

# Speciation & hybridization

Matías Gómez

# Outline

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What are species ?

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Speciation

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Hybridization

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Geographic modes of speciation

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Barriers to gene flow

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Genomics of speciation

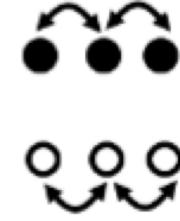
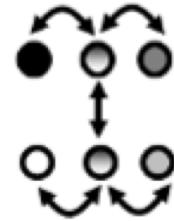
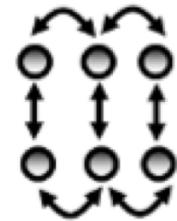
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Phylogenetics

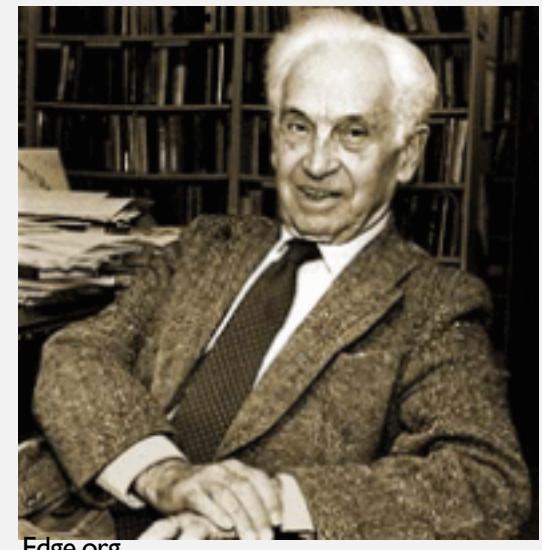
# What are species ?

- **Biological species concept (BSC):** Species are groups of actually or potentially interbreeding natural populations that are reproductively isolated from other such groups (Mayr, 1942).

