

Some implications of neutral evolution for ecology

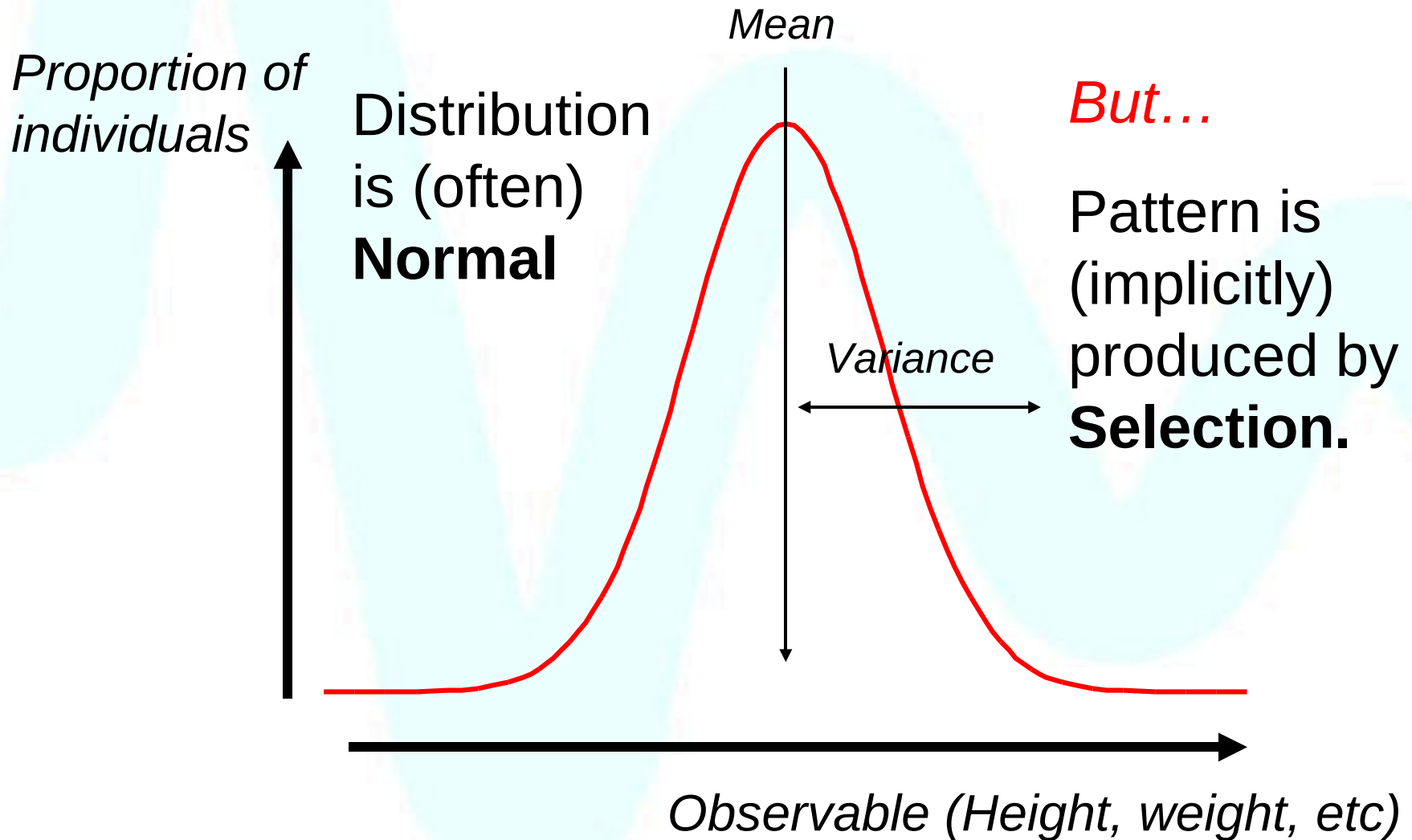
Daniel Lawson

Bioinformatics and Statistics Scotland

Work with

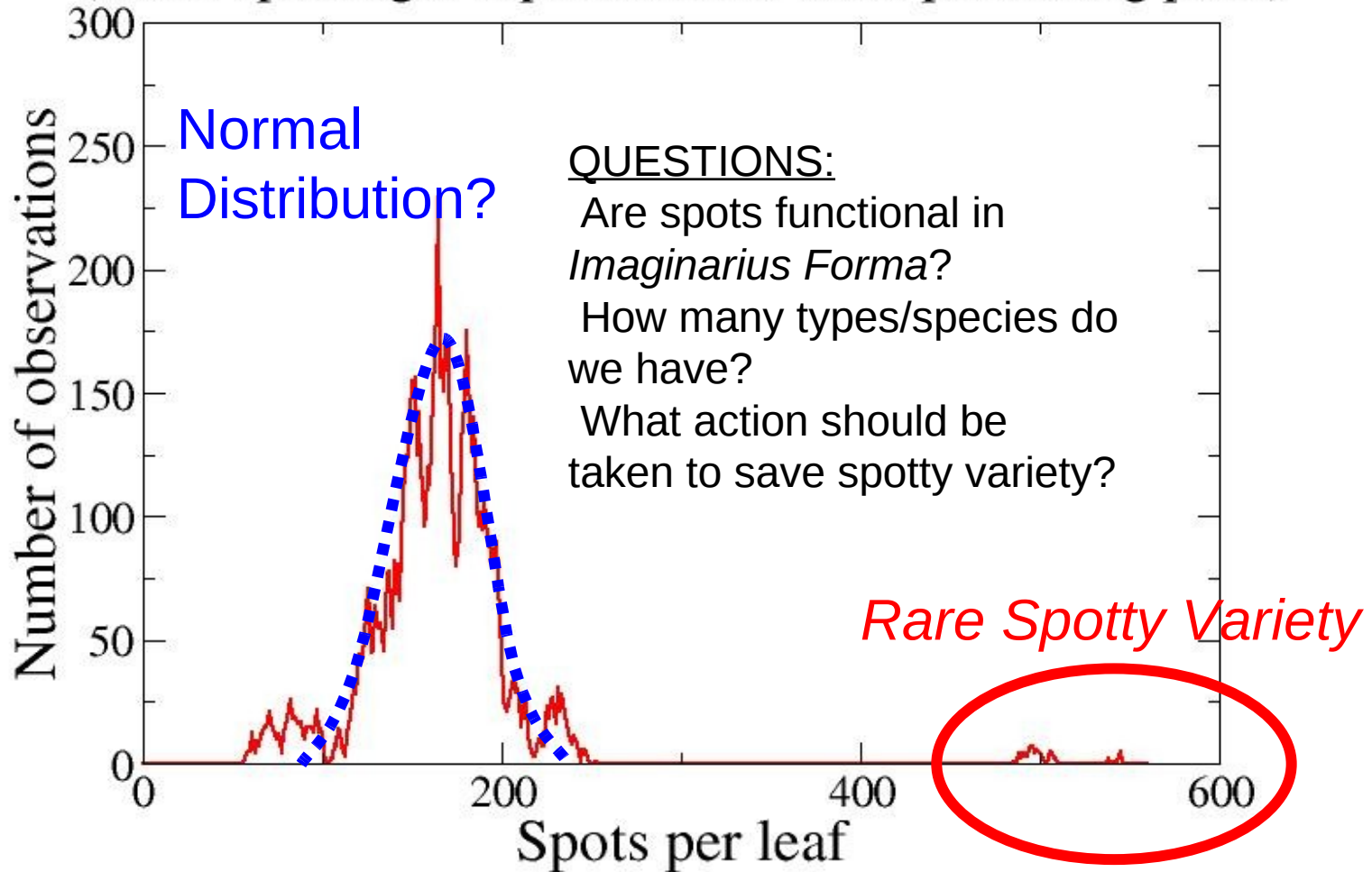
Henrik Jeldtoft Jensen,
Imperial College London

