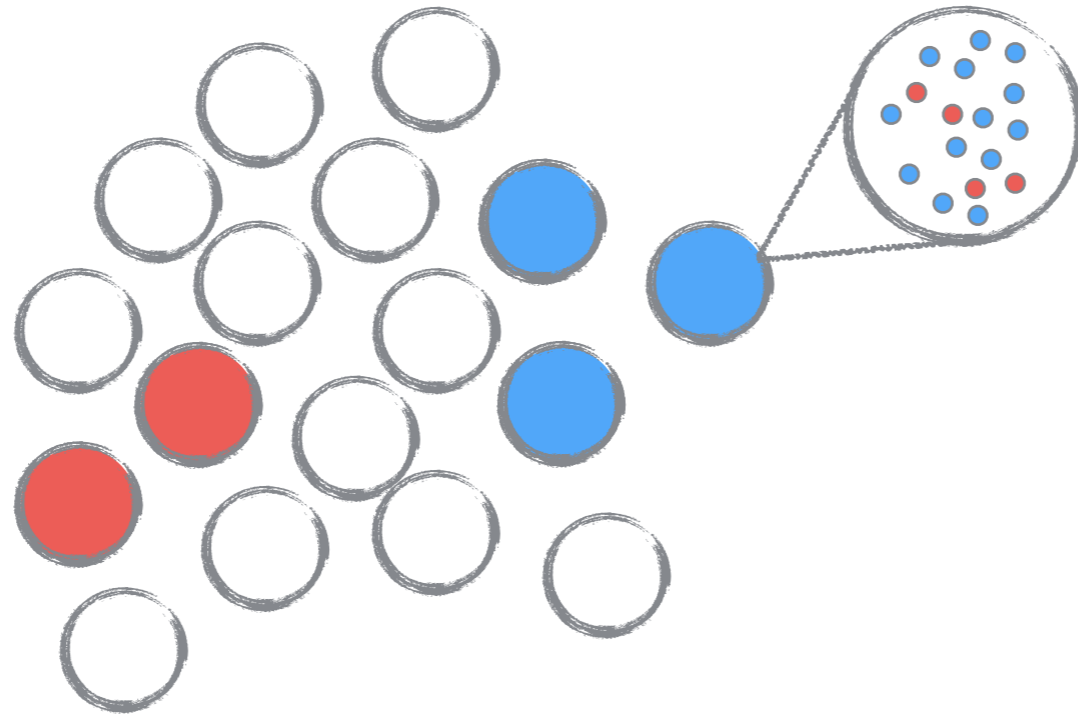


Probabilistic models for pathogen evolution within and between hosts

Shishi Luo
UC Berkeley

Pathogen evolution



- Influenza
- HIV
- General two-level selection

Antigenic drift of influenza

(with K Koelle, JC Mattingly, MC Reed)

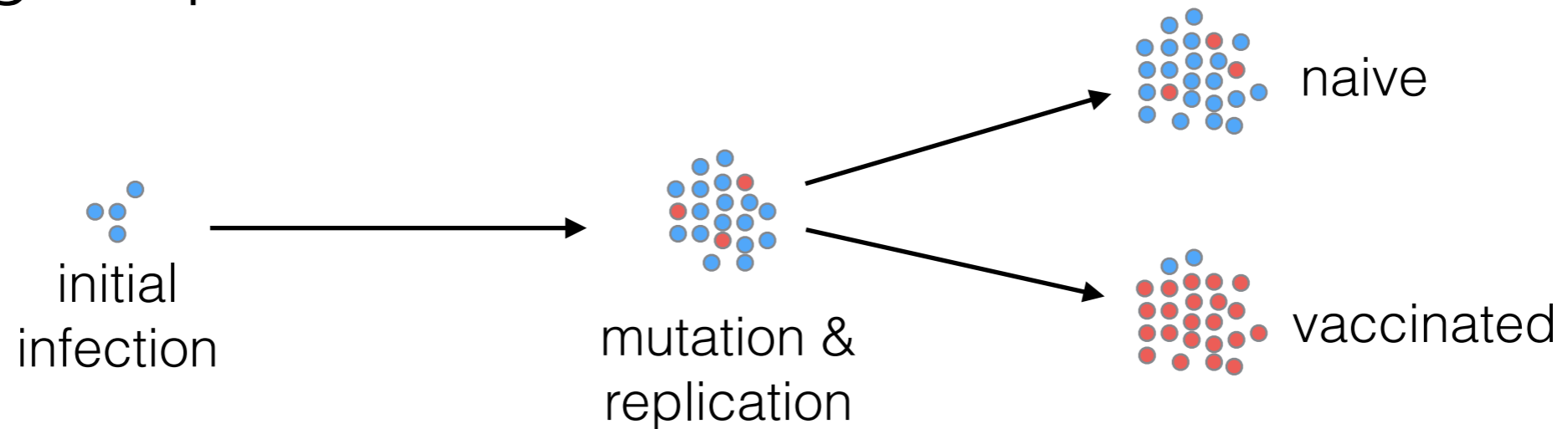


Vaccinate

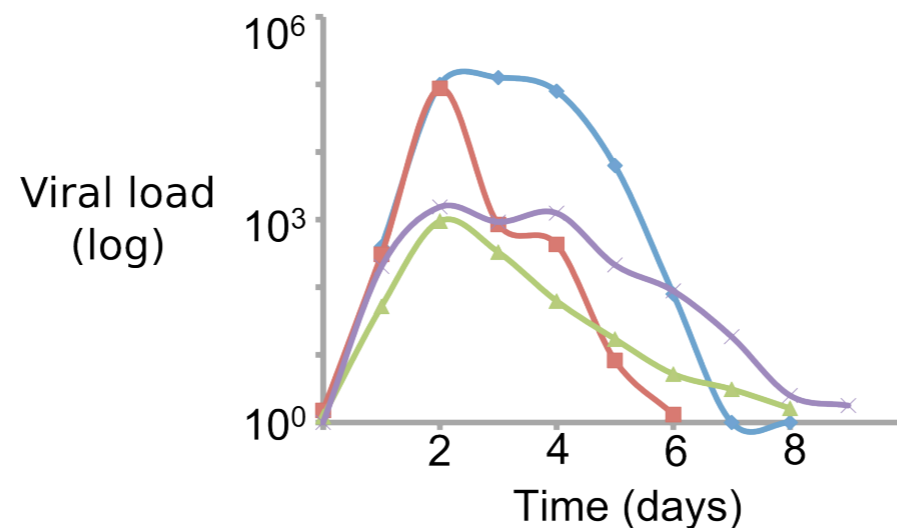


Viral dynamics and evolution

Passage experiments



Within-host influenza viral dynamics (colors are different studies)



ODEs with Poisson mutation

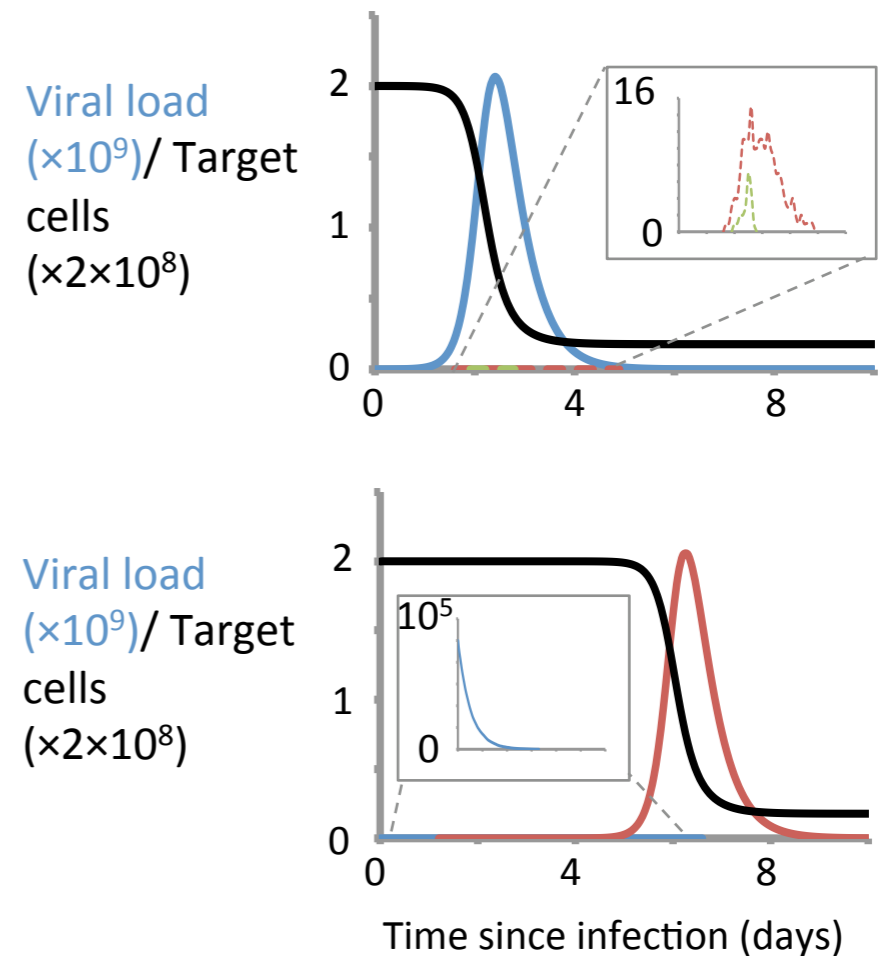
$$\dot{C} = -\beta C V_r - \beta C V_m$$