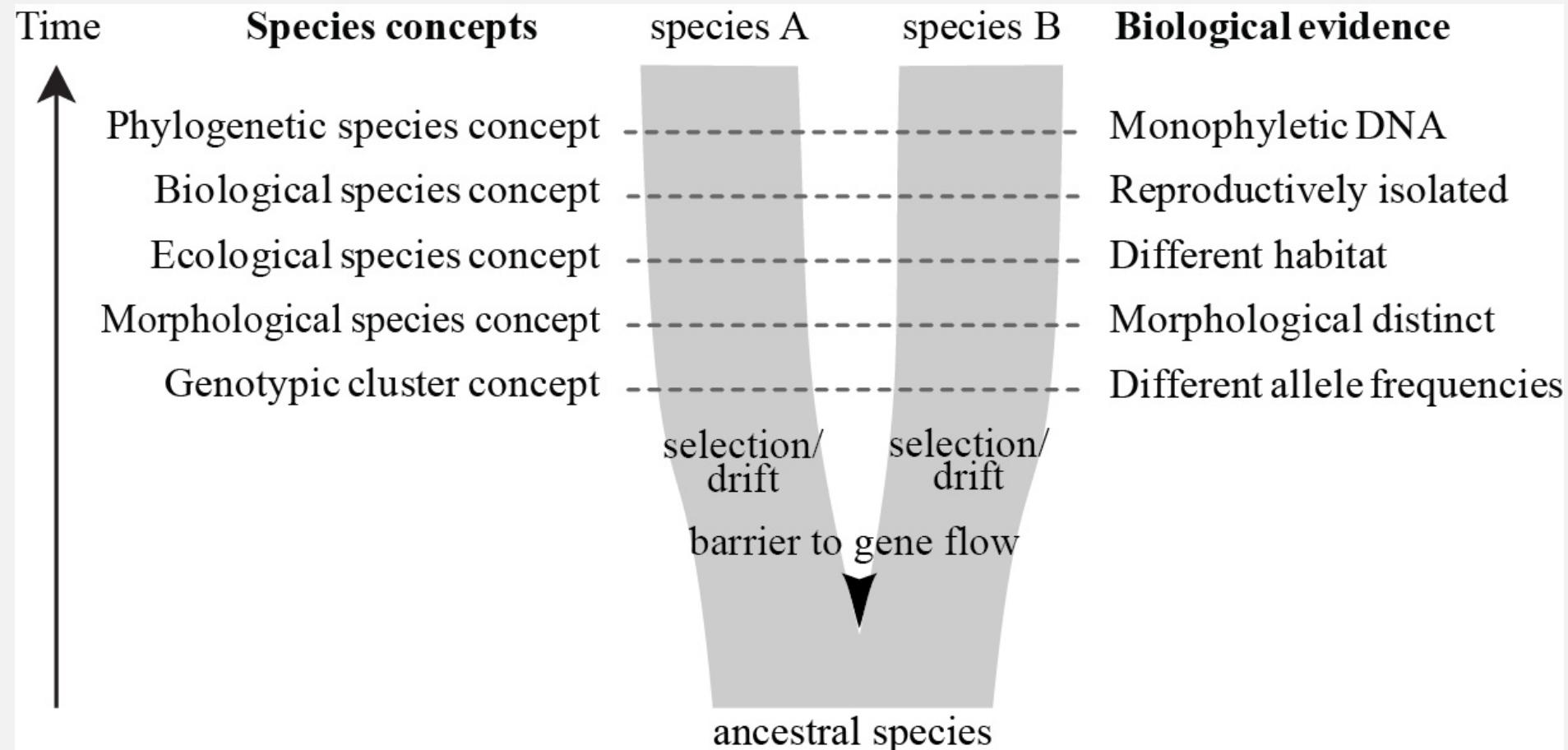
reproductive  
isolation



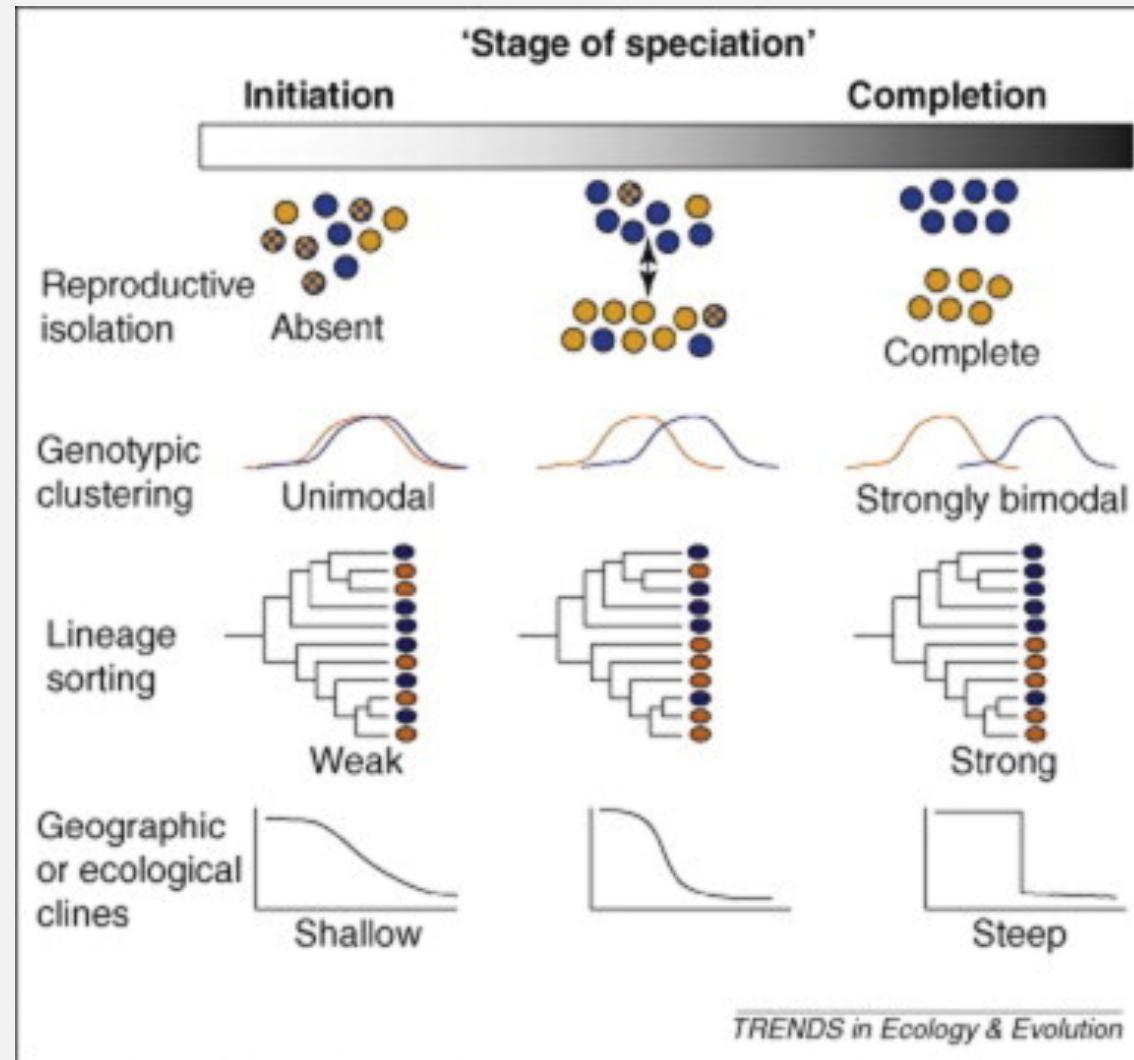
Nosil et al. (2009), *Trends in Ecology & Evolution*