Characterising a Species



Test Problem

Observations of number of spots per leaf of *Imaginarium Forma*
(Made up thought experiment for a self pollinating plant)

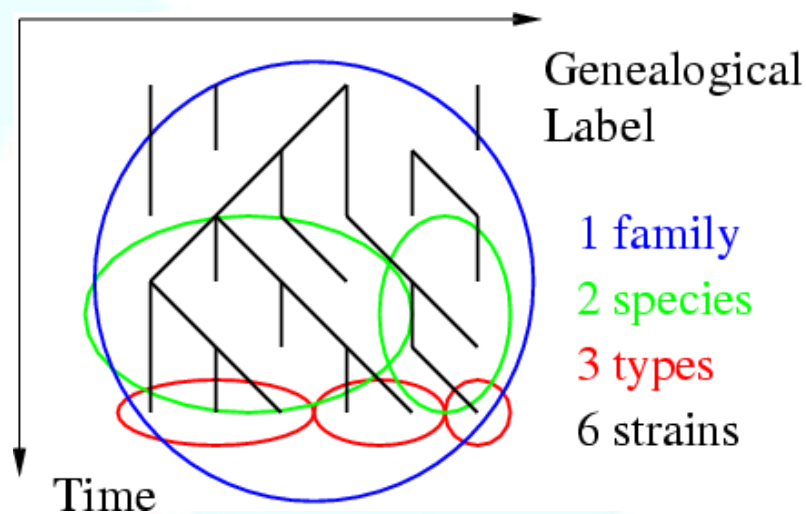


Defining Diversity

- Ecological Sense: “number” of different species or types

- Requires definition of species:

- *Biological Species concept?*
- *Phenotypically distinct?*
- *Genotypic species concept?*



- “Species” don't exist, but individuals form clusters
- Evolution definition of diversity: number of different clusters on a chosen threshold

The Neutral Model

- Assume that all individuals are 'equal'
 - Valid for Phenotypes that do not have function
 - Genotypes, considering the part of the genome that does not code for protein synthesis (**12% of Human DNA is variable!** *Redon et al. Nature. doi:10.1038/nature05329*)
- Each individual has the same probability to die (p_k), or give birth (p_b), in a time step
 - For simplicity, assume the total population (N) has reached equilibrium ($p_k = p_b$)
- Mutations (and/or colonisation) can occur, reproduction is asexual

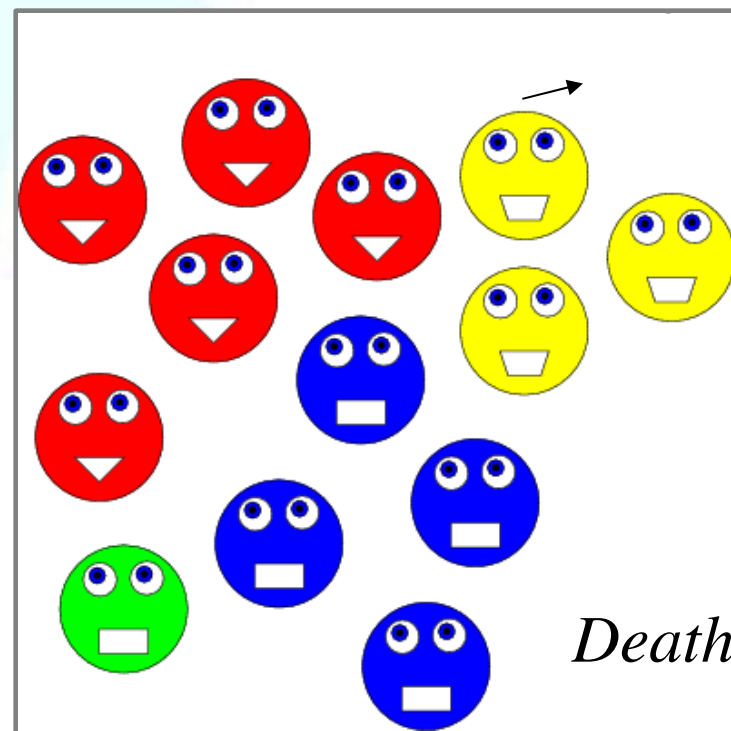
The ecological model

- Consider N individuals each labeled by species:

Birth or Colonise

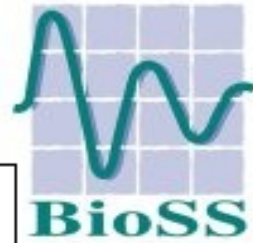
TIMESTEP:

- Pick an individual (from N) and mark it to die.
- Pick an individual (from N) and copy it, *or* with probability p_m , **colonise with a new species.**
- Kill the marked individual.