$$\dot{V}_r = \gamma C V_r - \delta_r V_r$$

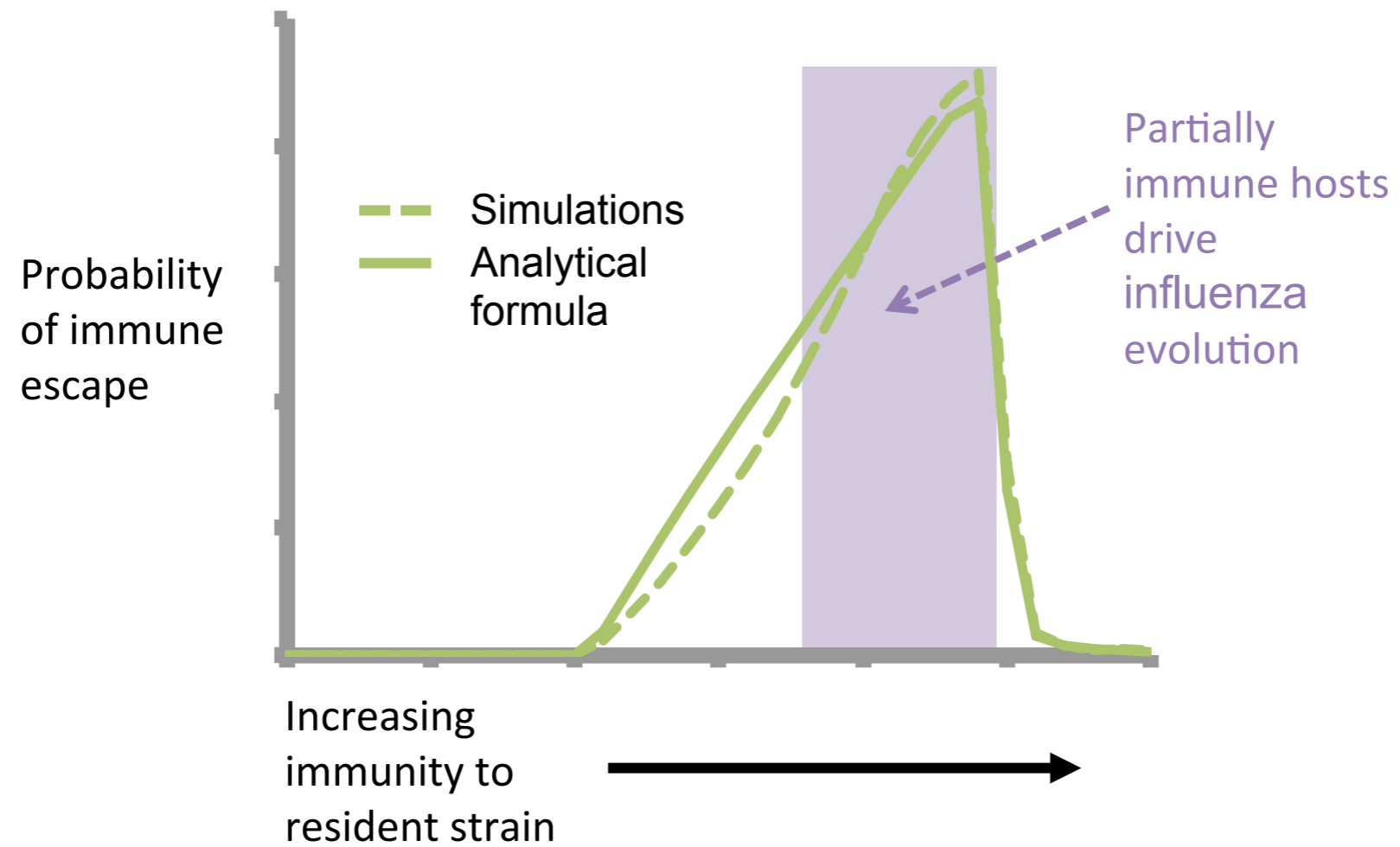
$$\dot{V}_m = \gamma C V_m - \delta_m V_m$$

$$V_r(0) = V_0 \quad C(0) = C_0$$

mutations occur at rate $\mu V_r(t)$

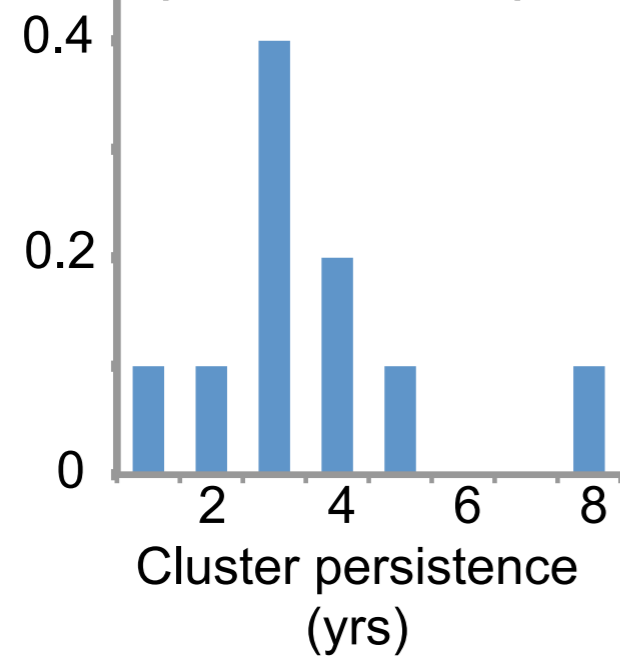


Results: Within hosts

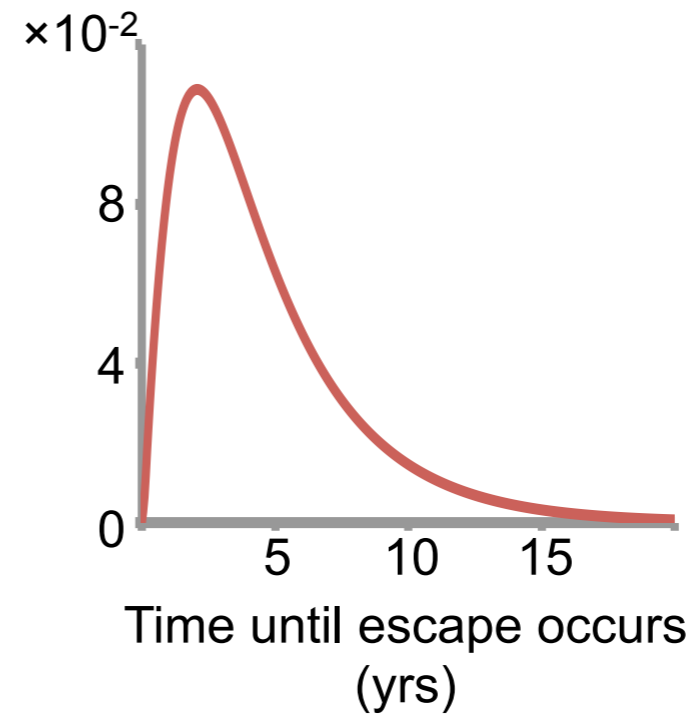


Results: Population level

**Proportion of clusters
(for influenza)**



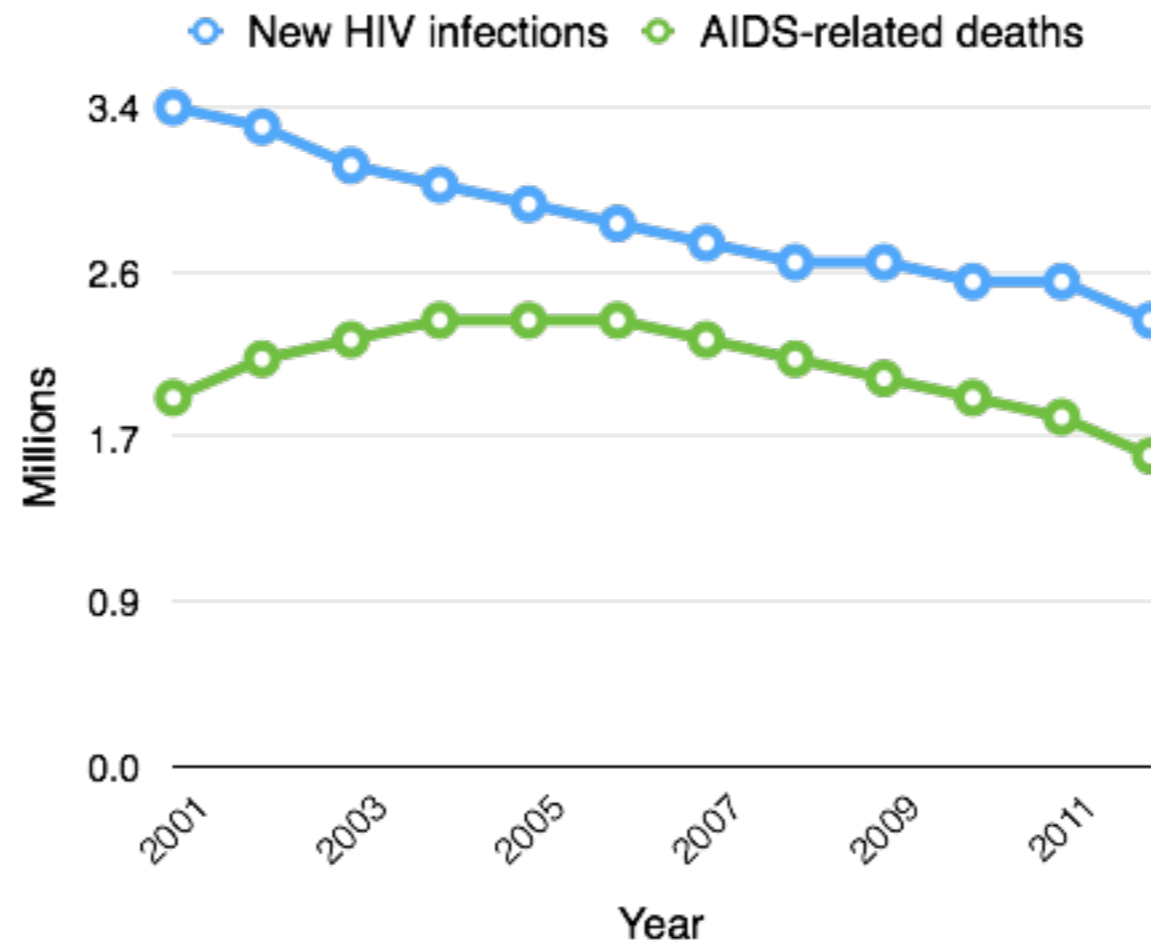
Probability density



Koelle et al *Epidemics* 2009

Broadly neutralizing HIV antibodies

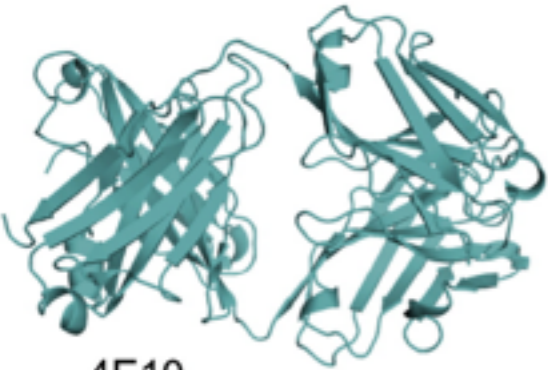
(with A Perelson)



