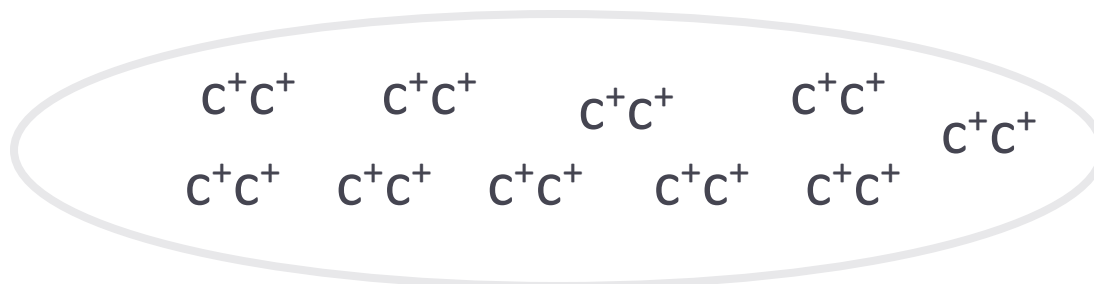
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Population 1: polymorphic loci (multiallelic: > two alleles)



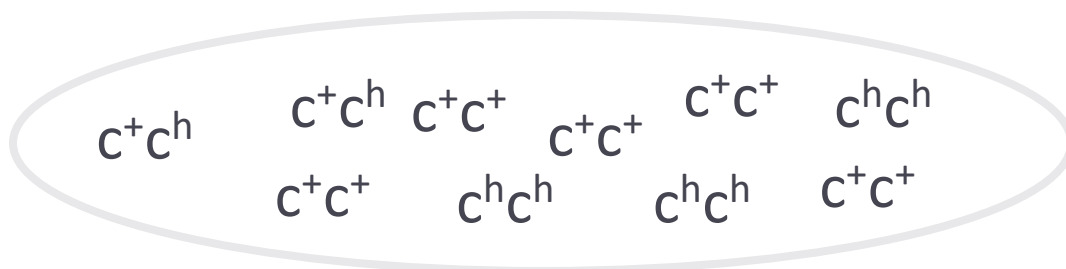
$N = 10$

Population 2: monomorphic loci



$N = 10$

Population 3: polymorphic locus (biallelic: only two alleles)



$N = 10$



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Biallelic locus: $p(A_1)$ & $q(A_2)$:

Base population

$$D = A_1A_1 \rightarrow d = \frac{D}{N}$$

$$H = A_1A_2 \text{ nebo } A_2A_1 \rightarrow h = \frac{H}{N}$$

$$R = A_2A_2 \rightarrow r = \frac{R}{N}$$

$$N = D + H + R$$



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$$N = D + H + R$$

Frequency of alleles

$$A_1 \rightarrow p = d + 0,5h$$

$$A_2 \rightarrow q = r + 0,5h$$



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$$N = D + H + R$$

Frequency of alleles

$$A_1 \rightarrow p = d + 0,5h$$

$$A_2 \rightarrow q = r + 0,5h$$

Next Generations

$$A_1A_1 \rightarrow A_{1\sigma} \times A_{1\phi} \rightarrow p \times p = p^2$$

$$A_1A_1 \rightarrow A_{1\sigma} \times A_{1\phi} \text{ nebo } A_{2\sigma} \times A_{1\phi} = (p \times q) + (q \times p) = 2pq$$

$$A_2A_2 \rightarrow A_{2\sigma} \times A_{2\phi} \rightarrow q \times q = q^2$$



Biallelic locus: $p(A_1)$ & $q(A_2)$:

Base population

$$D = A_1A_1 \rightarrow d = \frac{D}{N}$$

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$$N = D + H + R$$

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$$A_2A_2 \rightarrow A_{2\sigma} \times A_{2\phi} \rightarrow q \times q = q^2$$



Hardy-Weinberg equilibrium (HWE)

The Hardy–Weinberg principle states that both allele and genotype frequencies in a population remain constant – that is , they are in equilibrium – from generation to generation unless specific disturbing influences are introduced.