# Other species concepts



# Speciation continuum



# Biological species concept (BSC):

## PROS

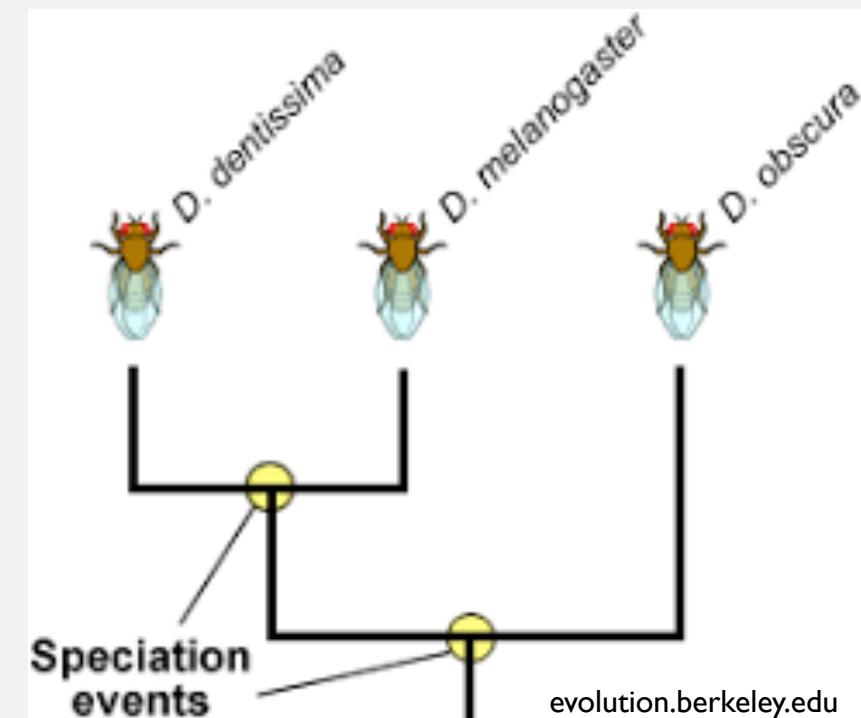
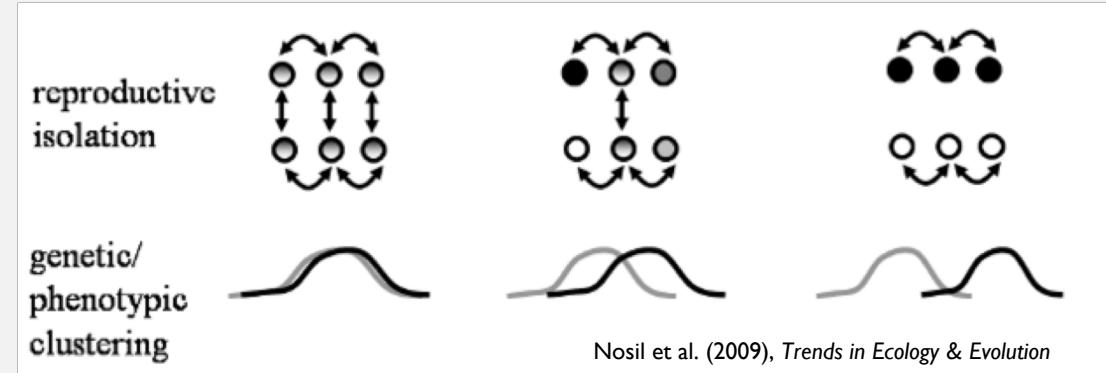
- Interbreeding explains why the members of a species resemble one another
- Reproductive isolation is caused by barriers to gene flow
- Gene pool = independent unit of evolution
- Organisms do not evolve but species do
- Species as a “harmonious” gene pool