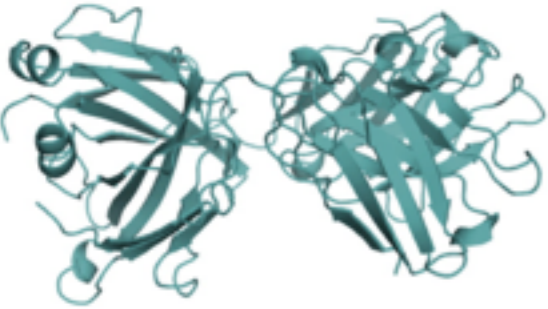
Data from UNAIDS Global fact sheet 2012

HIV-1 viral spike EM
Glycosylated
gp120 model

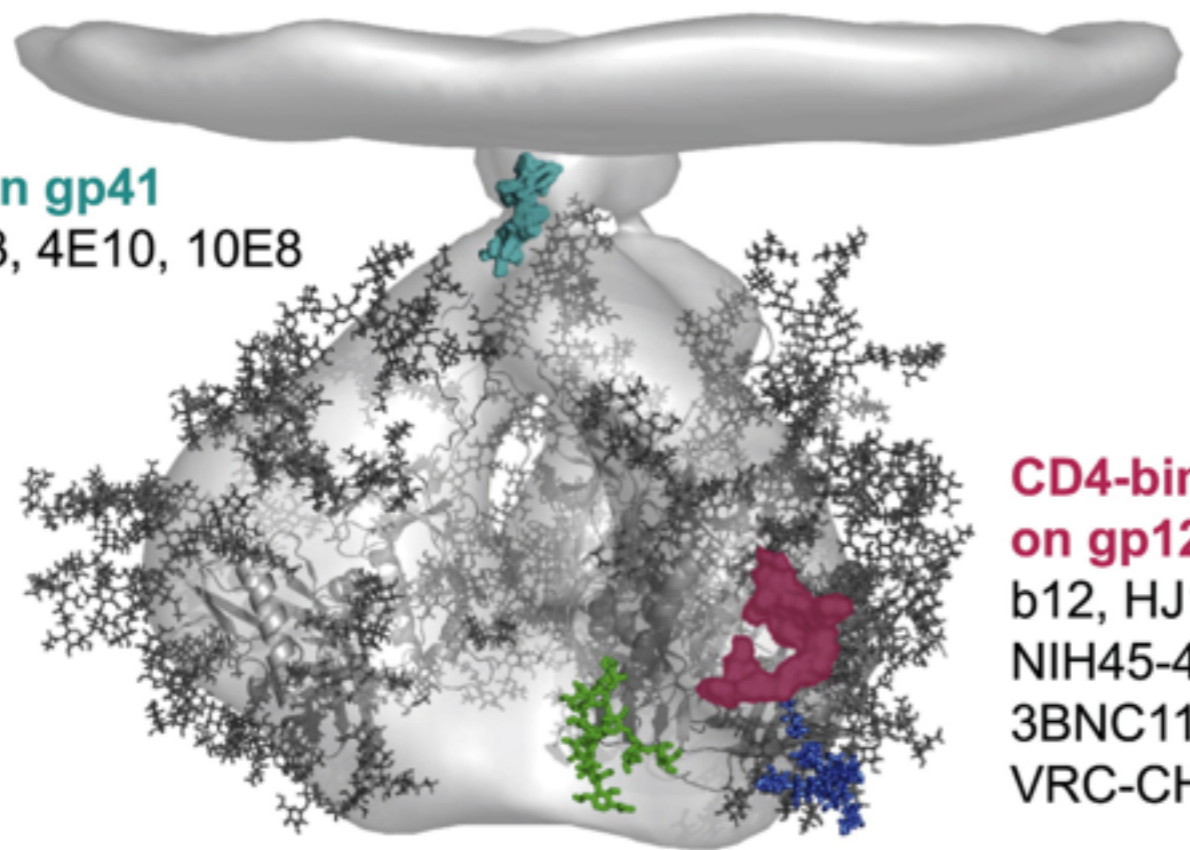
MPER on gp41
2F5, Z13, 4E10, 10E8



4E10



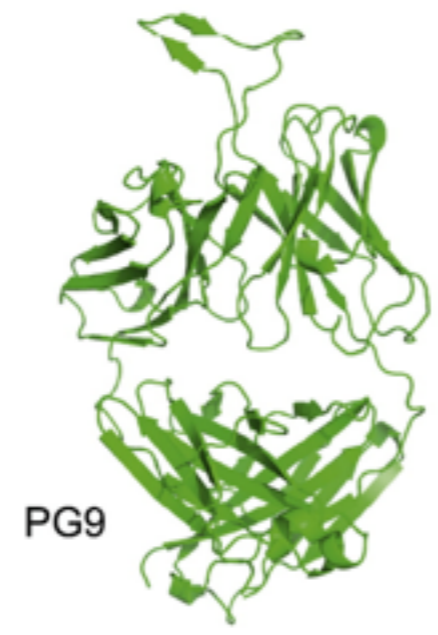
10E8



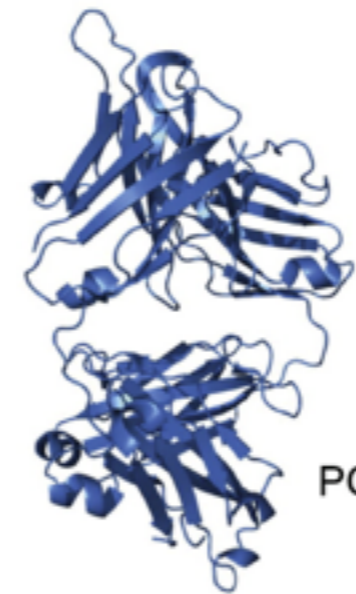
**CD4-binding site
on gp120**
b12, HJ16, VRC01,
NIH45-46, 12A12,
3BNC117, VRC-PG04,
VRC-CH31

V1V2 site on gp120
PG9, PG16, CH01-04,
PGT141-145

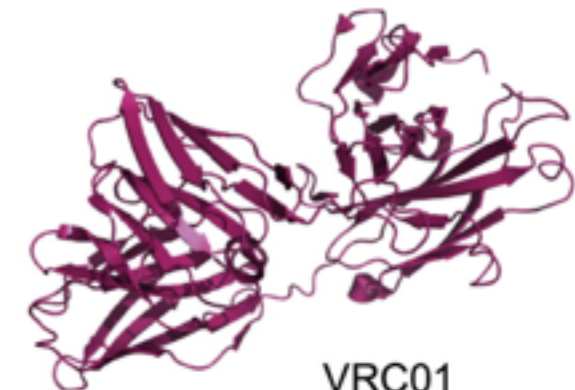
**Glycan-V3 site
on gp120**
2G12, PGT121-137



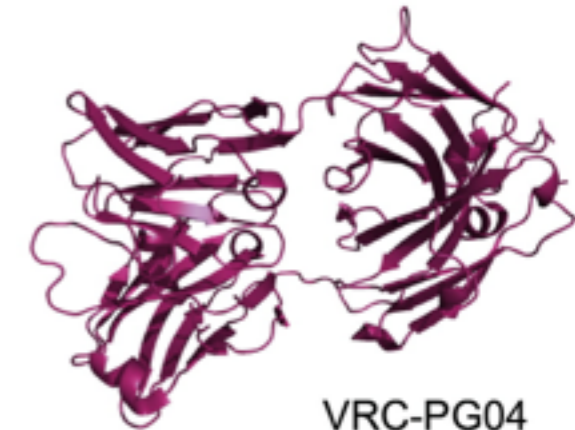
PG9



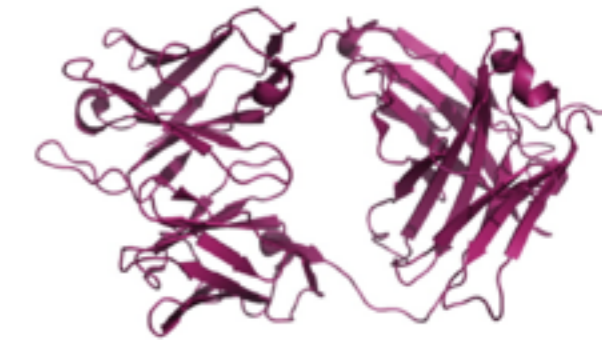
PGT128



VRC01

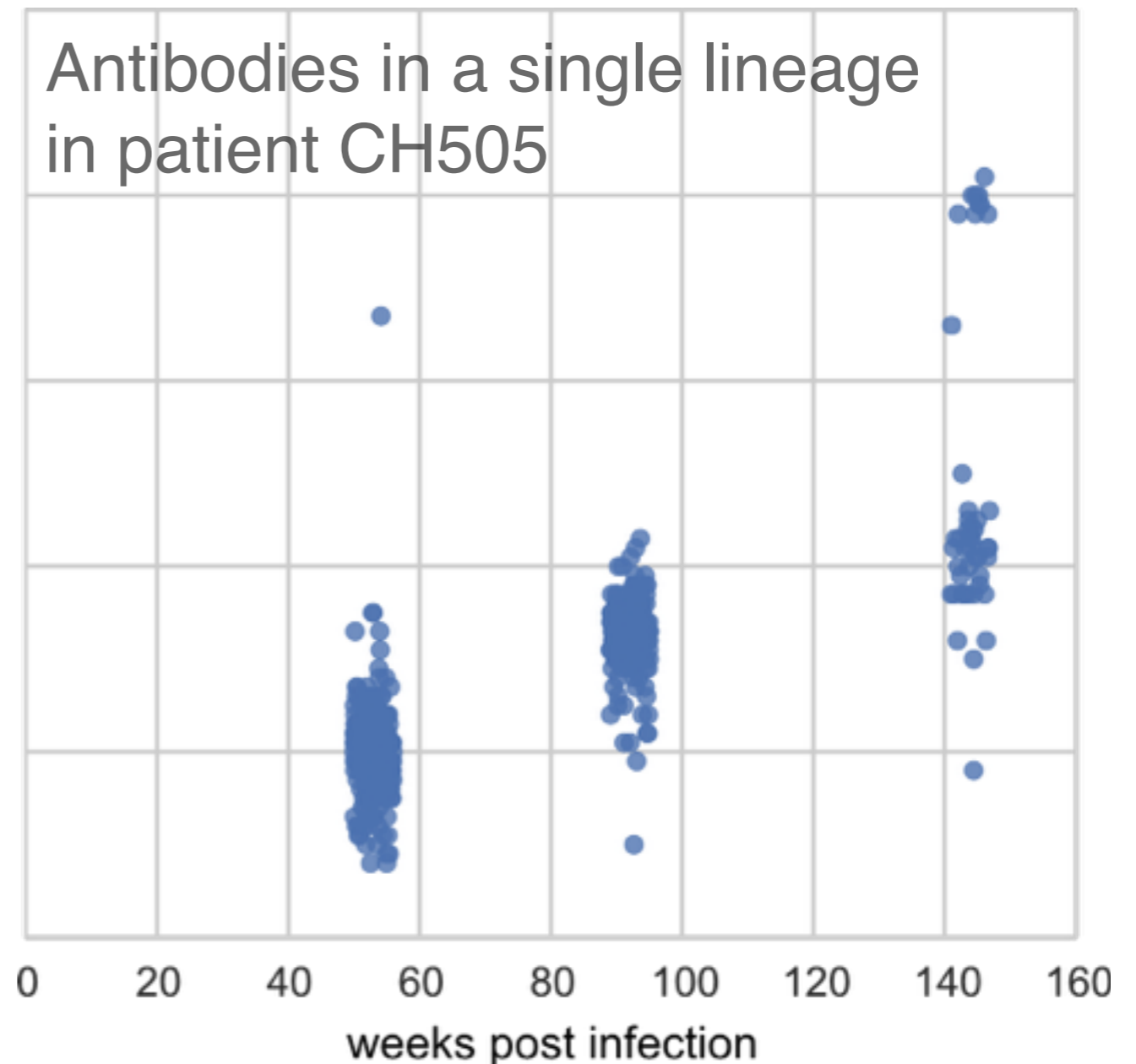
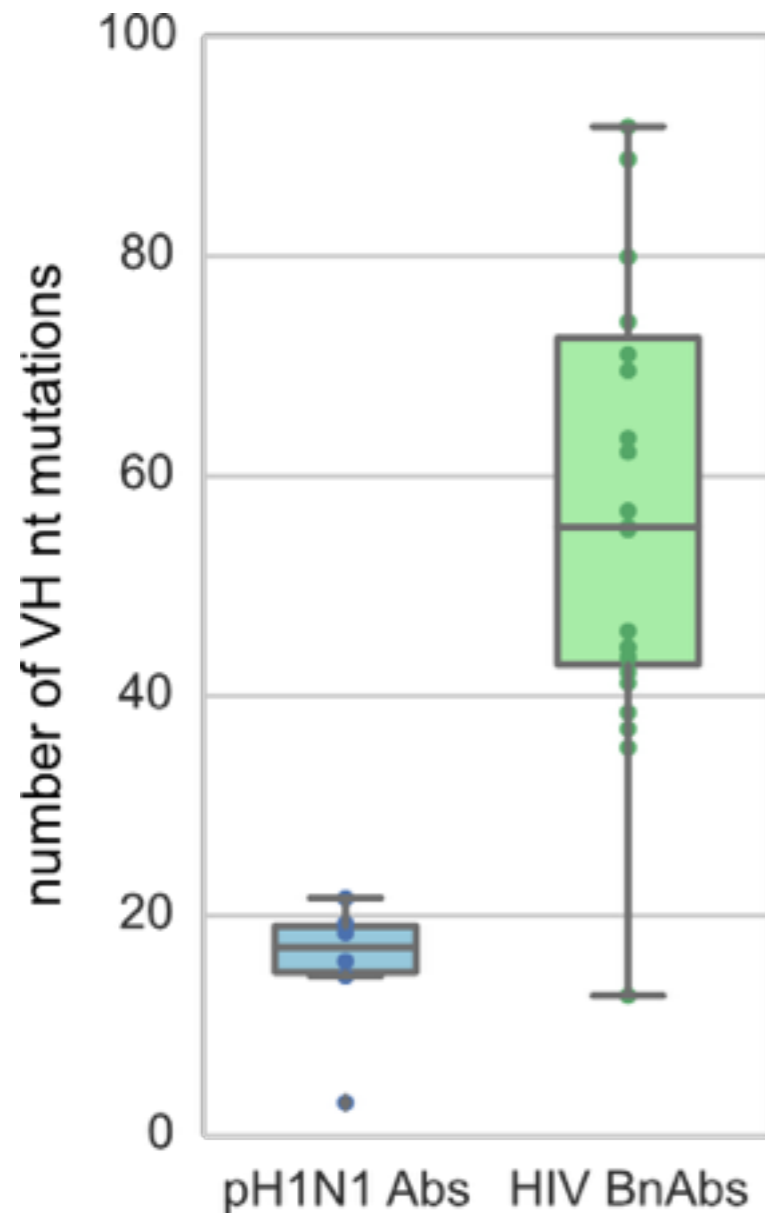


