

3rd term

Population genetics

Genetic drift and mutation

Problems

①

A: $p = 0.875$

a: $q = 0.125$

4 individuals \rightarrow 8 alleles

No a \rightarrow 8A $\rightarrow 0.875^8 = 0.344 \rightarrow$ probability of no a ($P(A) = P(\text{no a})$)

No A \rightarrow 8a $\rightarrow 0.125^8 = 5.96 \cdot 10^{-8} \rightarrow$ probability of no A ($P(a) = P(\text{no A})$)

$$p^2 + 2pq + q^2 = 1 \rightarrow 2pq = 1 - p^2 - q^2 \rightarrow 2pq = 1 - (0.125^2 + 0.875^2) = 0.219$$

$$P(A \text{ and } a) = P(Aa) = 1 - P(A) - P(a) \rightarrow P(Aa) = 1 - 0.344 - 5.96 \cdot 10^{-8} = 0.656$$

②

Aa \rightarrow 12 individuals \rightarrow 24 alleles

10% heterozygous \rightarrow num of generations \rightarrow $H_0 = 1$
 $H_t = 1 - 0.9 = 0.1 \rightarrow$ 10% heterozygous

$$H_t = H_0 \left(1 - \frac{1}{2N}\right)^t \rightarrow \ln\left(\frac{H_t}{H_0}\right) = t \cdot \ln\left(1 - \frac{1}{2N}\right) \rightarrow t = \frac{\ln(H_t/H_0)}{\ln(1 - (1/2N))} \rightarrow$$

$$t = \frac{\ln(0.1/1)}{\ln(1 - (1/24))} = 54.1 \rightarrow \text{it will take 54 generations for the population to be heterozygous in 10\%}$$

$$N = 240$$

$$t = \frac{\ln(0.1/1)}{\ln(1 - (1/2 \cdot 240))} = 1104.1 \rightarrow \text{in 1104 generations of a population of 240 individuals you will have 10\% heterozygosity}$$

③

$N = 95$

1 new allele

$P_0 \rightarrow \frac{1}{2N} = P_0 \rightarrow \frac{1}{2 \cdot 95} = 0.0053 \rightarrow$ there is a 0.53% probability that it will drift to fixation

$P_{\text{lost}} = 1 - P_0 \rightarrow P_{\text{lost}} = 1 - 0.0053 = 0.9947 \rightarrow$ There is a 99.47% probability of loss

$$T_{\text{fix}} = -4N \cdot \frac{(1-p) \ln(1-p)}{p} = -4 \cdot 95 \cdot \frac{(1-0.0053) \ln(0.9947)}{0.0053} = 3789 \rightarrow \text{Average fixation time is 3789 generations}$$

$$T_{\text{loss}} = -4N \cdot \frac{p \ln(p)}{1-p} = -4 \cdot 95 \cdot \frac{0.0053 \ln(0.0053)}{1-0.0053} = 357.6 \rightarrow \text{Average allele loss time is of 358 generations}$$

4

$$p_A = 0.75$$

$$N = 40$$

$$p'_A = 0.75$$

$$P(k \text{ alleles in } n \text{ alleles}) = \frac{n!}{k!(n-k)!} \cdot p^k \cdot q^{n-k}$$

$$k = 2N \cdot p'_A \rightarrow k = 2 \cdot 40 \cdot 0.75 = 60$$

$$q = 1 - p \rightarrow q = 1 - 0.75 = 0.25$$

$$n = 2N \Rightarrow n = 2 \cdot 40 = 80$$

$$P(60 \text{ in } 80) = \frac{80!}{60!(80-60)!} \cdot 0.75^{60} \cdot 0.25^{(80-60)} = 0.1025$$

There is a 10.25% chance that in the next generation the allele A frequency is 0.75

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$$N = 12 \begin{matrix} \nearrow 6M \\ \searrow 6F \end{matrix}$$

$$A = 0.5 = p_0$$

$$a = 0.5 = q_0$$

$$p_1^1 = 0.458 \quad q_1^1 = 1 - 0.458 = 0.542$$

$$p_2^1 = 0.583 \quad q_2^1 = 1 - 0.583 = 0.417$$

$$k = 2N \cdot p_0 \rightarrow k = 12 \cdot 0.5 = 6$$

$$n = 2N \Rightarrow 2 \cdot 6 = 12$$

$$P_1 = \frac{12!}{6!(12-6)!} \cdot 0.458^6 \cdot 0.542^{(12-6)} = 0.2162$$

$$P_2 = \frac{12!}{6!(12-6)!} \cdot 0.583^6 \cdot 0.417^{(12-6)} = 0.19077$$

$$p_d = 0.375$$

$$q_d = 1 - 0.375 = 0.625$$

$$k_d = 12 \cdot 0.375 = 4.5 \approx 5$$

$$P = \frac{12!}{5!(12-5)!} \cdot 0.375^5 \cdot 0.625^{(12-5)} = 0.22$$

$$P(0.375 \text{ in } P_1 \text{ and } P_2) = 0.22 \cdot 0.22 = 0.0484 \rightarrow 4.84\% \text{ that } P_1 \text{ and } P_2 \text{ have } p = 0.375 \text{ in the second generation}$$

