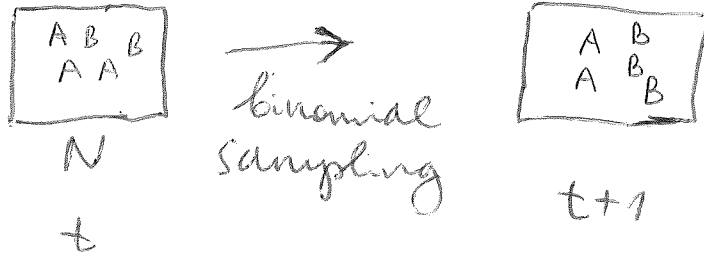


Fixation of mutant genes in a population



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.

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

With selection,

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

$p' + q' = 1$

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

$1+ps \approx 1$  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

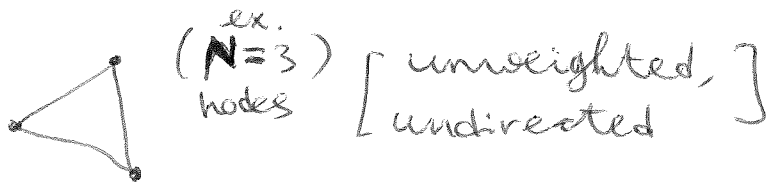
$$u\left(\frac{1}{N}\right) = \frac{1 - e^{-2s}}{1 - e^{-2Ns}} \approx \frac{2s}{1 - e^{-2Ns}}$$

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

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

Beltt, 2005

Consider a complete graph (all nodes connected to one another):



MFPT from node  $i$  to  $j$ :

$$T_{ij} = \underbrace{\frac{1}{N-1}}_{\text{go directly to } j} + \underbrace{\frac{N-2}{N-1}}_{\text{go to node } h \neq j} (1 + T_{hj}) =$$

$$= \frac{1}{N-1} + \frac{N-2}{N-1} (1 + T_{ij}) \Rightarrow \boxed{T_{ij} = N-1}$$

$\frac{N-2}{N-1} N + \frac{1}{N-1} = \frac{N^2 - 2N + 1}{N-1} = N-1$ , as expected

MRT:  $T_{ii} = 1 + \overset{N-1}{T_{ji}} = N$   
must step off

Furthermore, let  $F_{ij}(t)$  be the prob. that FPT  $i \rightarrow j$  is exactly  $t$ .

Then

$$F_{ij}(t) = \frac{1}{N-1} \delta_{t,1} + \underbrace{\frac{N-2}{N-1}}_{\text{prob. not to reach } j} F_{ij}(t-1) \quad (**)$$

Introduce a generating f'n:

$$\hat{F}_{ij}(x) = \sum_{t=0}^{\infty} F_{ij}(t) x^t$$

Then we can act with

$\sum_{t=0}^{\infty} x^t$  on (\*\*):

$$\begin{aligned}\hat{F}_{ij}(x) &= \frac{x}{N-1} + \sum_{t=0}^{\infty} x x^{t-1} \frac{N-2}{N-1} F_{ij}(t-1) = \\ &= \frac{x}{N-1} + \frac{N-2}{N-1} x \underbrace{\sum_{t'=0}^{\infty} x^{t'} F_{ij}(t')}_{\hat{F}_{ij}(x)}\end{aligned}$$

Then  $\hat{F}_{ij}(x) = \frac{x}{(N-1) - (N-2)x}$

Taylor expand to read off  $F_{ij}(t)$ :

$$\hat{F}_{ij}(x) \approx \frac{x}{N-1} \left( 1 + \frac{N-2}{N-1} x + \dots \right) \Rightarrow \left. \begin{array}{l} F_{ij}(1) = \frac{1}{N-1} \\ F_{ij}(2) = \frac{N-2}{(N-1)^2} \\ \dots \end{array} \right\}$$

Note that  $T_{ij} = \sum_{t=0}^{\infty} t F_{ij}(t) = N-1$ ,  
as expected