VRC-PG04

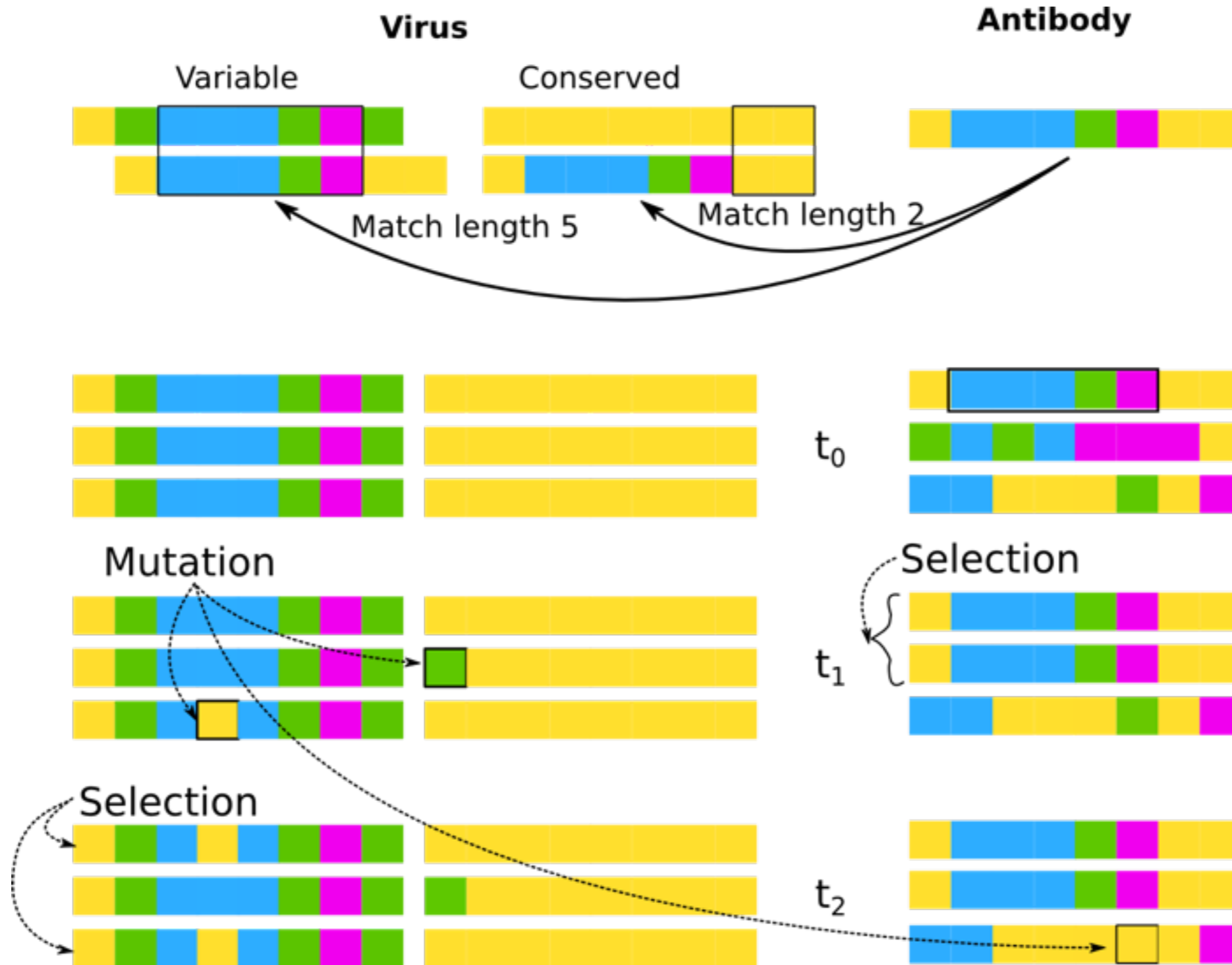


NIH45-46

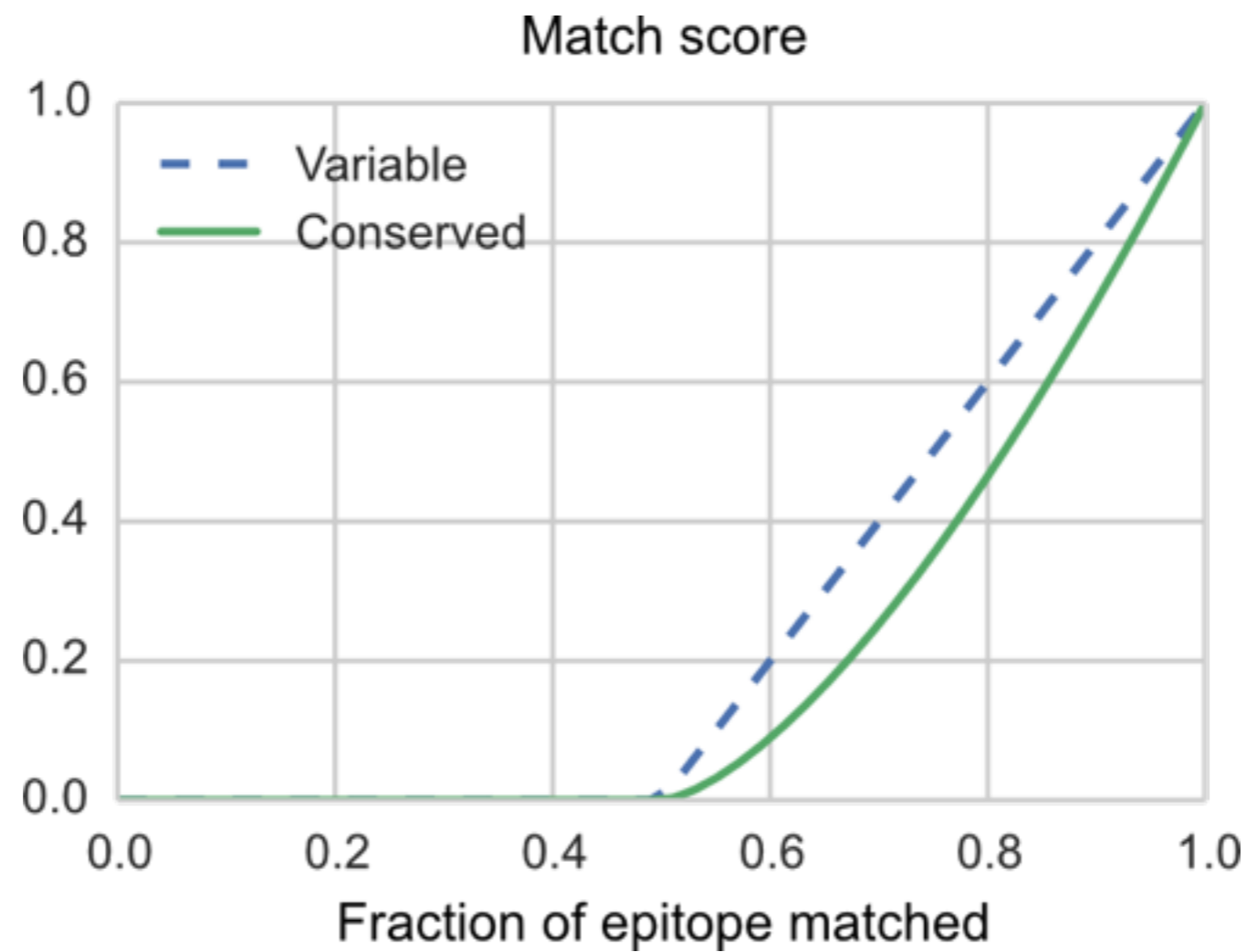
Broadly neutralizing HIV antibodies are highly mutated