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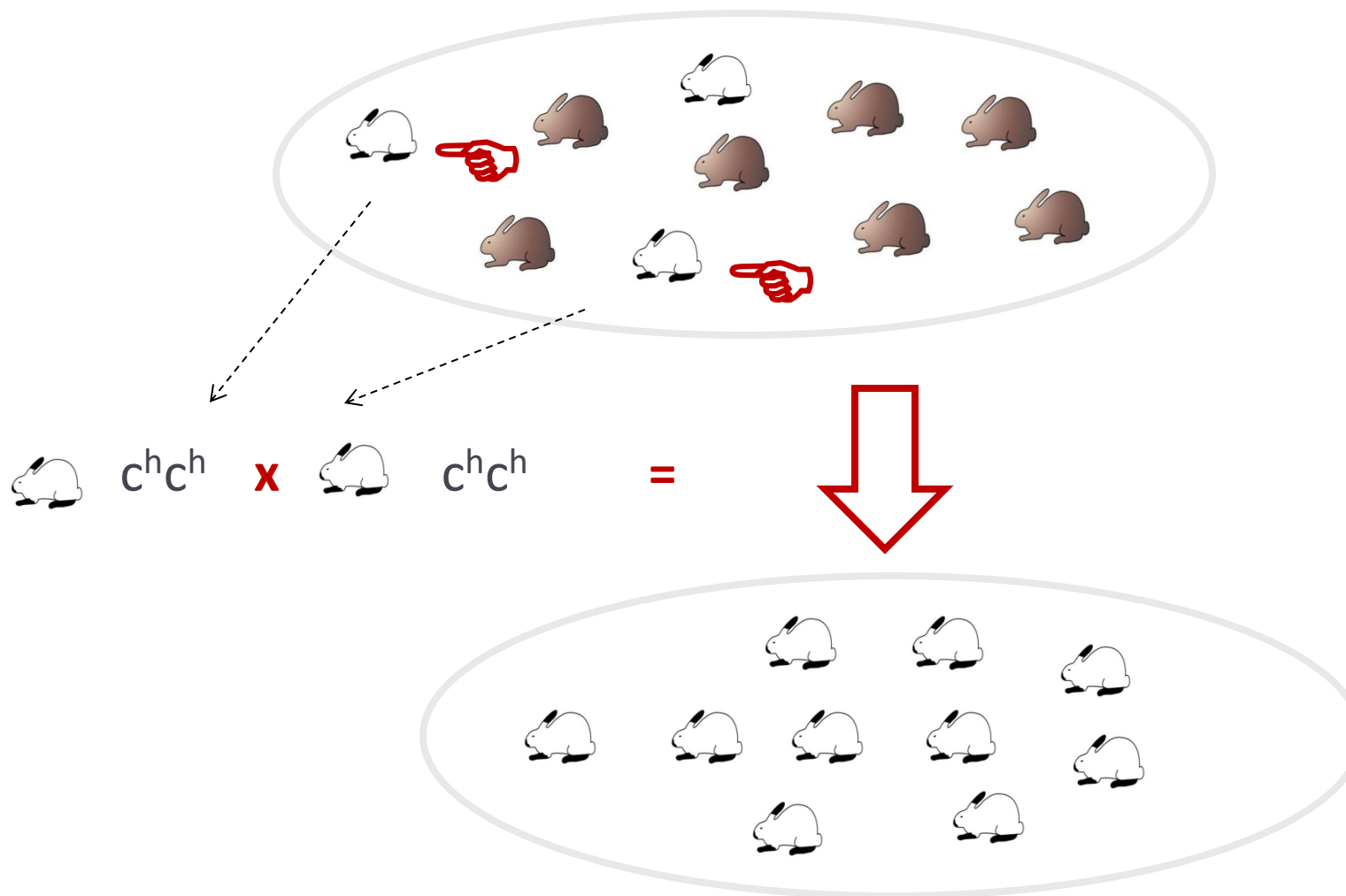
Those disturbing
influences include:

Selection,
Mutation
Migration,
Non-Random Mating,
Genetics Drift (finite population size),

