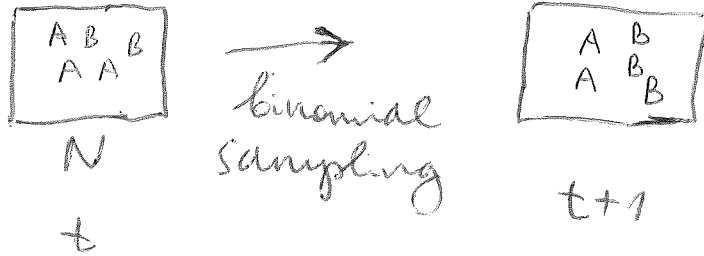


Fixation of mutant genes in a population



Consider a population of N genes in which the freq. of the allele A is p ($0 \leq p \leq 1$).

The freq. of allele B is then $q = 1 - p$

Define $u(p, t) = \text{prob. that } A \text{ is fixed (i.e. its freq.} = 1) \text{ in } \leq t \text{ generations (continuous time approx'n)}$

$$u(p, t + \delta t) = \int d(\delta p) \underbrace{f(p, p + \delta p; \delta t)}_{\text{prob. density } p \rightarrow p + \delta p \text{ in } \delta t} \times u(p + \delta p, t)$$

Introduce

$$\int \lim_{\delta t \rightarrow 0} \frac{1}{\delta t} \int (\delta p) f(p, p + \delta p; \delta t) d(\delta p) \equiv M,$$

$$\int \lim_{\delta t \rightarrow 0} \frac{1}{\delta t} \int (\delta p)^2 f(p, p + \delta p; \delta t) d(\delta p) \equiv V$$

Just as before, we obtain:

$$\frac{\partial u(p,t)}{\partial t} = \frac{V}{2} \frac{\partial^2 u}{\partial p^2} + M \frac{\partial u}{\partial p}$$

BCs: $u(0,t) = 0, \quad u(1,t) = 1$

Measure time in generations \rightarrow
 $\rightarrow M, V$ are the mean & variance of the change of p per generation.

Consider $u(p) = \lim_{t \rightarrow +\infty} u(p,t)$:

$$\frac{\partial u}{\partial t} = 0 \Rightarrow \frac{V}{2} \frac{d^2 u}{dp^2} + M \frac{du}{dp} = 0 \quad (*)$$

"steady state" $u(0) = 0, \quad u(1) = 1$

(*) can be solved:

Indeed, $\frac{du}{dp} = \frac{G(p)}{C_1}$; $\frac{d^2 u}{dp^2} = G(p) \left[-\frac{2M(p)}{V(p)} \right] \frac{1}{C_1}$ $u(p) = \frac{\int_0^p G(x) dx}{\int_0^1 G(x) dx}$

$-G(p)M(p) \frac{1}{C_1} + \frac{MG(p)}{V(p)C_1} = 0$, as expected $\int_0^x \frac{2M(y)}{V(y)} dy$ $\underbrace{\int_0^1 G(x) dx}_{C_1}$

$$G(x) = e^{-\int_0^x \frac{2M(y)}{V(y)} dy}$$

Note that $u(\frac{1}{N})$ is a chance of fixation of a "novel" mutant gene.

Now, for the binomial distr'n we have: $\int \mu = np$, $\left\{ \begin{array}{l} \leftarrow \# \text{ trials} \\ \leftarrow \text{prob. of success} \end{array} \right.$
 $\sigma^2 = npq$

Prob. (i alleles of type A) =

$$= \frac{N!}{i!(N-i)!} p^i (1-p)^{N-i}$$

No selection:

$$E(\Delta p | p) = \frac{E(i)}{N} - p = \frac{\overset{\text{freq. of allele A}}{Np}}{N} - p = \underline{\underline{0}}$$

new freq. on average
old freq.

$$\text{Var}(\Delta p | p) = \text{Var}\left(\frac{i}{N} - p | p\right) = \frac{\text{Var}(i)}{N^2} = \frac{Np(1-p)}{N^2} = \frac{p(1-p)}{N}$$

With selection,

	A	B	
	$1+s$	1	\leftarrow prob. of survival
before selection \Rightarrow	p	q	
after selection \Rightarrow	$p' = \frac{p(1+s)}{p(1+s) + q}$	$q' = \frac{q}{p(1+s) + q}$	$= q'$

$p' + q' = 1$

$$p' - p \equiv \Delta_s p = \frac{p(1+s) - p^2(1+s) - pq}{p(1+s) + q} = \frac{pq s}{p(1+s) + q} \approx pq s$$

$1+ps \approx 1$ if s is small

So, $E(\Delta p | p) = sp(1-p)$.

In our notation, $\frac{2M(y)}{V(y)} = 2Ns$ & $G(x) = e^{-2N \frac{V(y)}{s} x}$ " $\frac{2[spq]}{p\theta/N}$

Finally, $u(p) = \frac{1 - e^{-2Nsp}}{1 - e^{-2Ns}}$ Kimura's formula

$$u\left(\frac{1}{N}\right) = \frac{1 - e^{-2s}}{1 - e^{-2Ns}} \approx \frac{2s}{1 - e^{-2Ns}}$$

$$\lim_{s \rightarrow 0} u\left(\frac{1}{N}\right) = \frac{2s}{2Ns} = \frac{1}{N}, \text{ as expected.}$$

$u(p) = p$ is a well-known neutral result.