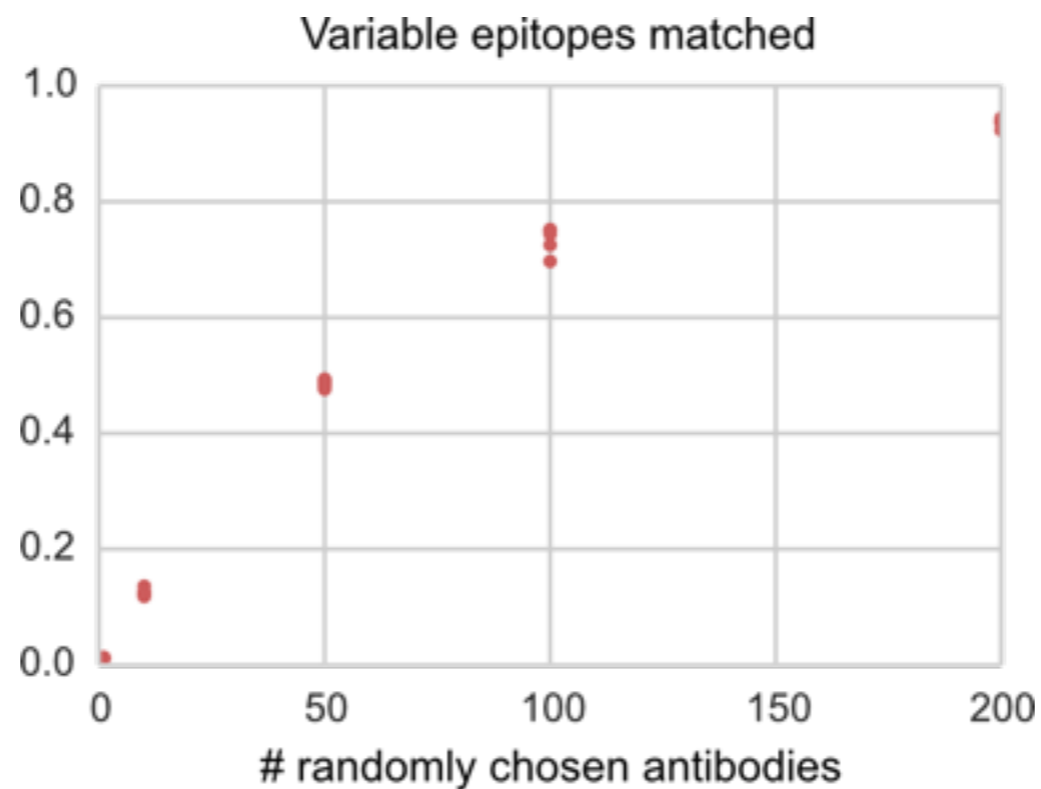
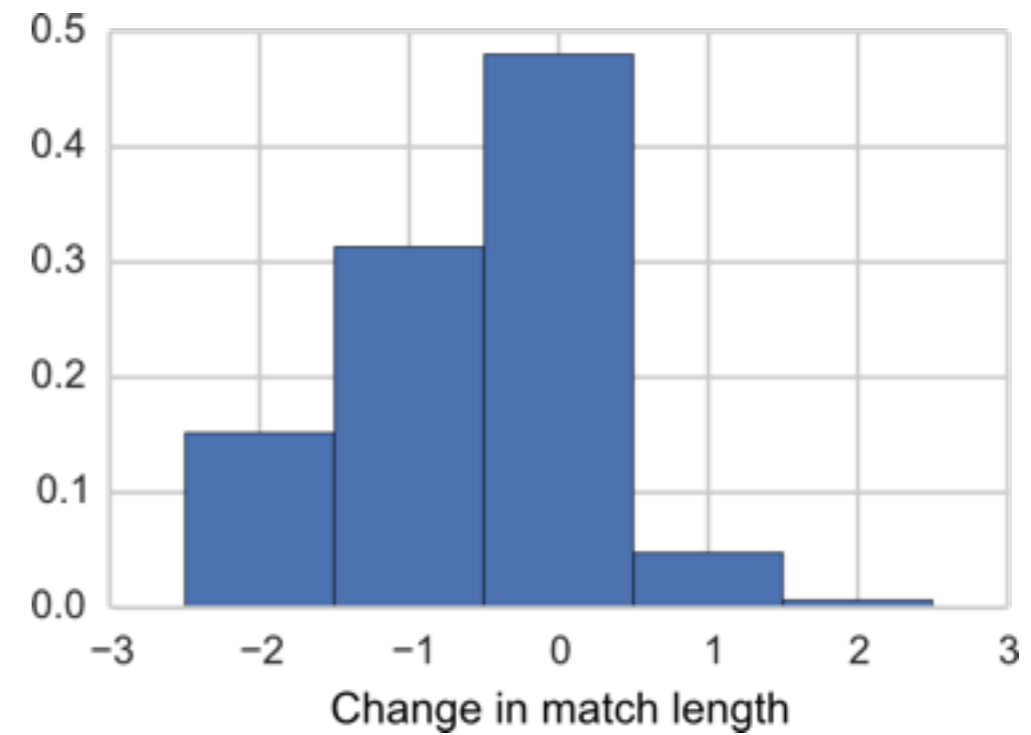
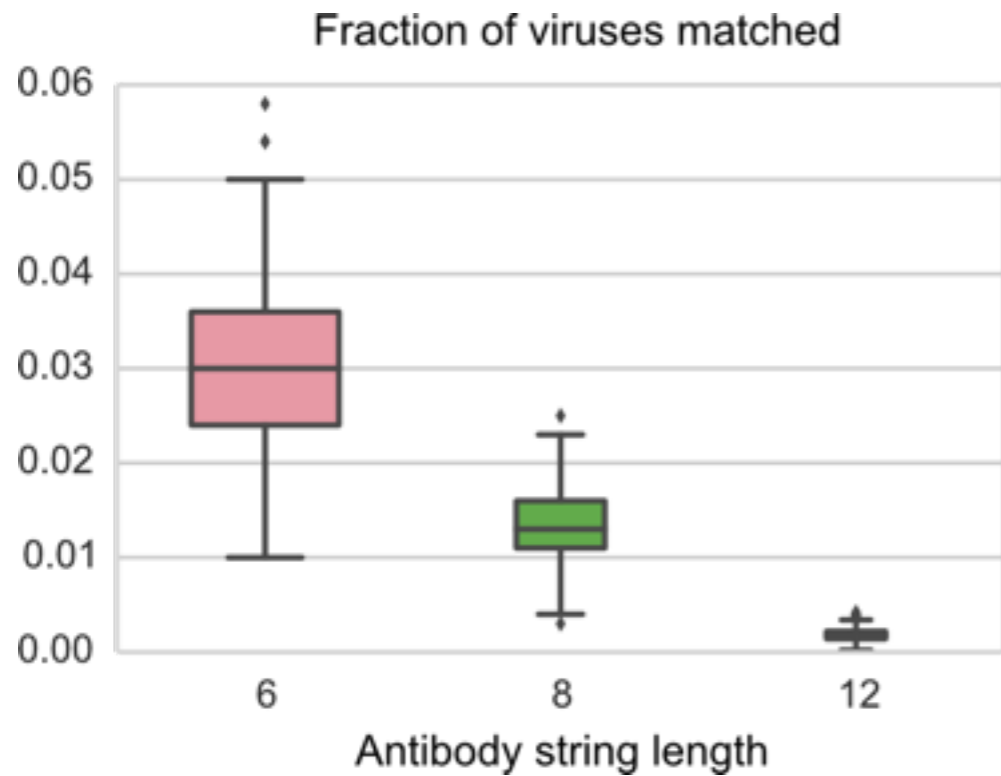
'Bitstring' model