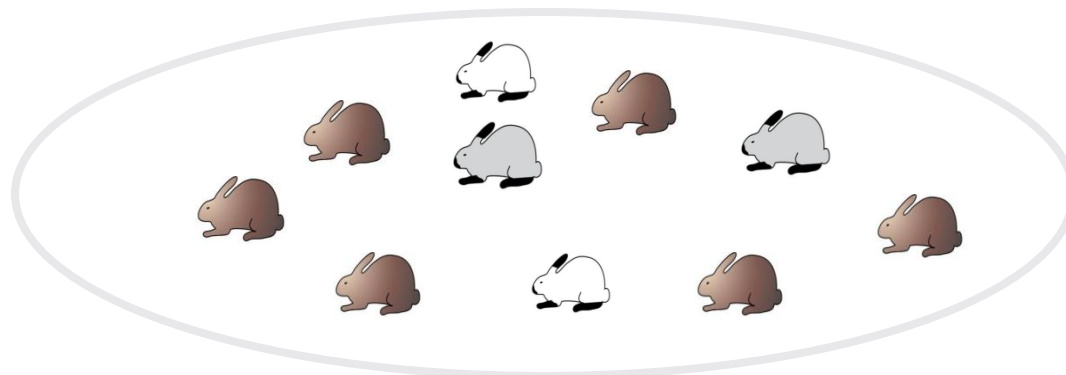
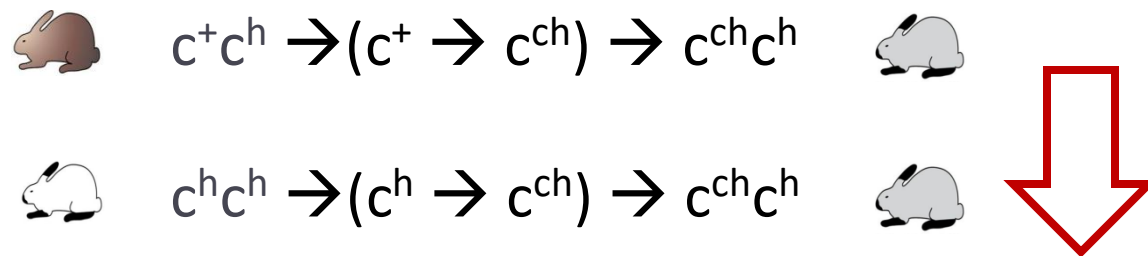
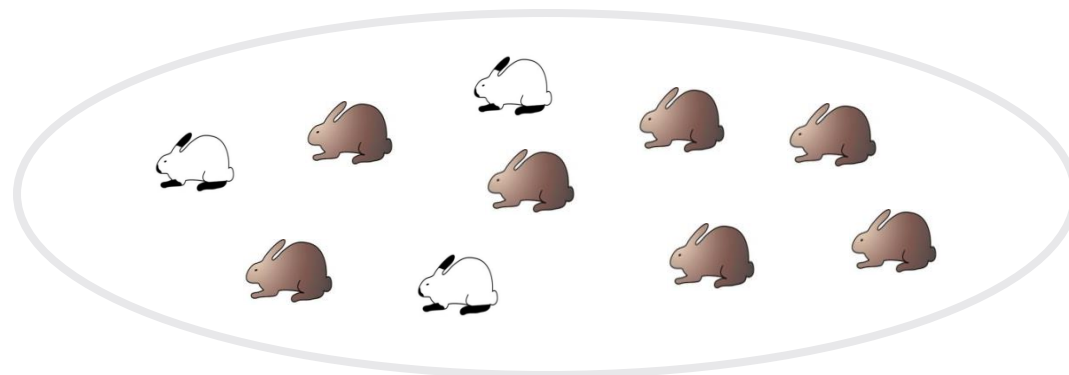
It is important to understand that outside the lab, one or more of these „disturbing influences“ are always in effect.

That is, Hardy-Weinberg Equilibrium is impossible in nature.
Genetic equilibrium is an ideal state that provides a baseline to measure genetic change against.