## CONS

- Asexual reproduction
- Fossil species
- Hybrids

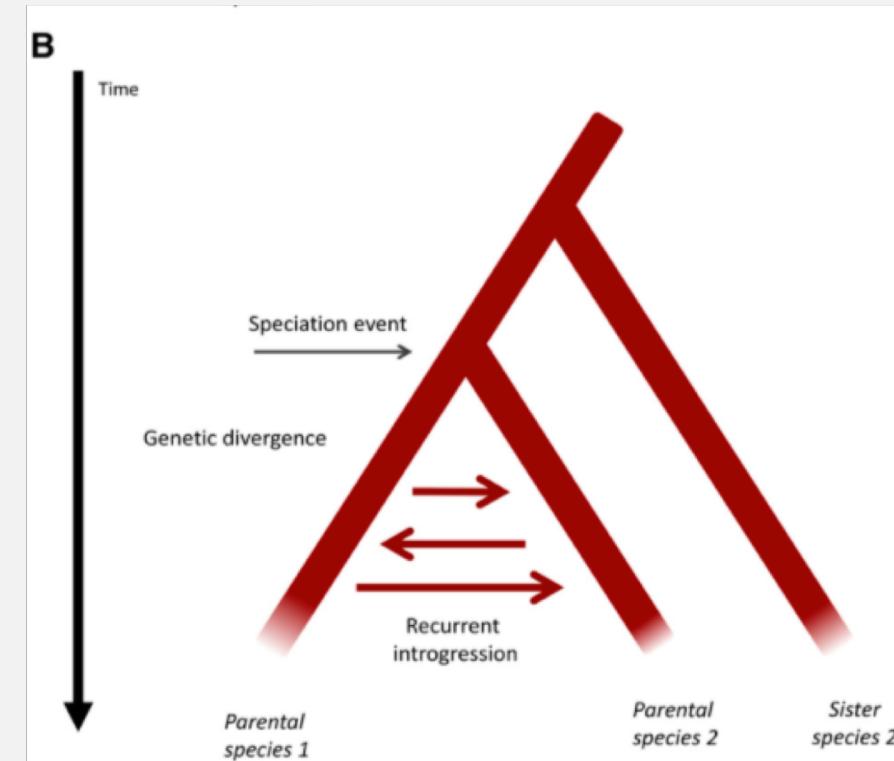
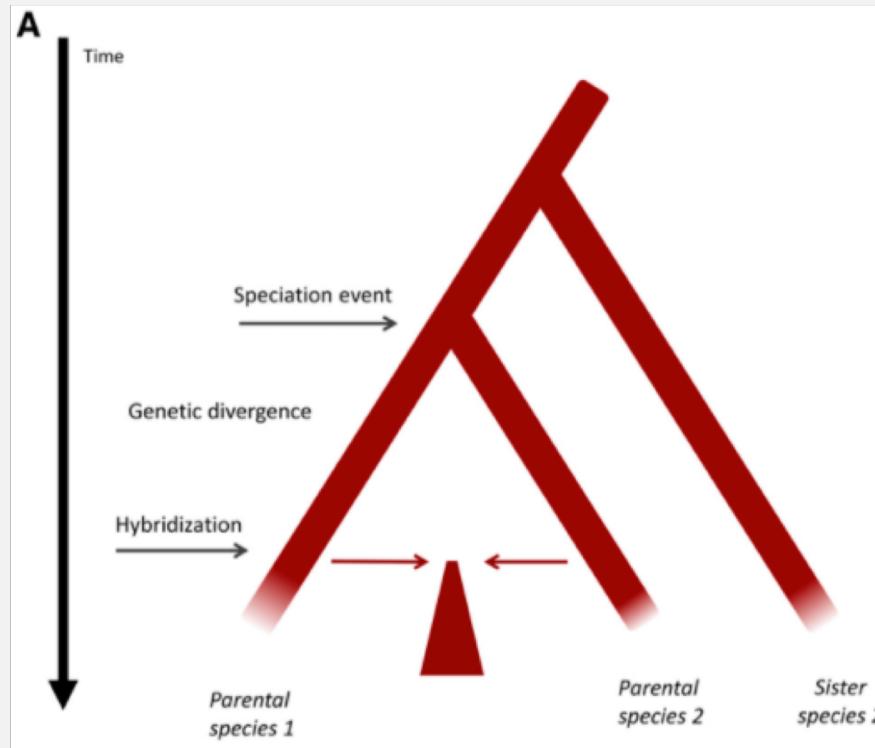
# Speciation