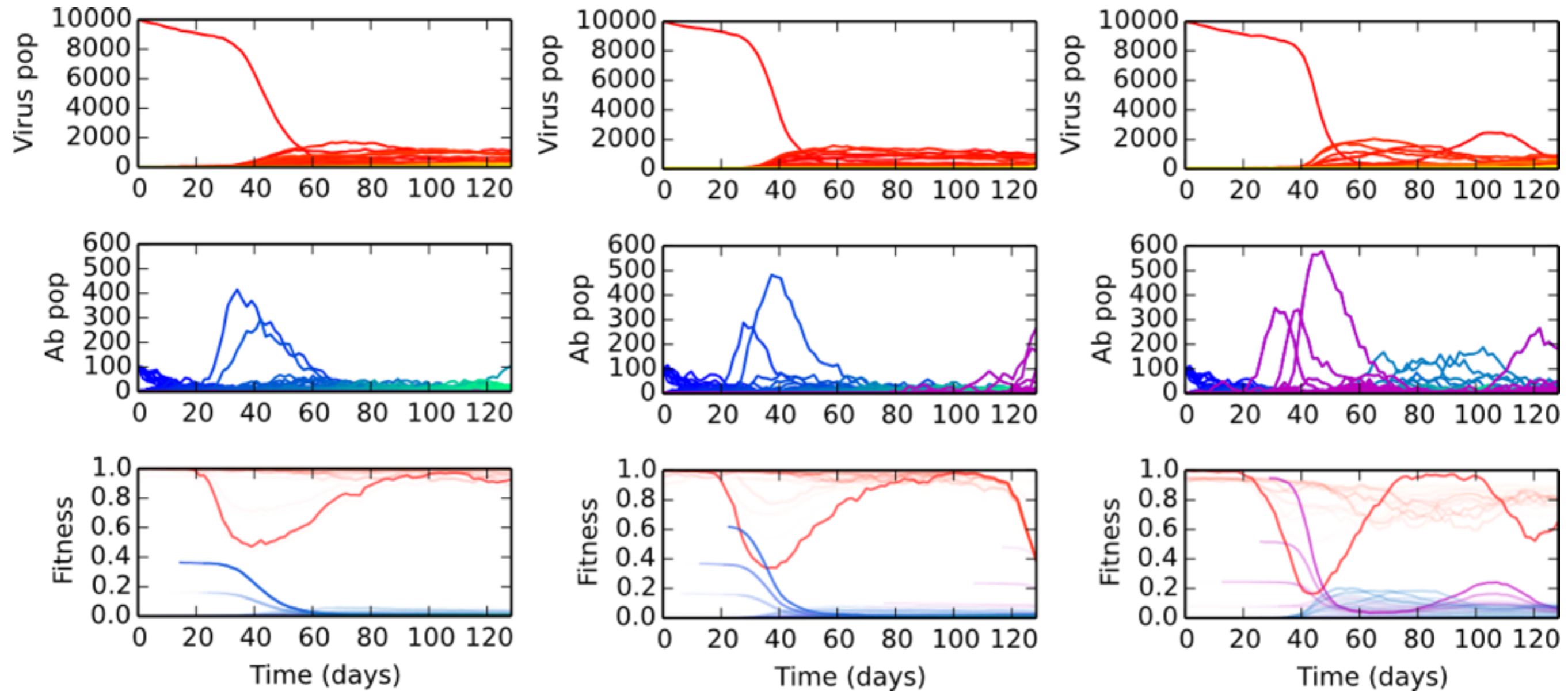
Conserved region is less accessible



Binding properties

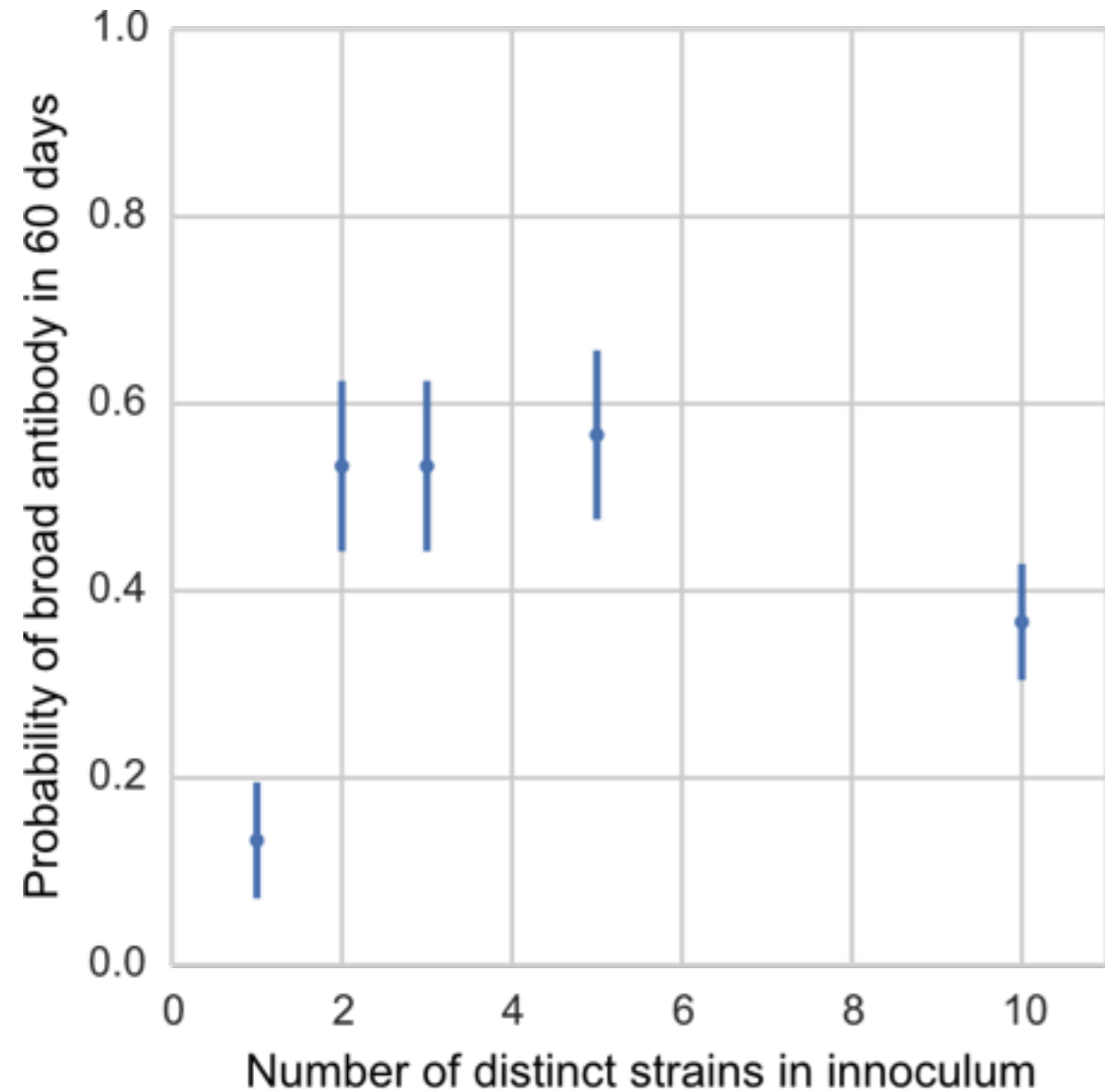


Infection with single strain

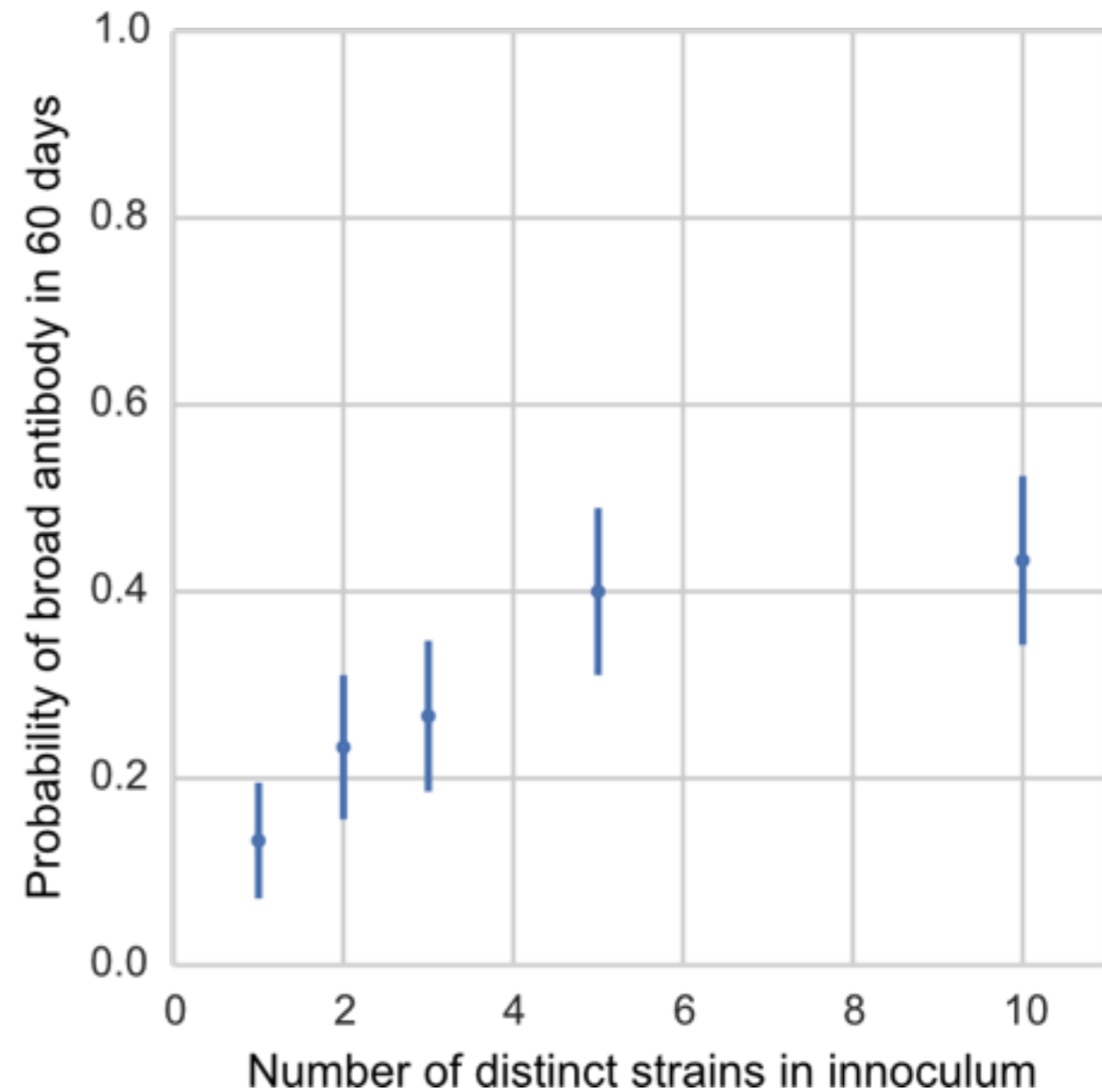


Vaccine strategies

Strains selected uniformly at random



Strains are nearest neighbor mutants

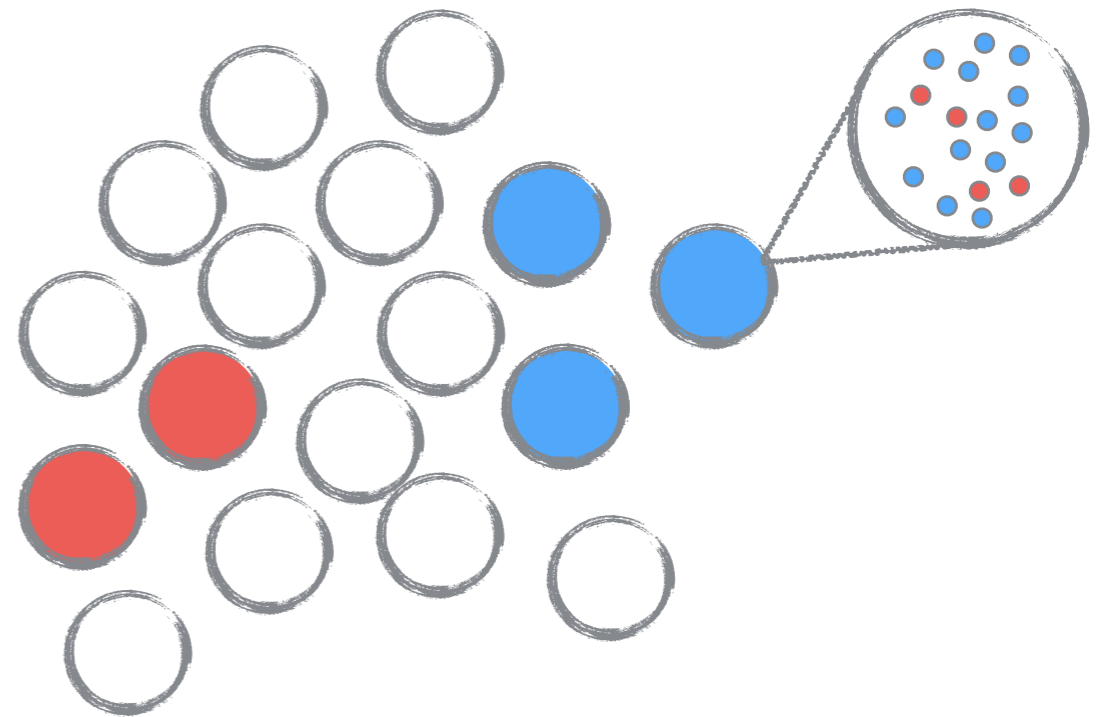


Bars show +/- 1 SE

Multilevel selection in host-pathogen systems

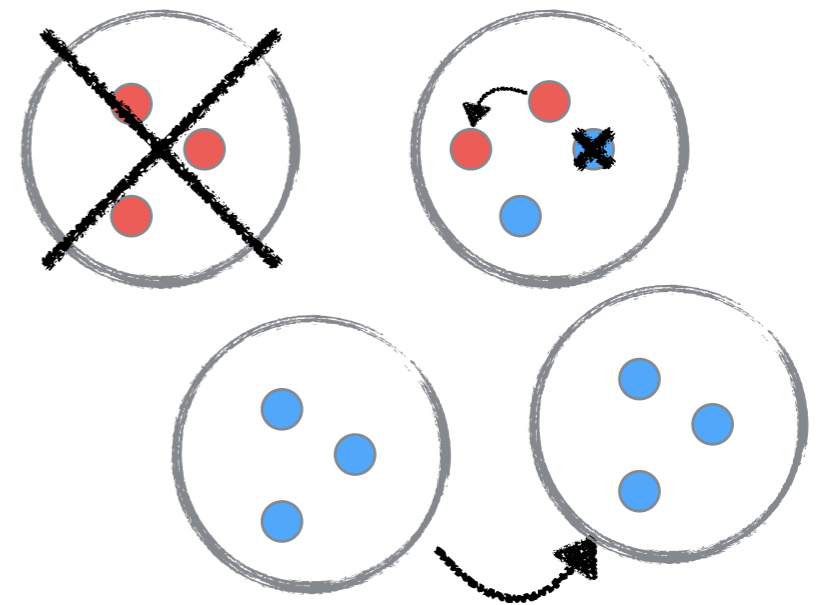
(with JC Mattingly)