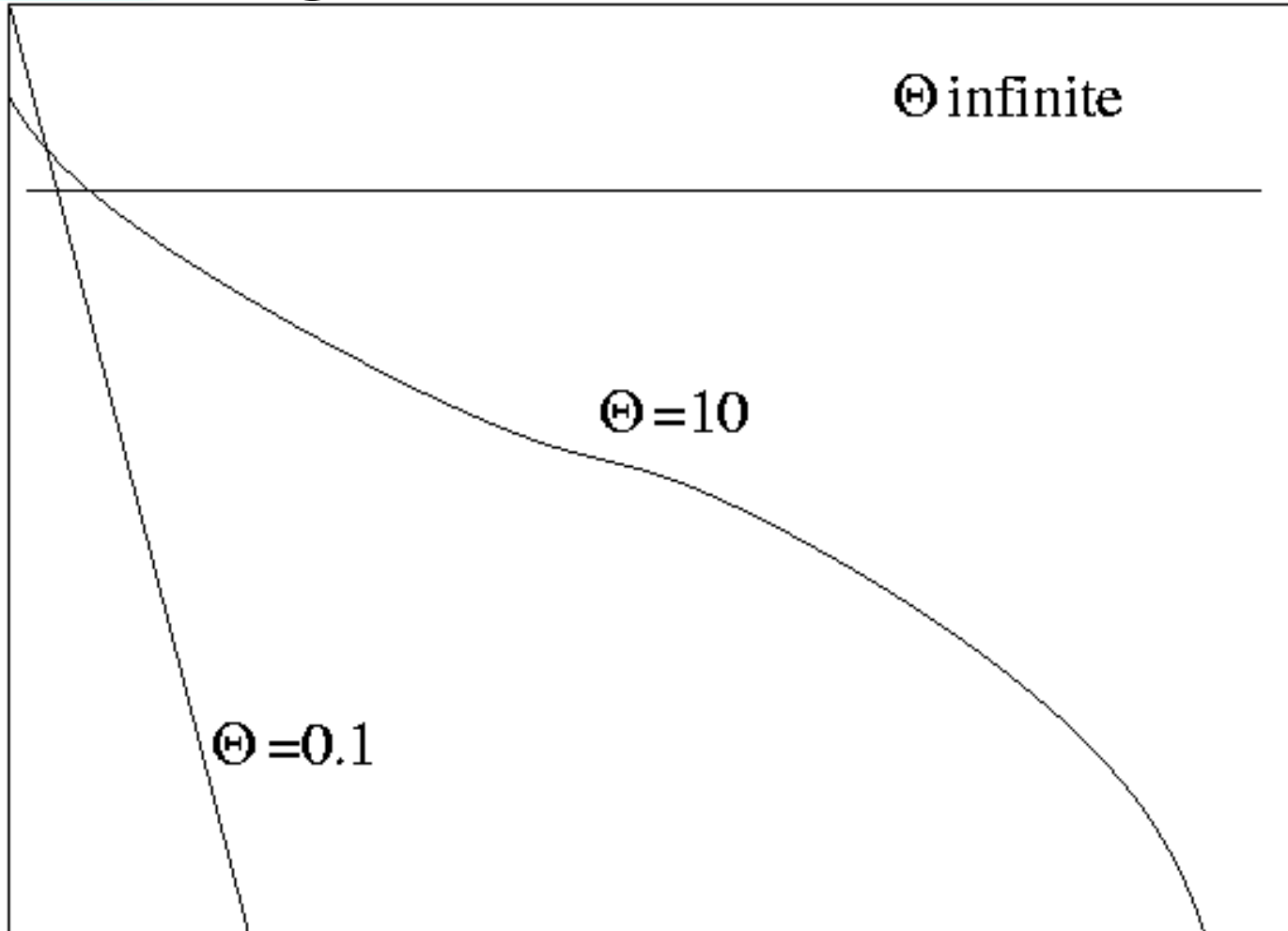


Same as a mutation!

Ecological Model Results



Number of individuals
Log of Equilibrium Species Abundance



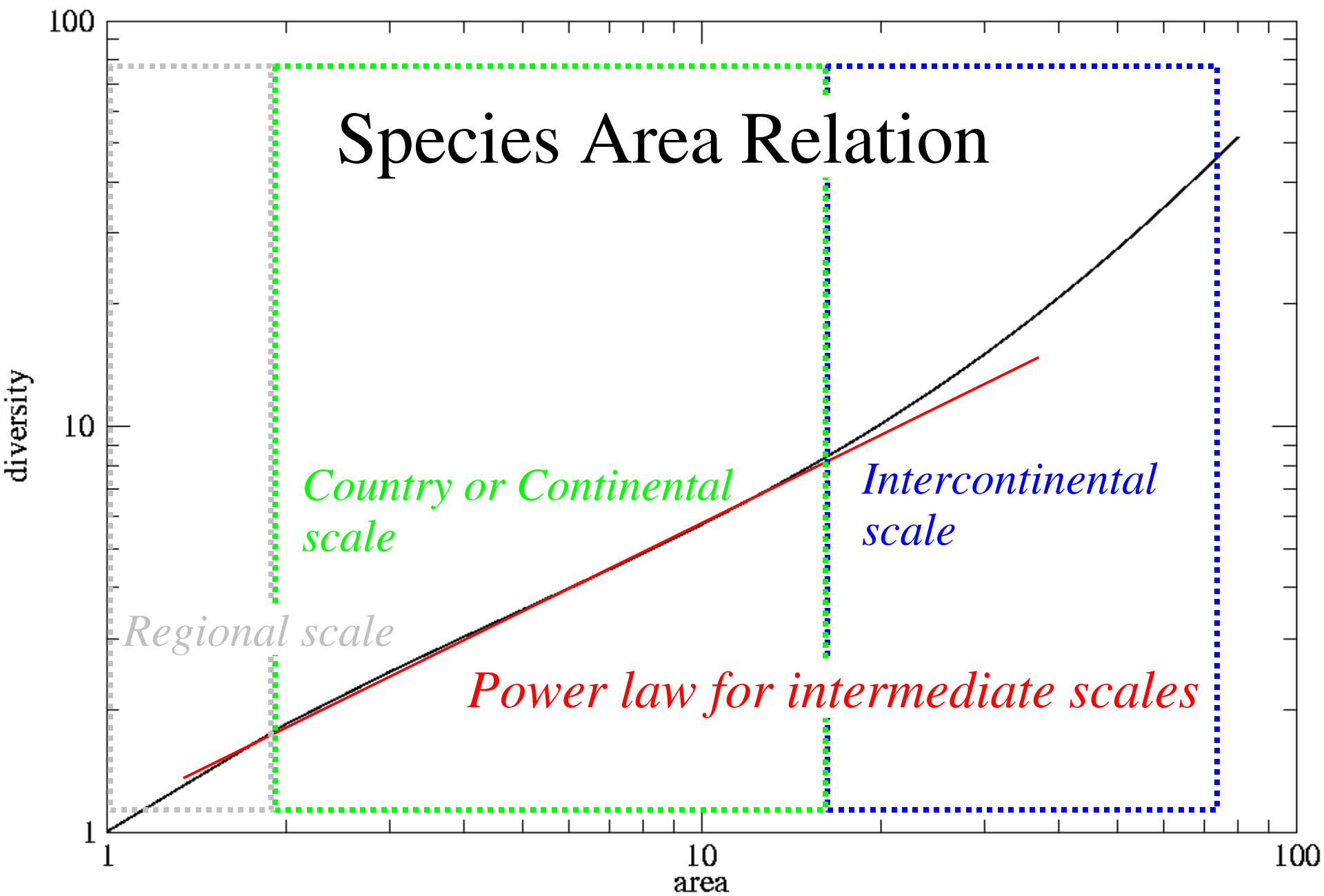
Species Rank in Abundance

*Species
number*

$\Theta = (\text{population size})(\text{probability of a new species})$ *ordered by size*

