

Muller's ratchet: model of accumulating delet. mutations in populations w/ no recombination

imagine a population w/ diff. genetic variants. The variant w/ k deleterious mutations has a growth rate:

$$f_k = f_0 (1 - k s) \quad k = 0, 1, 2, \dots$$

$\uparrow$ growth rate w/o any mutations $\rightarrow$ cost per mutation

additive model of fitness: additional mutations add linearly to cost

model of growth (ignore conversion b/t types):

$$\frac{dn_k(t)}{dt} = f_k n_k(t)$$

$n_k(t)$ = # organisms of type k

for now: populations assume large, ignore stochastic fluct.

$\rightarrow n_k(t) = n_k(0) e^{f_k t}$

derive an equ. for $X_k(t) \equiv \frac{n_k(t)}{\sum_{k=0}^{\infty} n_k(t)}$ } $N(t)$ = total popul.

frac. of k in total

$$\frac{dx_k}{dt} = \frac{d}{dt} \left(\frac{n_k}{N} \right) = \frac{1}{N} \frac{dn_k}{dt} - \frac{n_k}{N^2} \frac{d}{dt} N$$

$$= f_k \frac{n_k}{N} - \frac{n_k}{N^2} \frac{d}{dt} \sum_{l=0}^{\infty} n_l$$

$$\bar{f} = \frac{1}{N} \sum_{l=0}^{\infty} f_l n_l$$

= mean fitness
(growth rate)
of entire popul.

$$= f_k x_k - \frac{n_k}{N^2} \sum_{l=0}^{\infty} f_l n_l$$

$$= f_k x_k - x_k \bar{f}$$

$$\Rightarrow \frac{dx_k}{dt} = (f_k - \bar{f}) x_k \quad \text{replicator equation}$$

add conversion between types:

$$\frac{dx_k}{dt} = (f_k - \bar{f}) x_k - \mu x_k + \mu x_{k-1}$$

loss to

$k \rightarrow k+1$

mutations

(w/ rate μ)

↑

total mut.

rate for

deleterious mutations

↑

gain from

$k-1 \rightarrow k$

mutations

• no back mutations

• no adaptive mutations

(can't undo a mutation,

$k+1 \rightarrow k$ is
not allowed)

in our case: $f_k = f_0 (1 - k \phi)$

$$\bar{f} = f_0 (1 - \bar{k} \phi)$$

avg.
of
mutations

$$\bar{k} = \frac{1}{N} \sum_{l=0}^{\infty} l n_l = \sum_{l=0}^{\infty} l x_l$$

$$f_k - \bar{f} = (\bar{k} - k) \phi$$

$$\frac{dx_k}{dt} = [(\bar{k} - k) \phi - \mu] x_k + \mu x_{k-1} \quad k=0,1,2,\dots$$

look for steady state sol'n $\frac{dx_k}{dt} = 0$

check: $x_k = e^{-\lambda} \frac{\lambda^k}{k!}$ Poisson distrib. $\lambda = \frac{\mu}{\phi}$

$$\sum_{k=0}^{\infty} x_k = 1$$

$$\bar{k} = \sum_{k=0}^{\infty} k x_k = \lambda$$

plug in & check it works:

$$\begin{aligned} & [(\lambda - k) \phi - \mu] \frac{e^{-\lambda} \lambda^k}{k!} + \mu \frac{e^{-\lambda} \lambda^{k-1}}{(k-1)!} \\ &= \frac{e^{-\lambda} \lambda^{k-1}}{(k-1)!} \left\{ \underbrace{[(\lambda - k) \phi - \mu] \frac{\lambda}{k}}_{\phi [\cancel{\lambda - k - \lambda}] \cdot \frac{\lambda}{k}} + \mu \right\} = 0 \quad \checkmark \end{aligned}$$

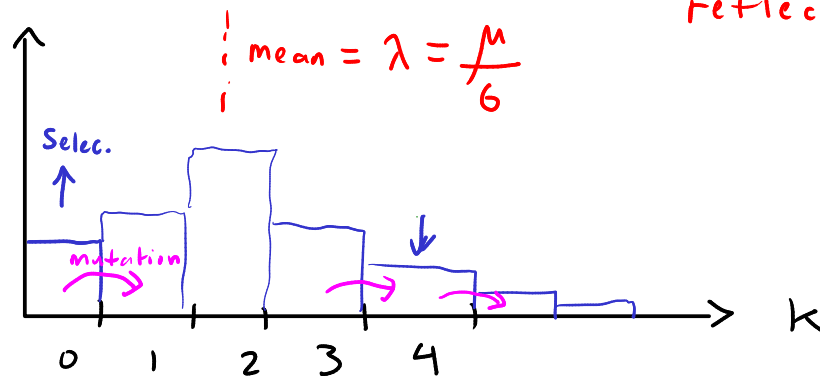
$$\overbrace{-\mu} + \mu$$

stationary distrib. is not unique.

Any shifted Poisson works:

$$x_k = \begin{cases} 0 & \text{for } k \leq k_{\min} \\ \frac{e^{-\lambda} \lambda^{k-k_{\min}}}{(k-k_{\min})!} & \text{for any integer } k_{\min} \geq 0 \end{cases}$$

$k_{\min} = 0$ solution:



reflects mutation-selection balance

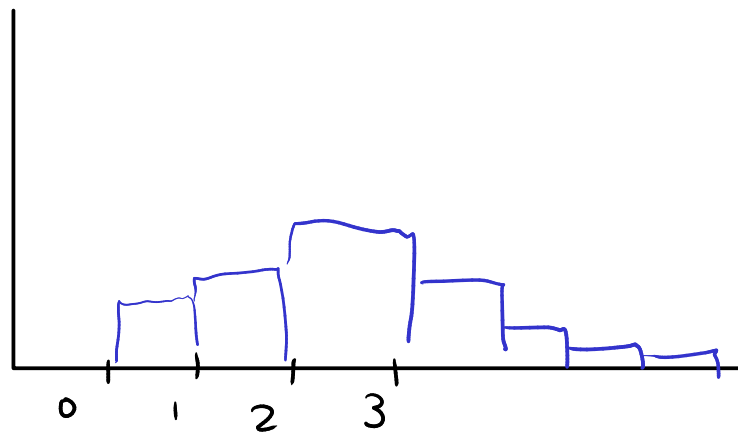
$k = k_{\min}$ class has fraction $x_{k_{\min}} = e^{-\lambda} = e^{-\mu/6}$

if $\mu \gg 6$ we can have $x_{k_{\min}}$ quite small

if $x_{k_{\min}}$ is small enough & we add stochastic fluctuations, $n_{k_{\min}}$ can fluctuate to zero

loss of $k_{\min}$ category $\Rightarrow$ "click" of ratchet

$$k_{\min} = 1$$



loss of $k_{\min} = 1 \Rightarrow$ another "click"

w/ enough clicks, all f_k become negative
 $\Rightarrow$ whole population goes extinct
 $\Rightarrow$ "mutational meltdown"

What about small rate of adaptive mutations going in other direction? $\left. \vphantom{\begin{matrix} \text{adaptive mutations} \\ \text{going in} \\ \text{other direction?} \end{matrix}} \right\} \begin{matrix} \text{ad.} \\ \text{rate} \\ \mu \end{matrix}$

$$\frac{dx_k}{dt} = (f_k - \bar{f}) x_k - \mu x_k + \mu x_{k-1} - \nu x_k + \nu x_{k+1}$$