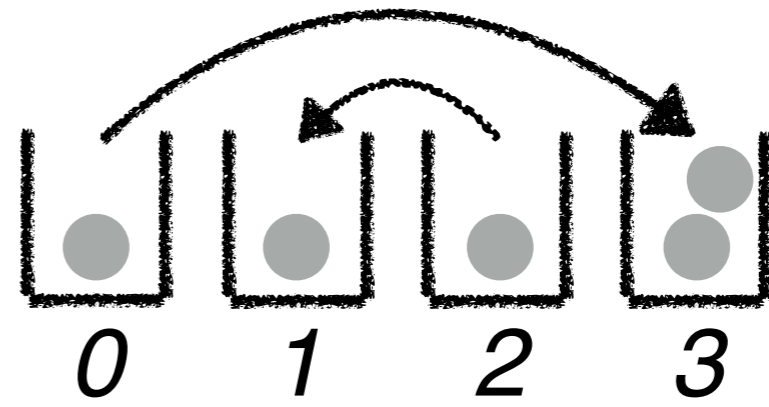
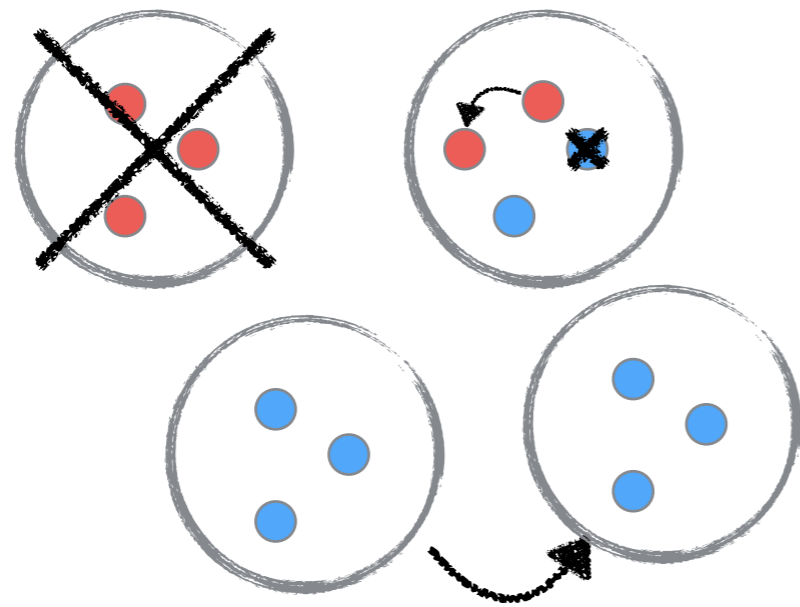
- myxomatosis-rabbit
- human papillomavirus
- malaria

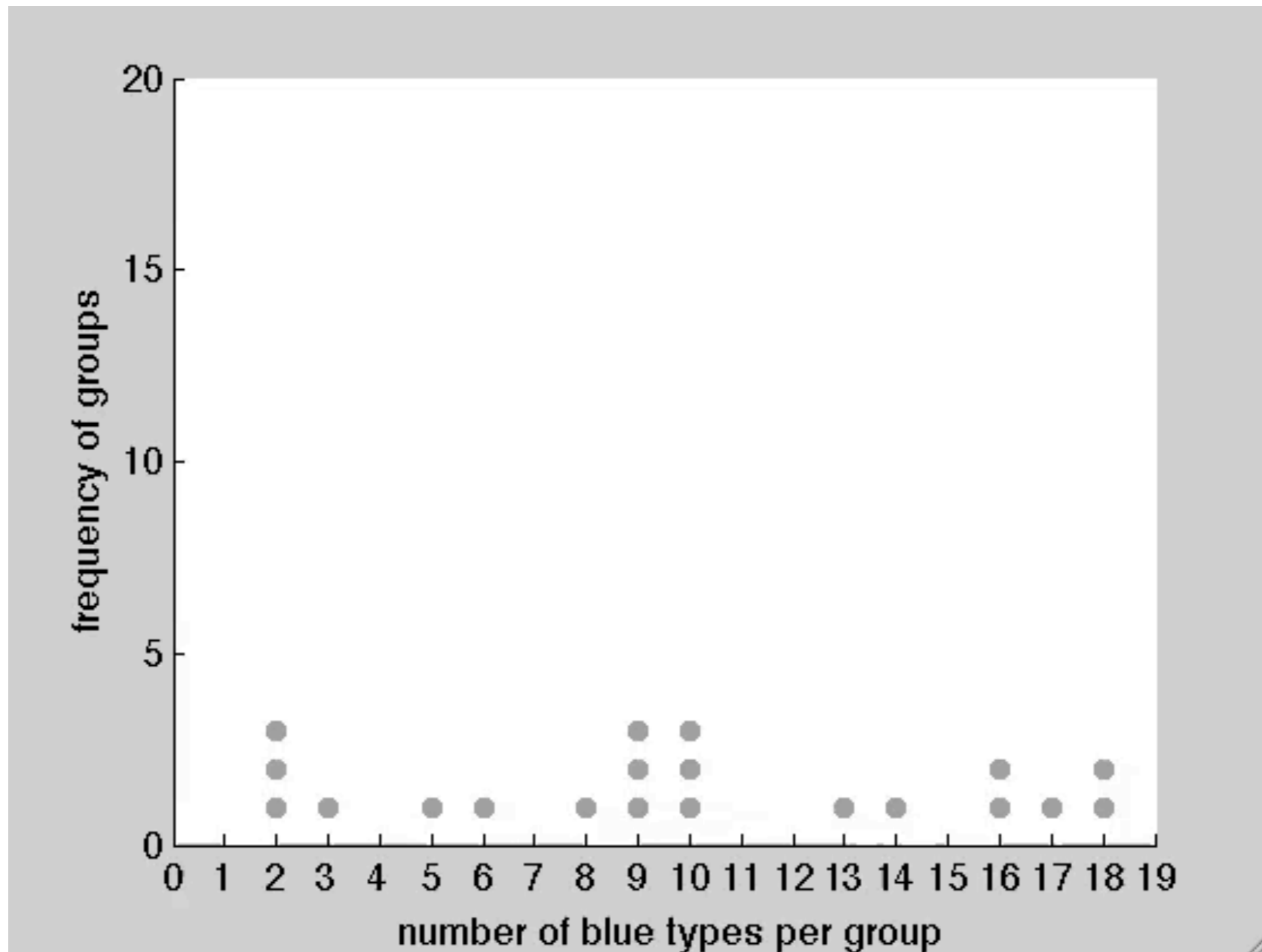


Suppose:

- m groups, n individuals per group
- red beats blue in each group,
 $1 + s$ vs 1
- bluer groups beat redder groups,
 $1 + r \frac{k}{n}$, where $\frac{k}{n}$ is fraction of blue
- $w = \frac{\text{rate of group-level events}}{\text{rate of individual-level events}}$







Recall: blue types beneficial at group level

- Let X_t^j be the position of ball j on the (rescaled) 1-d lattice $\{0, \frac{1}{n}, \dots, 1\}$ at time t .
- Define the measure-valued process:

$$\mu_t^{m,n} = \frac{1}{m} \sum_{j=1}^m \delta_{X_t^j}$$

$\mu_t^{m,n}$ has transition rate matrix $R = R_1 + wR_2$ where:

$$R_1 \left(v, v + \frac{1}{m} (\delta_{\frac{j}{n}} - \delta_{\frac{i}{n}}) \right) = \begin{cases} mv \binom{i}{n} i \left(1 - \frac{i}{n} \right) (1 + s) & \text{if } j = i - 1, i < n \\ mv \binom{i}{n} i \left(1 - \frac{i}{n} \right) & \text{if } j = i + 1, i > 0 \\ 0 & \text{otherwise} \end{cases}$$

$$R_2 \left(v, v + \frac{1}{m} (\delta_{\frac{j}{n}} - \delta_{\frac{i}{n}}) \right) = mv \binom{i}{n} v \binom{j}{n} (1 + r \frac{j}{n})$$

s : individual-level selection, r : group-level selection, w : relative rate of events, m : number of groups, n : number of individuals in each group

The deterministic limit of $\mu_t^{m,n}$:

$$\frac{\partial}{\partial t} \mu(t, x) = s \frac{\partial}{\partial x} [x(1-x)\mu(t, x)] + wr\mu(t, x) \left[x - \int_0^1 y\mu(t, y) dy \right]$$

The stochastic (Fleming-Viot) limit:

$$\begin{aligned} \partial_t \nu_t = & \sigma \partial_x [x(1-x)\nu_t] + \partial_{xx} [x(1-x)\nu_t] \\ & + w\alpha\rho\nu_t \cdot \left[x - \int_0^1 y\nu_t(y) dy \right] + w\alpha\sqrt{2\nu_t(1-\nu_t)}\dot{W}_t \end{aligned}$$

Long-term behavior of PDE

Suppose μ_0 is a density and k^* is such that

$$\mu_0^{(k)}(1) = 0 \text{ for all integers } k < k^* \text{ and } \mu_0^{(k^*)}(1) \neq 0$$