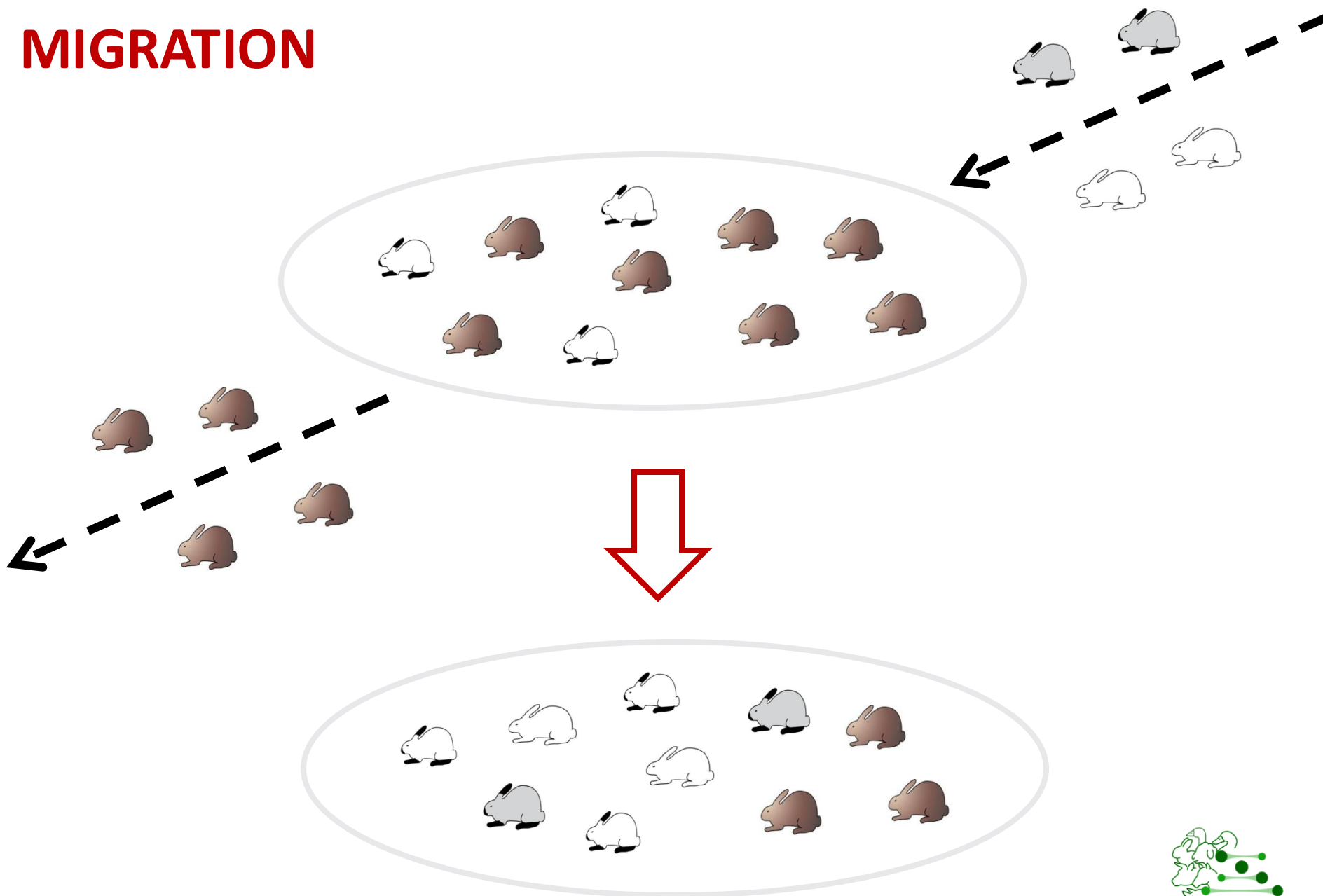
SELECTION



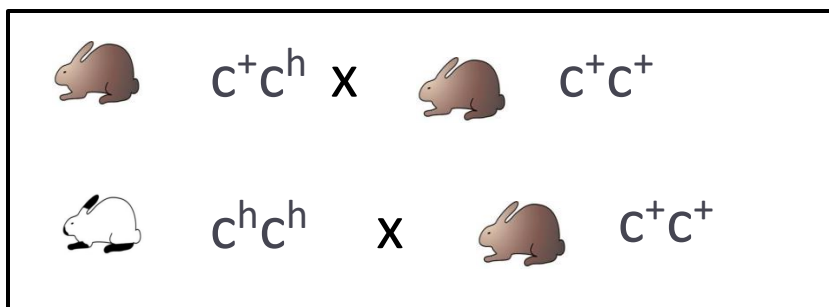
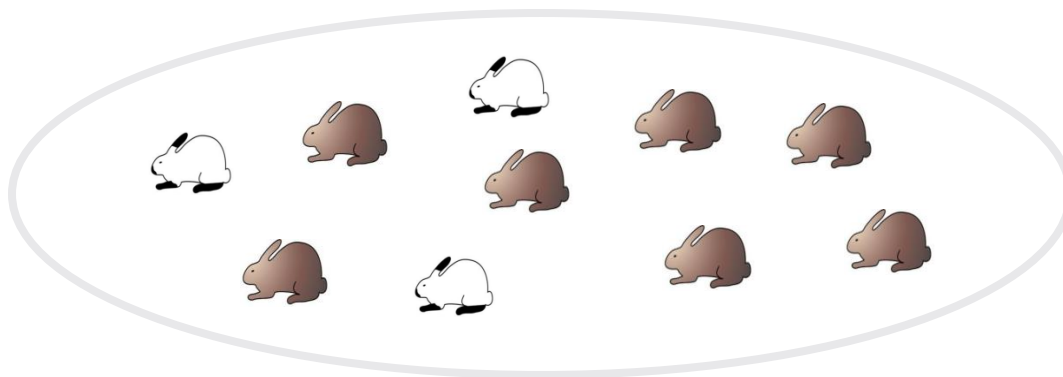
MUTATION