neutral model after 500 gens

correctly curved SAR



Species Area Relation

Country or Continental scale

Intercontinental scale

Regional scale

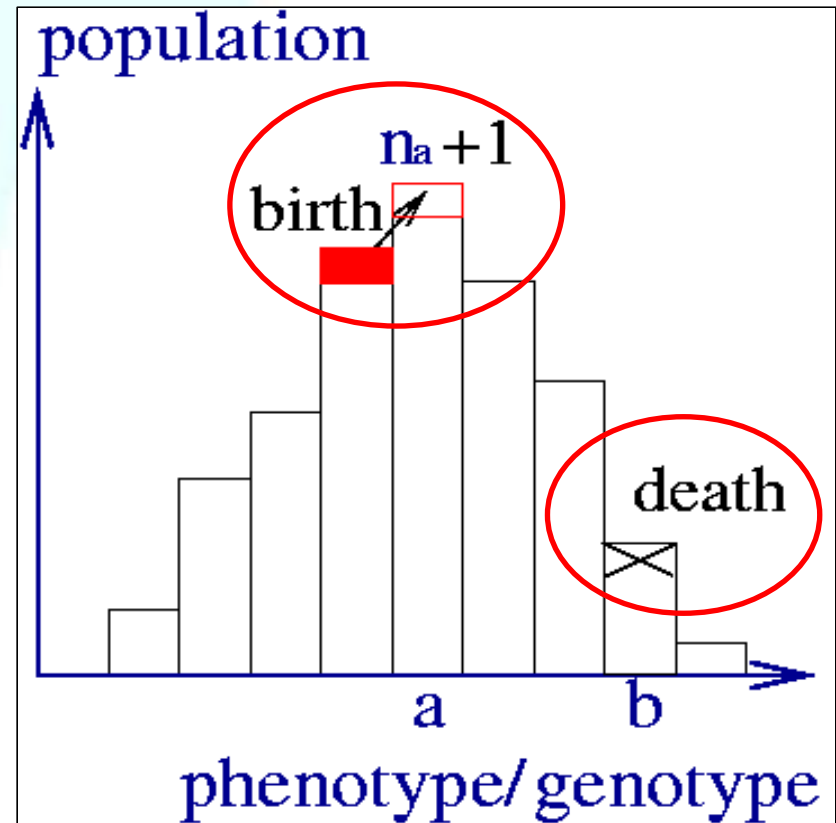
Power law for intermediate scales

An evolution model

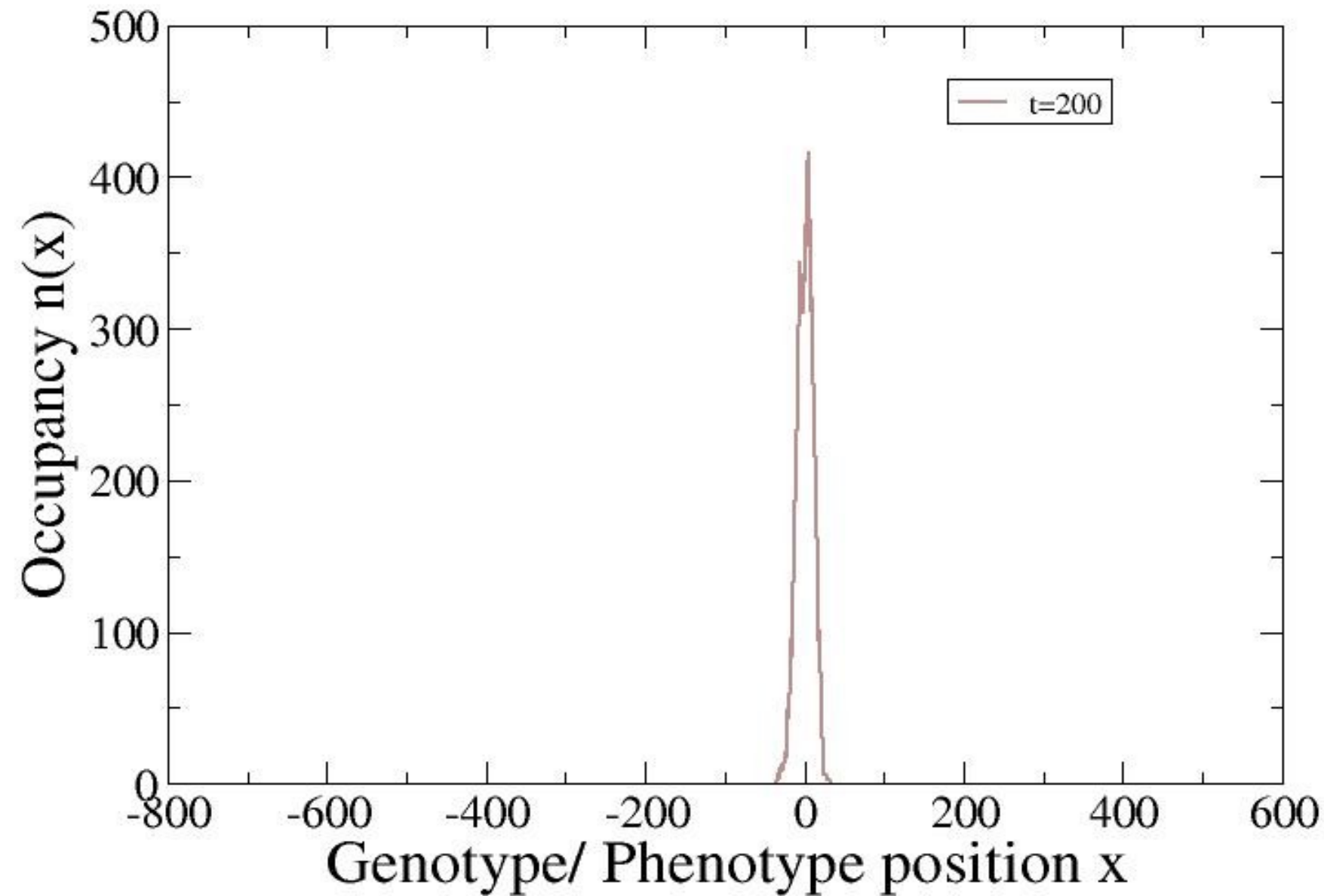
- Consider N individuals each labeled by phenotype position:

TIMESTEP:

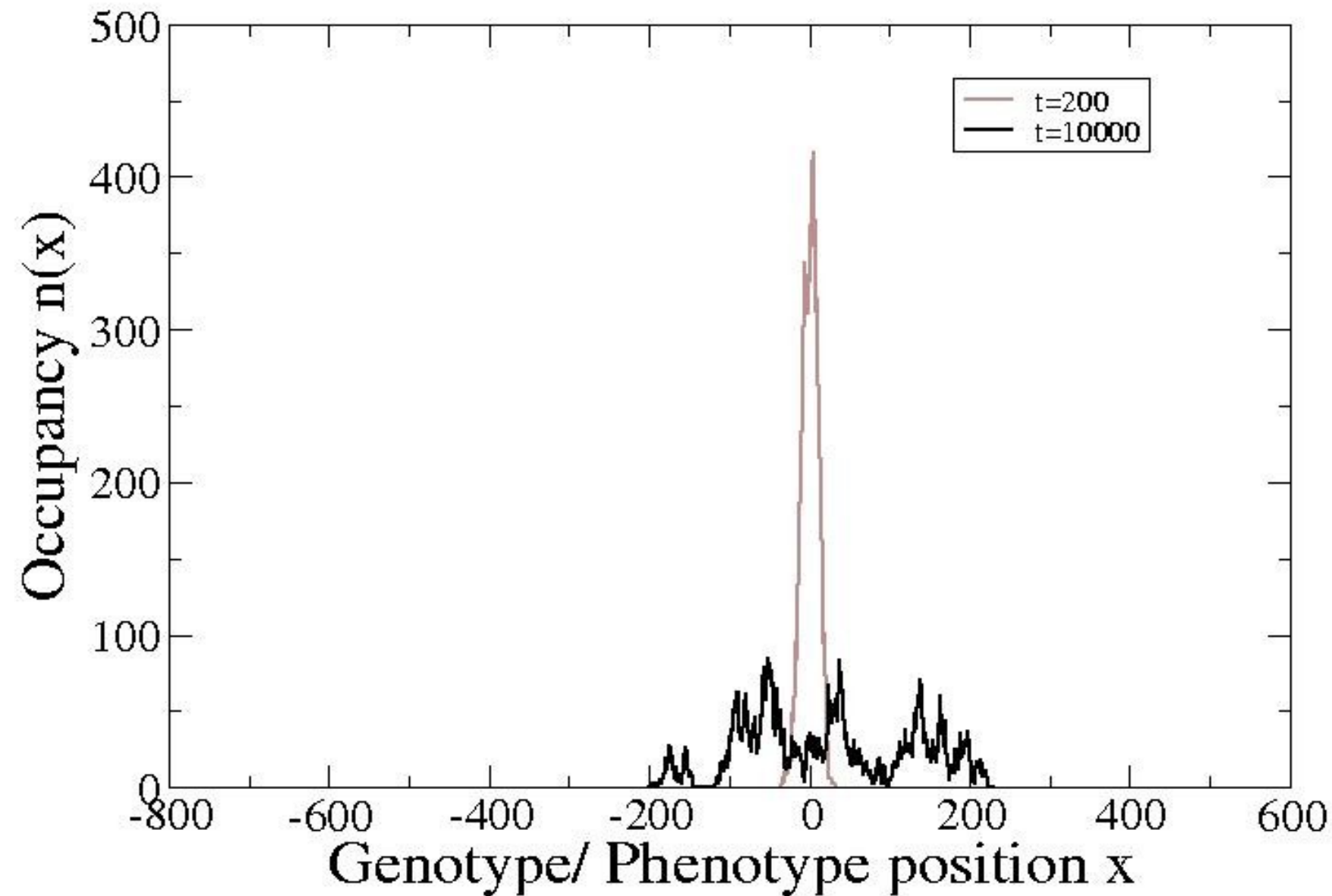
- Pick an individual (from N) and mark it to die.
- Pick an individual (from N) and copy it. With probability p_m **Mutate to a similar type.**
- Kill the marked individual.