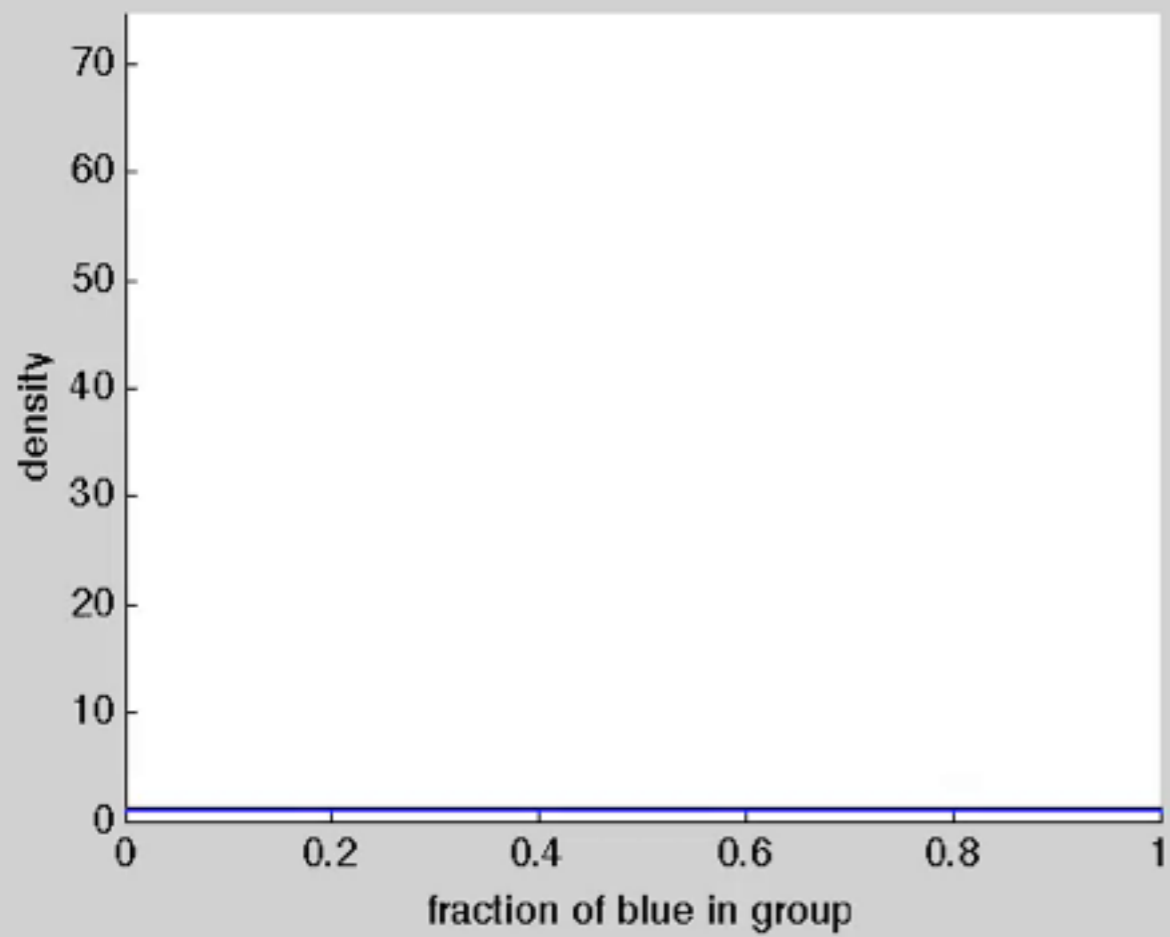
If $\lambda - 1 > k^*$,

$$\mu(t, x) \rightarrow \text{Beta}(\lambda - k^* - 1, k^* + 1) \text{ as } t \rightarrow \infty$$

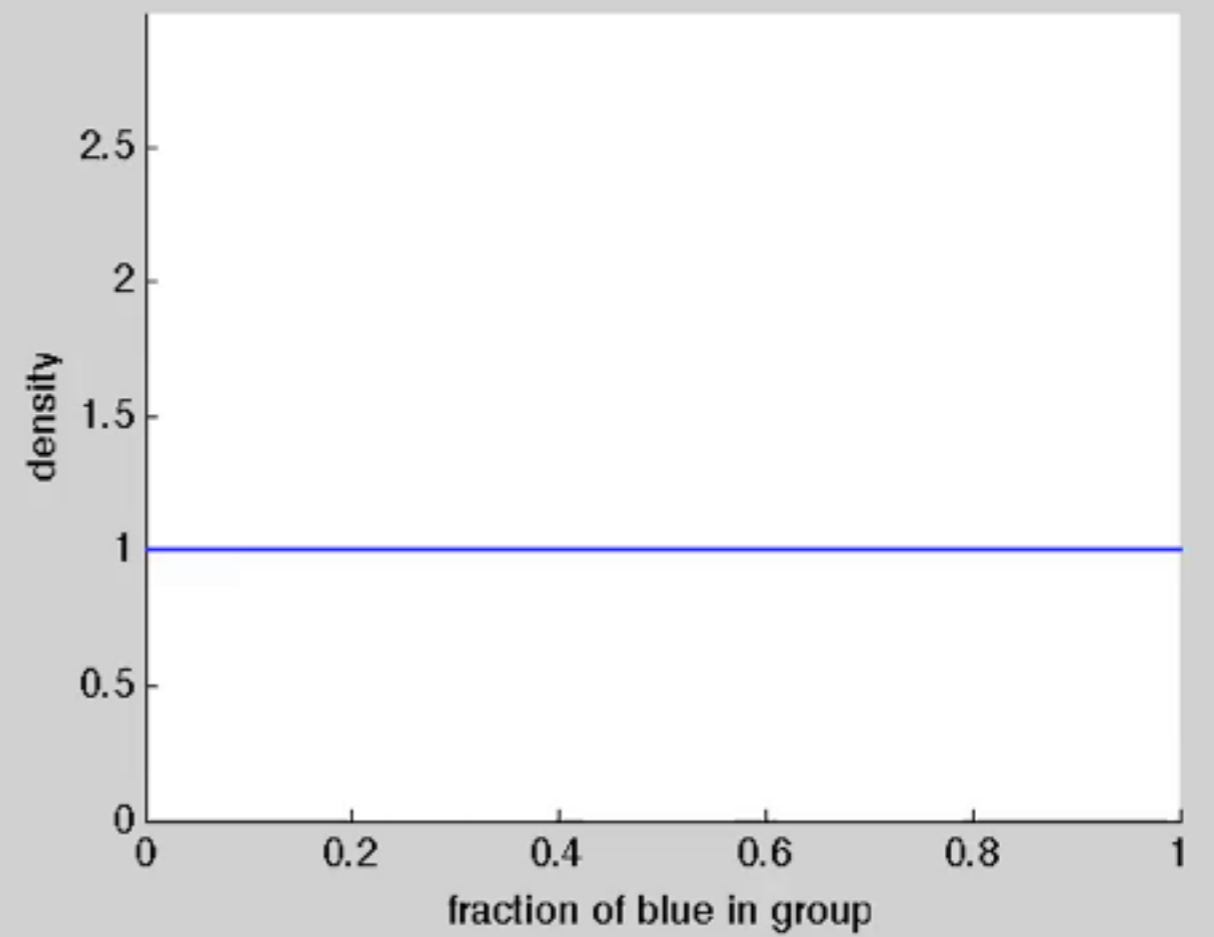
If $\lambda - 1 \leq k^*$,

$$\mu(t, x) \rightarrow \delta_0(x) \text{ as } t \rightarrow \infty$$

$$\lambda = 1.5$$



$$\lambda = 4$$

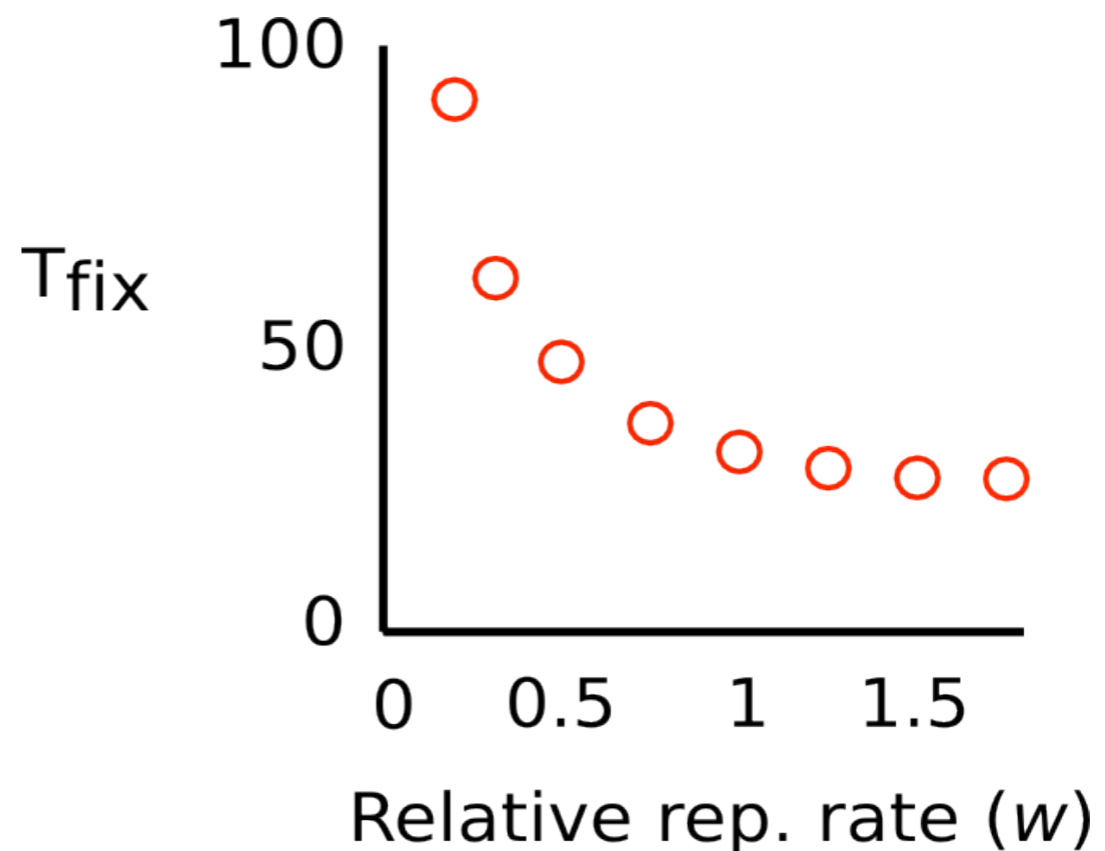


Properties of particle system

Relative rate of replication acts like group-level selection

$$\frac{\partial}{\partial t} \mu(t, x) = s \frac{\partial}{\partial x} [x(1-x)\mu(t, x)] + wr\mu(t, x) \left[x - \int_0^1 y\mu(t, y) dy \right]$$

$$\lambda := \frac{wr}{s}$$



What if m and n are finite?

$$\frac{\partial}{\partial t} \mu(t, x) = s \frac{\partial}{\partial x} [x(1-x)\mu(t, x)] + wr\mu(t, x) \left[x - \int_0^1 y\mu(t, y) dy \right]$$