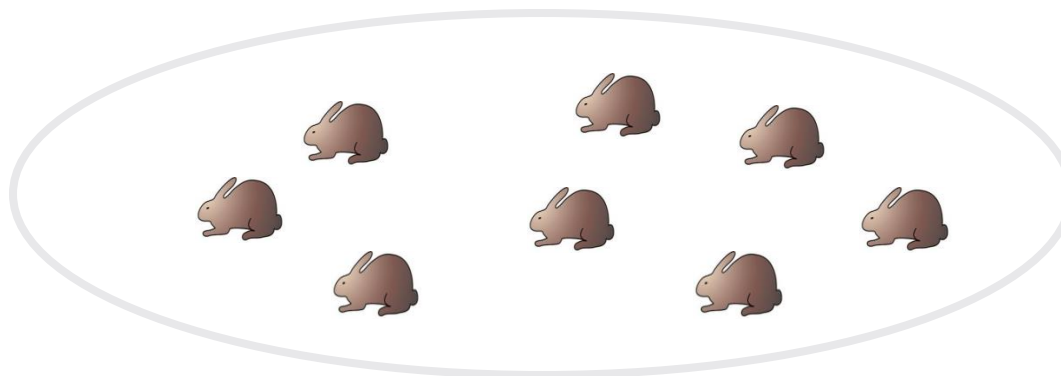
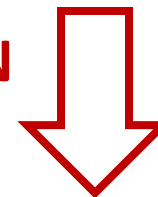
MIGRATION



RANDOM MATING



ONLY THEN



WITHOUT GENETIC DRIFT (infinite population size)