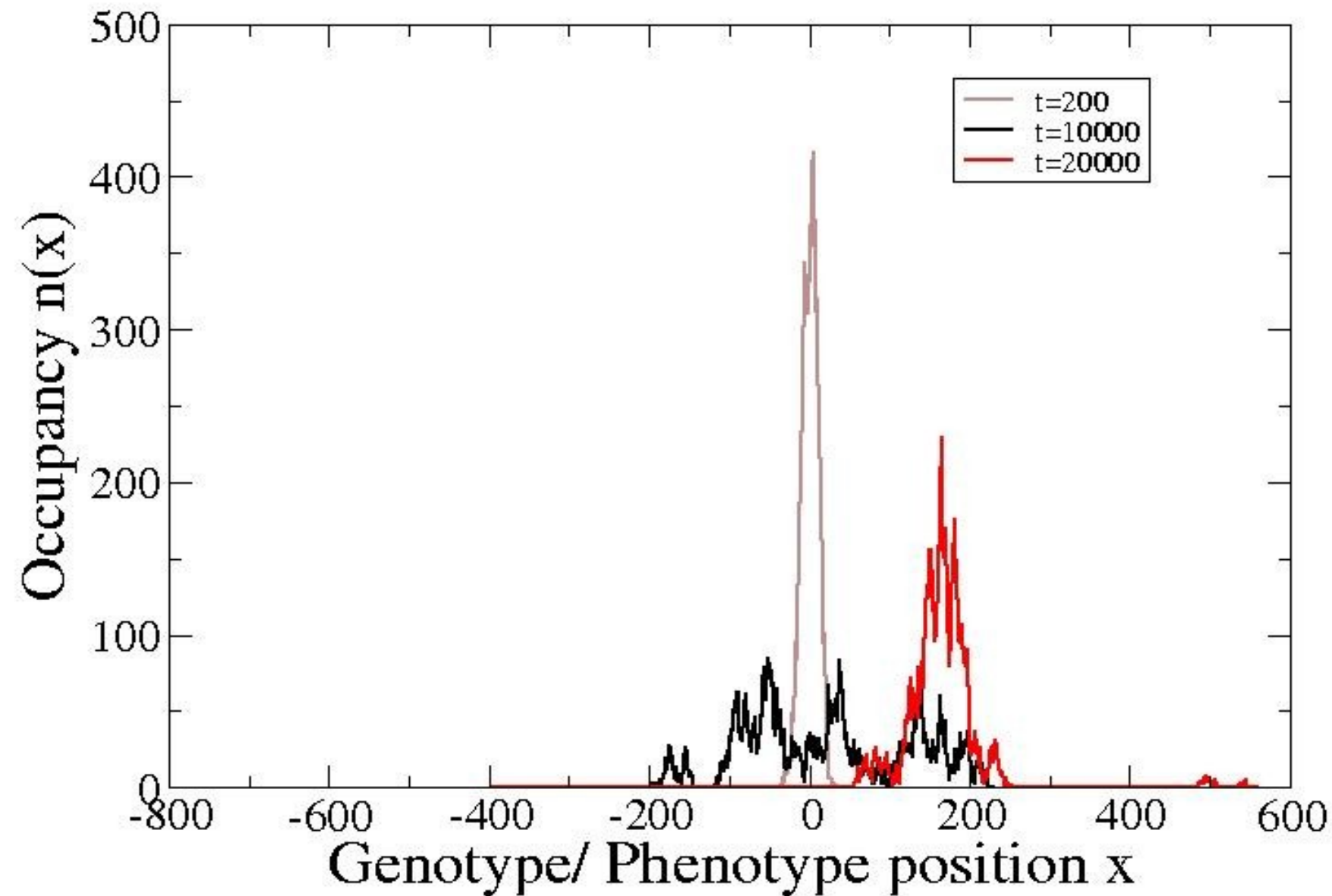
Evolution of 10000 particles



Evolution of 10000 particles

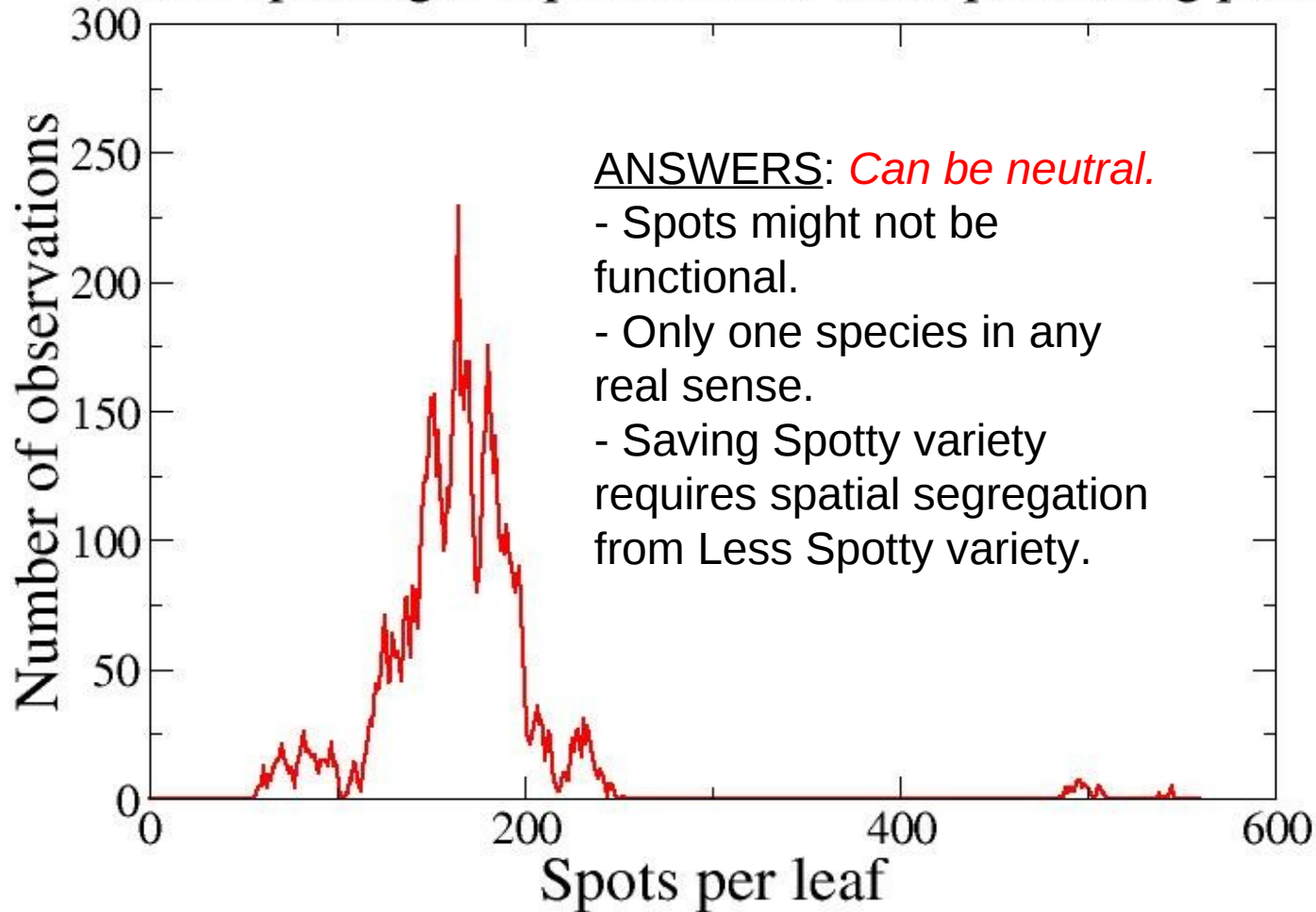


Evolution of 10000 particles

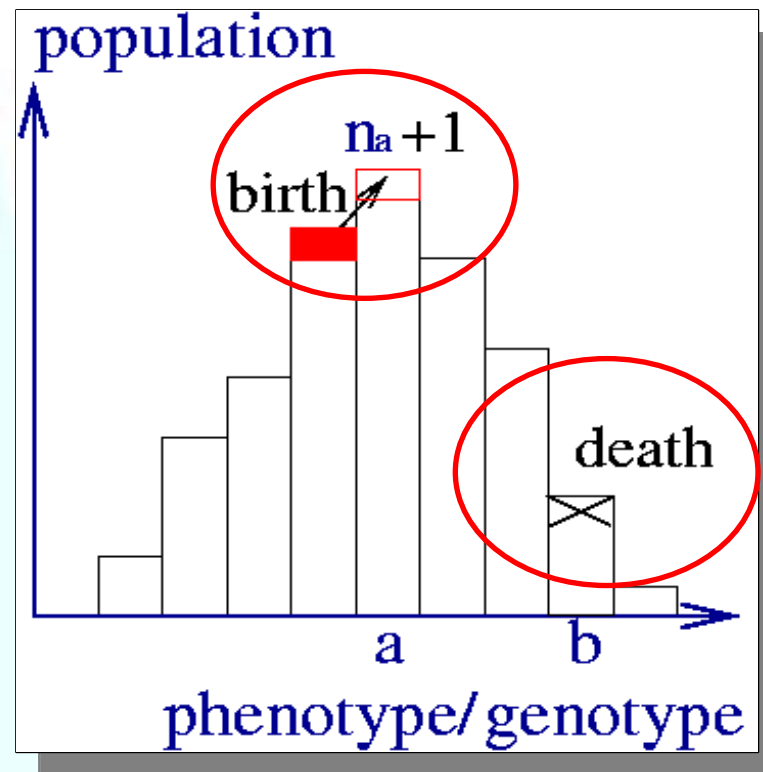


Test Problem

Observations of number of spots per leaf of *Imaginarium Forma*
(Made up thought experiment for a self pollinating plant)



Solution: first try (1)



$$p_b(a) = \frac{n_a}{N} (1 - p_m) + \frac{p_m}{2} \frac{n_{a+1} + n_{a-1}}{N}$$

$$p_k(a) = \frac{n_a}{N}$$

Solution: first try (2)