⑥ p1s Genetic drift theory

$$p_0 = 0.5 \quad p_{q_0} = 0.5 \rightarrow 2A \ 2a$$

$$t = 2$$

$$p_2 = 0.25 \quad q_2 = 0.75 \rightarrow 1A \ 3a$$

$$P(0.25|0.5) = P(1A|2A) = P(1 \text{ allele } A \text{ in gen } 2) = 0.4219$$

→ table

	0	1	2	3	4	n th gen
n th alleles	1	0	0.4219	0.25	0.0419	0
ingen	0	0.4219	0.25	0.375	0.25	0.0625

$$P(1 \text{ all in gen } 2) = (0 \cdot 0.0625) + (0.4219 \cdot 0.25) + (0.25 \cdot 0.375) + (0.0419 \cdot 0.25) + (0 \cdot 0.0625) = 0 + 0.1055 + 0.0938 + 0.01173 + 0 = 0.21103$$

The probability that a population of 2 individuals (4 alleles) that start with $p_0 = 0.5$ and have $p_2 = 0.25$ in 2 generations is of 0.21103

⑦

$$N_m = \frac{55000}{50} = 1100 \text{ males}$$

$$N_f = 55000 - 1100 = 53900 \text{ females}$$

assuming $N_{\text{cont}} = 1$
adding individuals all in groups of 50 F to 1 M

$$N_e = \frac{4N_m N_f}{N_m + N_f} \rightarrow N_e = \frac{4 \cdot 1100 \cdot 53900}{1100 + 53900} = 4312$$

Assuming equal probability of M/F individual independency

$$27500 = M \quad 27500 = F \quad (\text{there will be males without a group})$$

⑧

$$N_0 = 104 \quad N_1 = 62 \quad N_2 = 10 \quad N_3 = 110 \quad t = 4$$

$$\frac{1}{N_e} = \frac{1}{t} \cdot \left(\frac{1}{N_0} + \frac{1}{N_1} + \frac{1}{N_2} + \frac{1}{N_3} \right) \Rightarrow \frac{1}{N_e} = \frac{1}{4} \cdot \left(\frac{1}{104} + \frac{1}{62} + \frac{1}{10} + \frac{1}{110} \right) = 0.03371 \rightarrow$$

$$\rightarrow N_e = \frac{1}{0.03371} = 29.6658 \rightarrow \text{The effective size of the population is } 30$$

Genetic drift

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$$N = 55000$$

Assuming that all males controls a group of exactly 50 females
, that all individuals are adults and that there are no
lone individuals. Otherwise you could assume an even (50/50) split.

$$N_m = \frac{55000}{50} = 1100$$

all others must be females

$$N_f = 55000 - 1100 = 53900$$

$$a) N_e = \frac{4 \cdot 1100 \cdot 53900}{1100 + 53900} = 4312 \approx N_e \rightarrow N_e = \frac{4 N_m \cdot N_f}{N_m + N_f}$$

$$b) N_m = \frac{20}{50} = 0.4 \rightarrow \text{round up, for the species to continue reproducing you need 1 male at least}$$

$$N_m = 1$$

$$N_f = N - N_m \rightarrow N_f = 20 - 1 = 19$$

$$N_e = \frac{4 \cdot 1 \cdot 19}{1 + 19} = 3.8 \approx 4$$

$$c) H_0 = 0.028$$

$$t = 5$$

Assuming that the bottleneck didn't change the heterozygosity proportion

Assuming that the population grows to contain 50 individuals in 5 generations

$$H_t = H_0 \left(1 - \frac{1}{2N}\right)^t \rightarrow H_5 = 0.028 \left(1 - \frac{1}{2 \cdot 50}\right)^5 = 0.0266$$

Mutation

①

$$A \rightarrow a$$

$$\mu = 0.00002$$

$$p_0 = 0.5$$

$$p_{10} ? \quad p_t = p_0 (1 - \mu)^t$$

$$p_{10} = 0.5 (1 - 0.00002)^{10} = 0.499900009 \rightarrow \text{Frequency of A in 10 generations}$$

$$p_{100} = 0.5 (1 - 0.00002)^{100} = 0.49900099 \rightarrow \text{Frequency of A in 100 generations}$$

$$p_{1000} = 0.5 (1 - 0.00002)^{1000} = 0.49009 \rightarrow \text{Frequency of A in 1000 generations}$$

$$p_{10000} = 0.5 (1 - 0.00002)^{10000} = 0.409365 \rightarrow \text{Frequency of A in 10000 generations}$$

It's a very slow change in allele frequency

②

$$\mu = 0.005 \quad v = 0.0015$$

$$p_0 = 0.8 \quad q_0 = 0.2$$

$$p_t = \frac{v}{\mu + v} + \left(p_0 - \frac{v}{\mu + v} \right) (1 - \mu - v)^t$$

$$q_1 = \frac{0.0015}{0.005 + 0.0015} + \left(0.2 - \frac{0.0015}{0.005 + 0.0015} \right) (1 - 0.005 - 0.0015)^1 = 0.2002$$

In the first generation $q = 0.2002$

$$q_2 = \frac{0.0015}{0.005 + 0.0015} + \left(0.2 - \frac{0.0015}{0.005 + 0.0015} \right) (1 - 0.005 - 0.0015)^2 = 0.200398$$

In the second generation $q = 0.200398$

$$q_3 = \frac{0.0015}{0.005 + 0.0015} + \left(0.2 - \frac{0.0015}{0.005 + 0.0015} \right) (1 - 0.005 - 0.0015)^3 = 0.2005961$$

In the third generation $q = 0.2005961$

$$p = \frac{v}{\mu + v} \rightarrow p = \frac{0.0015}{0.005 + 0.0015} = 0.2308$$

$$q = 1 - p \rightarrow q = 1 - 0.2308 = 0.7692$$

Allele frequencies in equilibrium are $q = 0.7692$ $p = 0.2308$