samo-oplození:
Population s $N=1$

$$[c^+c^h] \rightarrow P(c^+):P(c^h) = 0.5 : 0.5$$

$$2 \text{ times} \rightarrow (c^+, c^+), (c^+, c^h), (c^h, c^+), (c^h, c^h)$$

$$1 : 2 : 1$$

$$P(2 \text{ times } c^+ \text{ or } 2 \text{ times } c^h) = 0,5$$

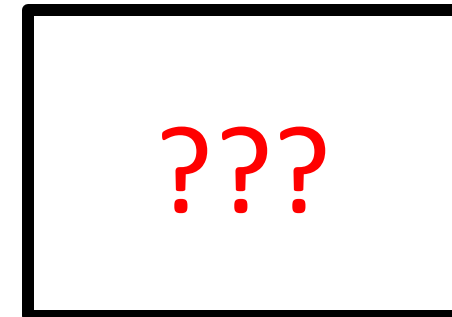
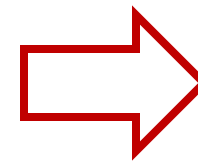
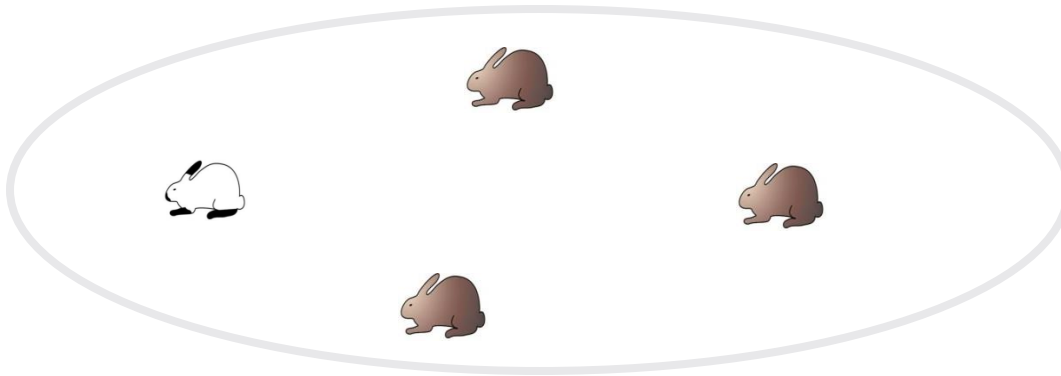
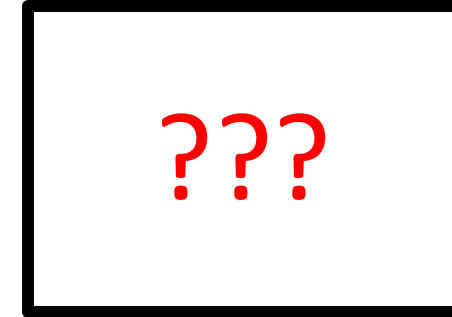
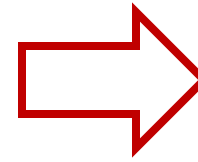
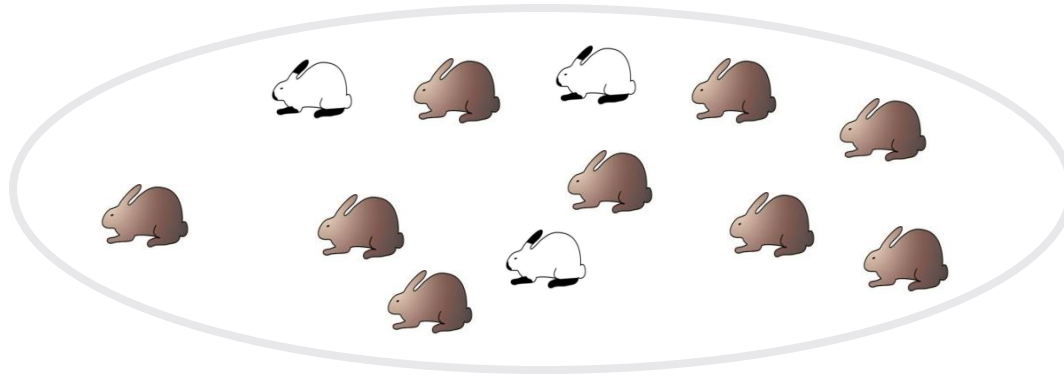
$$P(10 \text{ times } c^+ \text{ or } 10 \text{ times } c^h) < 0,5$$