- Mean field argument:

$$\langle n_a(t+1) - n_a(t) \rangle = p_b(a) - p_k(a)$$

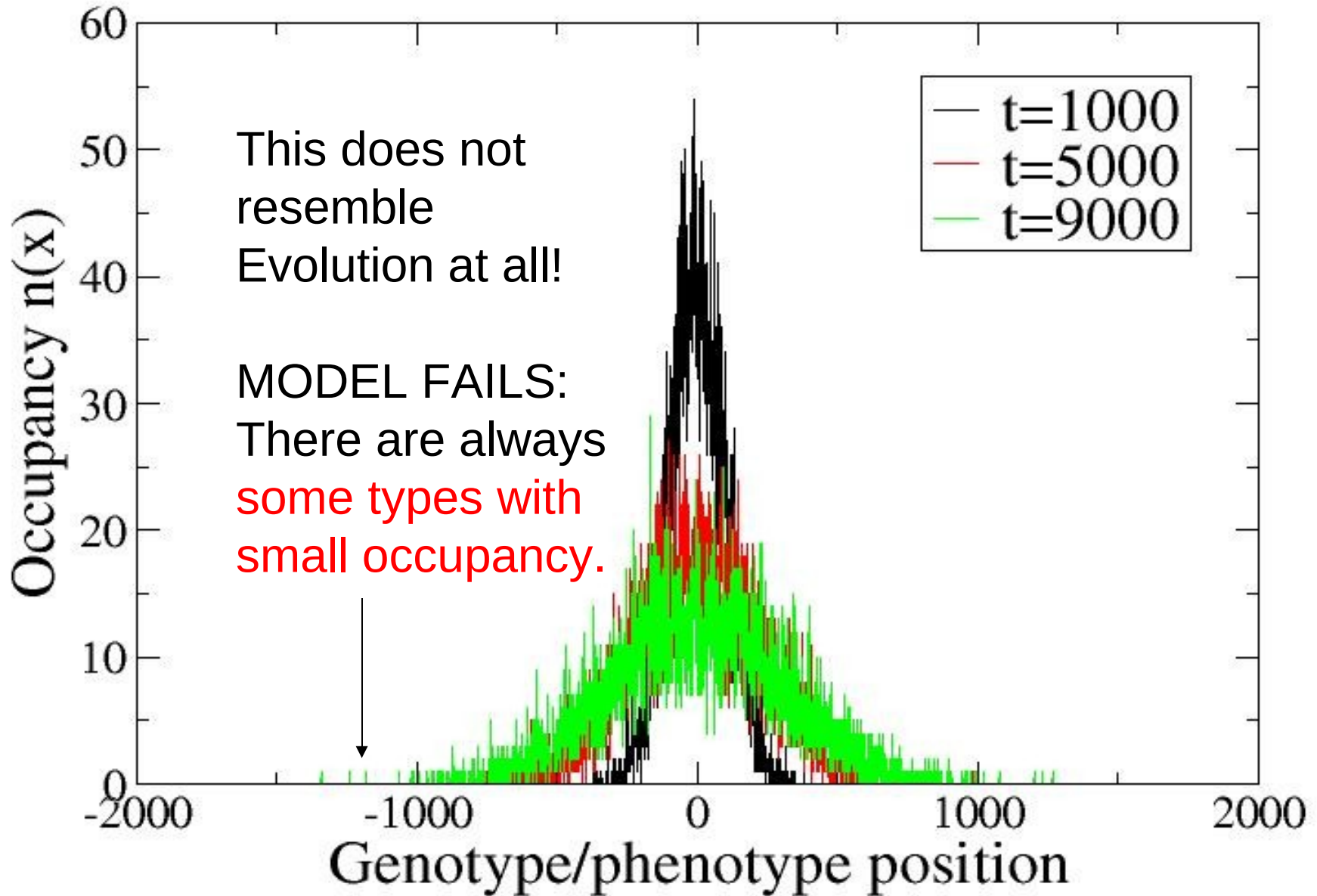
– Expect this to be valid when N large?

- This becomes:

$$\langle n_a(t+1) - n_a(t) \rangle = \frac{p_m}{2N} [n_{a+1}(t) + n_{a-1}(t) - 2n_a(t)]$$

- This is just the diffusion of N particles!

Diffusion of 10000 particles



Solution

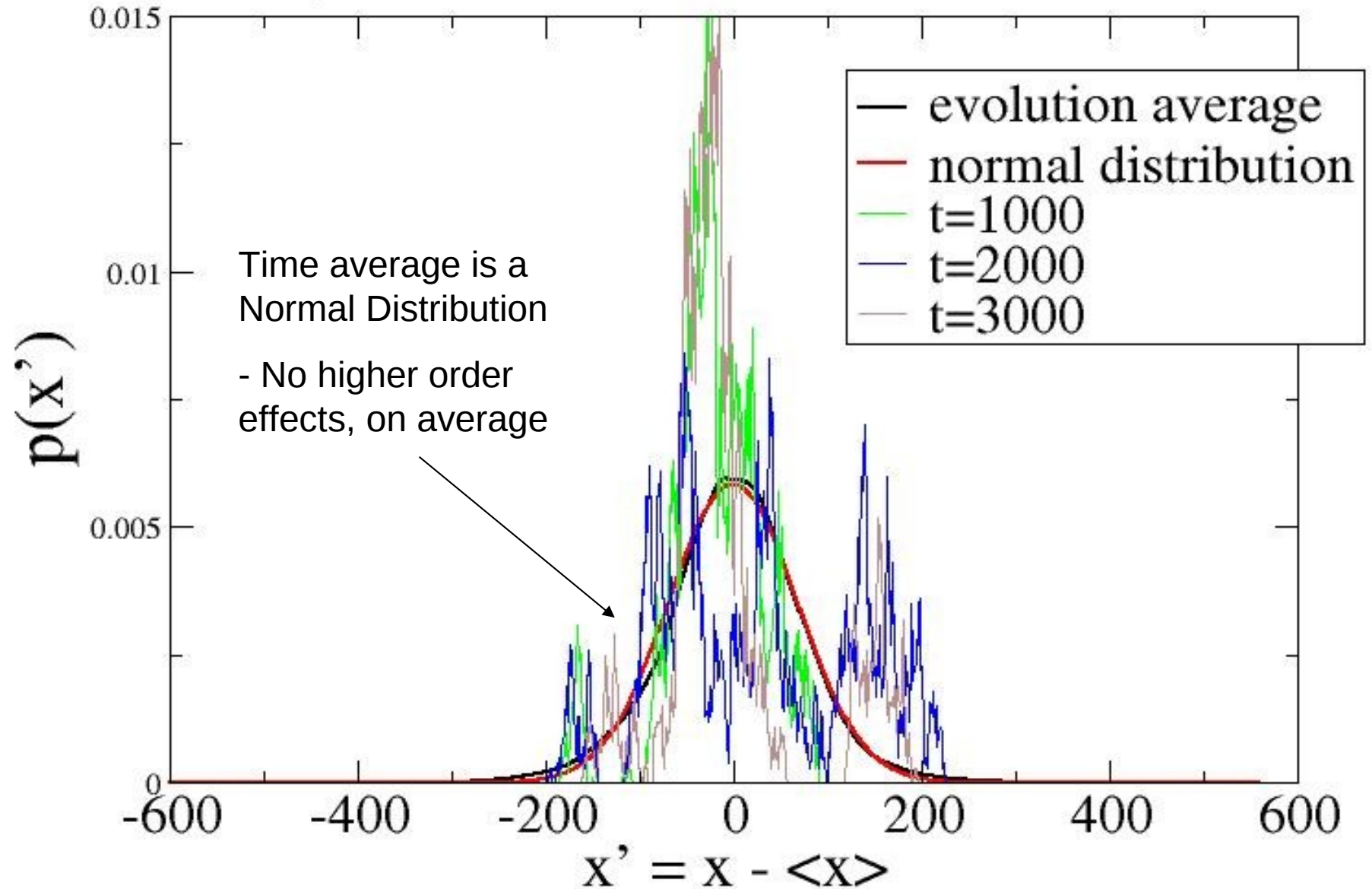
- Simplify the model – consider only first two moments of the distribution
- Peak is a Gaussian distribution of area 1 with dynamic mean μ and width w .

- Select death location x
- Select birth location y , mutated by 1 with probability P_m
- Remove $1/N$ from death location and place at birth location
- Update μ and w

Solution method

- Write down equations for the change in the mean and the variance of the peak position μ and the width w
- Take continuous limit to obtain Stochastic Differential Equations
- Solve!
- This works because the distribution is normal only when *averaged over time*, measured *relative* to the *current mean position*

Time average distribution of evolution process around the mean compared with a normal distribution (N=10000)



Solving for the width

$$d(w^2) = \left(p^* - \frac{2w^2}{N} \right) dT + \frac{2w^2}{\sqrt{N}} dW$$

Mutation distance
Generation time

dW is Random, mean 0

Change in variance (in a timestep)

Deterministic part

+ Noise part

Solution at steady state:

$$p(w)dw = \frac{(Np_m)^2}{2w^5} e^{-\frac{Np_m}{2w^2}} dw$$

Power-law decay at large w

Neutral Clustering results

- Mean width: $\langle w \rangle = \sqrt{\frac{N p_m \pi}{8}}$

Fluctuations in w also $\sim N^{0.5}$

- Position: $\langle x \rangle_{\text{RMS}} = \sqrt{T(p_m + w^2)} \sqrt{\frac{p_m T}{2}}$

*With time in generations...
 $\langle x \rangle_{\text{RMS}}$ is independent of N !*

- Compare with diffusion:

$$\langle x \rangle_{\text{RMS}} = \sqrt{\frac{p_m T}{N}} \qquad \langle w \rangle_{\text{RMS}} = \sqrt{\frac{p_m T}{N}}$$

- Diffusion “does nothing” in infinite populations... evolution does “more”!

Neutral evolution results

- Selection produces a stable peak, neutrality produces an *unstable* peak.
- Characteristic peak width, but large fluctuations (*multiple clusters*).
- Peak centre diffuses *independently* of population size!
- Provides null model for expected phenotype distributions, *non-trivial* distributions are probable.



Reference

Lawson and Jensen:

***“Neutral Evolution as Diffusion in phenotype space:
reproduction with mutation but without selection”***

Physics Review Letters, March 07 (98, 098102)

www.arxiv.org/abs/q-bio/0609009

Thank you for your attention!

Diversity in models

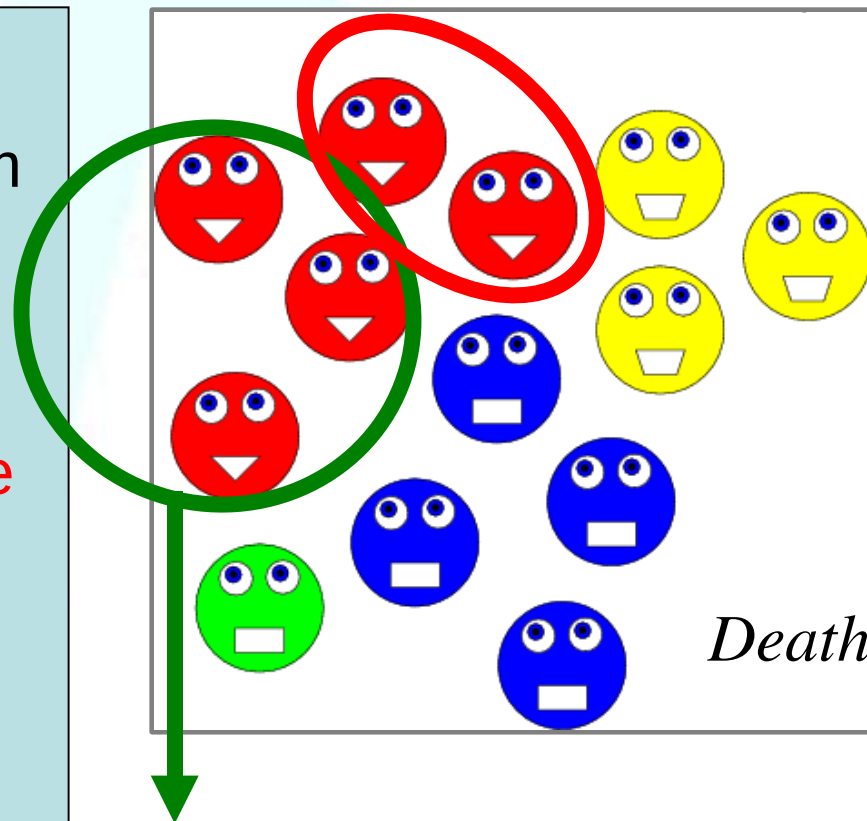
- We expect a “species”:
 - To be “different enough” from other species.
 - To be constant in time. An individual of a species today is comparable with an individual of that species in the past.
- A model discussing species diversity must accurately represent:
 - Species Abundance Distribution
 - Species Lifetime Distribution
 - Species Area Relation
 - etc....

Updated ecological model

- Consider N individuals each labeled by species:

TIMESTEP:

- Pick an individual (from N) and mark it to die.
- Pick an individual (from N) and with probability p_a **a proportion speciate allopatrically**.
- Kill the marked individual.



Mutate proportion of population allopatrically

How long do species survive?



Cutoff time determined by probability of mutation