$$P(100 \text{ times } c^+ \text{ or } 100 \text{ times } c^h) \ll 0,5$$



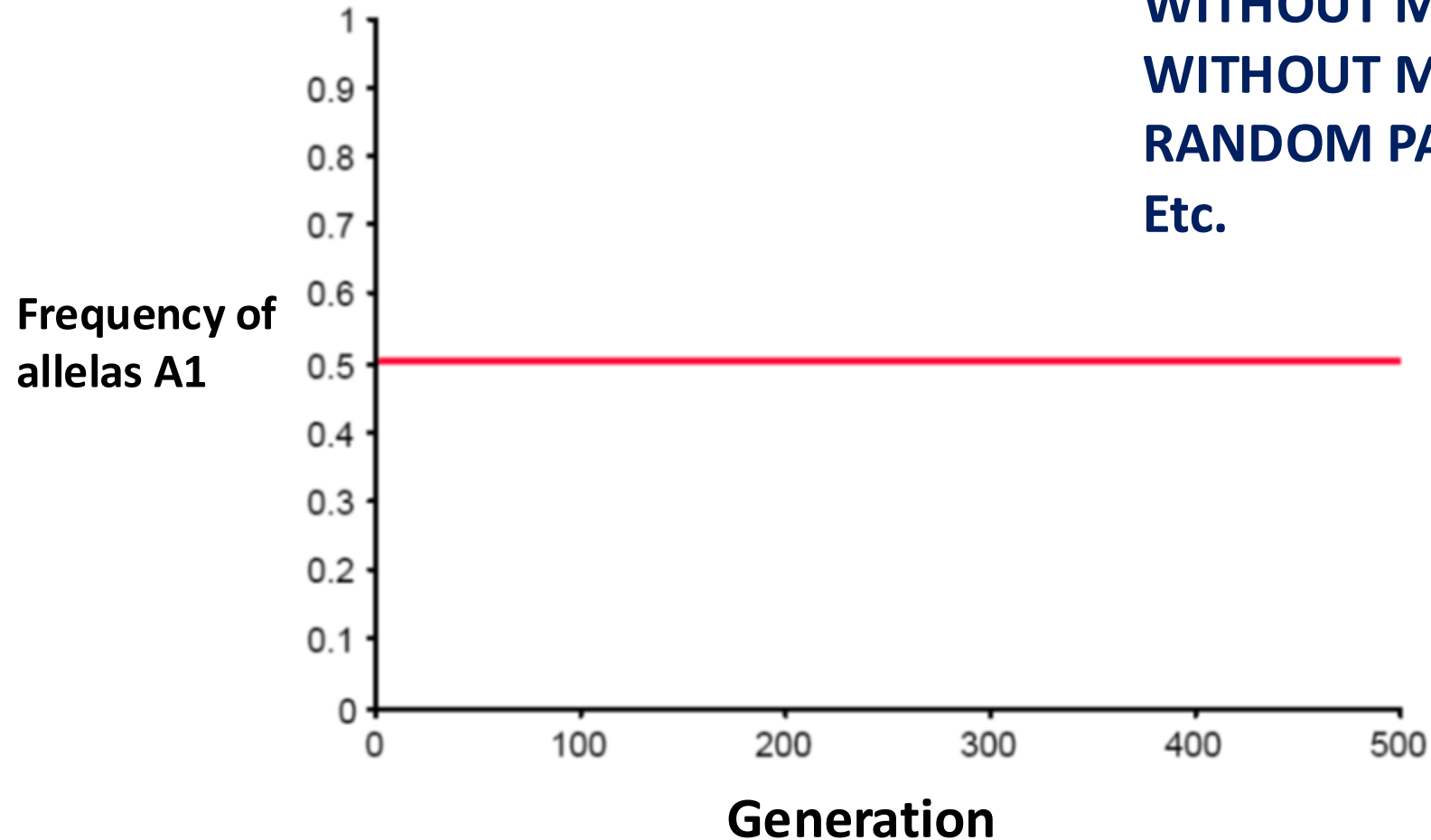
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WITHOUT GENETIC DRIFT (infinite population size)



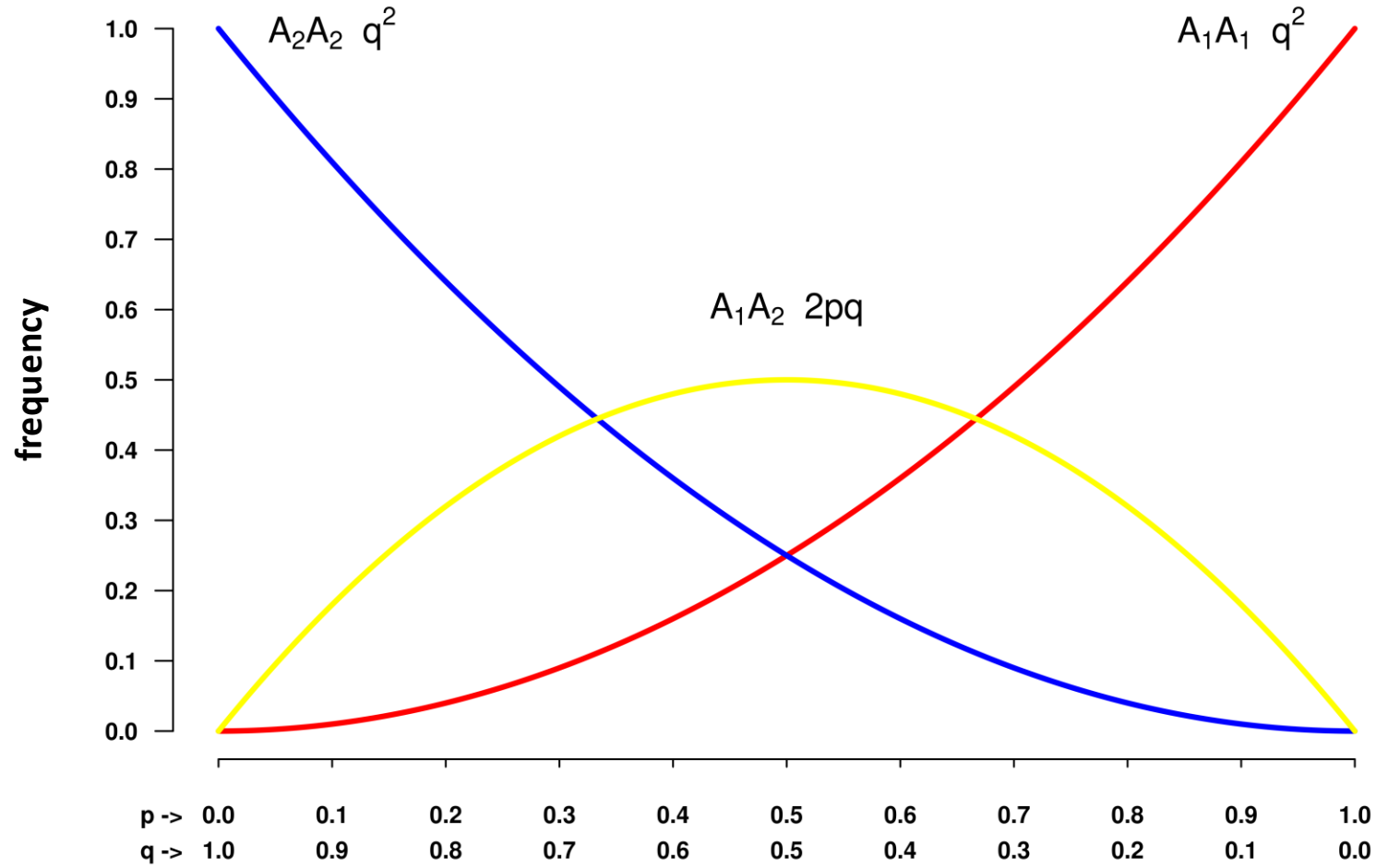
POPULATION SIZES: + infinite

WITHOUT SELECTION
WITHOUT MUTATION
WITHOUT MIGRATION
RANDOM PAIRING
Etc.



$$p_0 = p_1 = p_2 = p_3 = \dots = p_t$$

Hardy-Weinberg equilibrium (HWE)



Partners:



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Thank you for your attention!

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