- The origin of two or more species from a common ancestral one
- Barriers to gene flow
- Reproductive isolation (RI)
- Discrete and interbreeding genetic pools

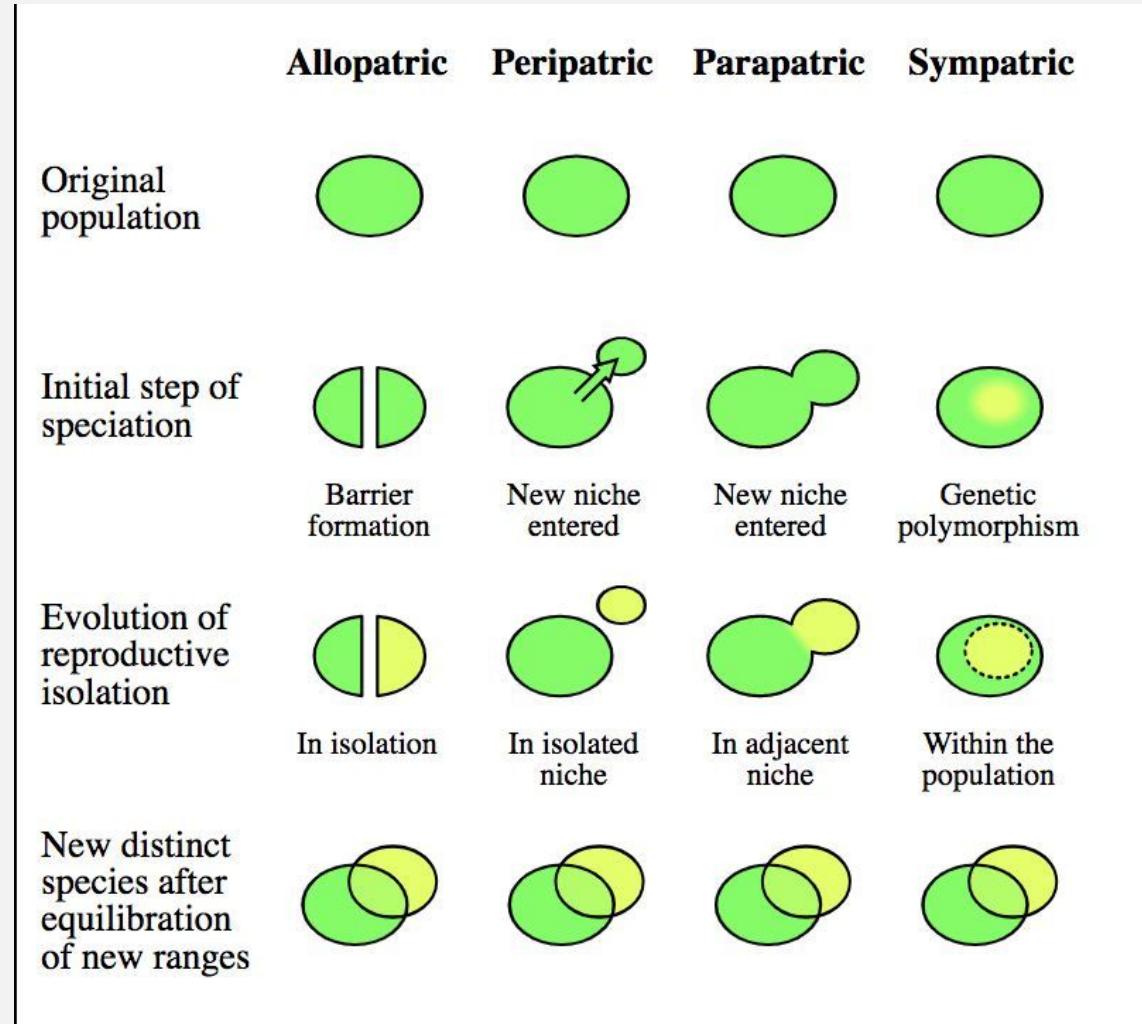


# Hybridization

- Interbreeding between species or divergent populations
- Introgression: incomplete postmating isolation

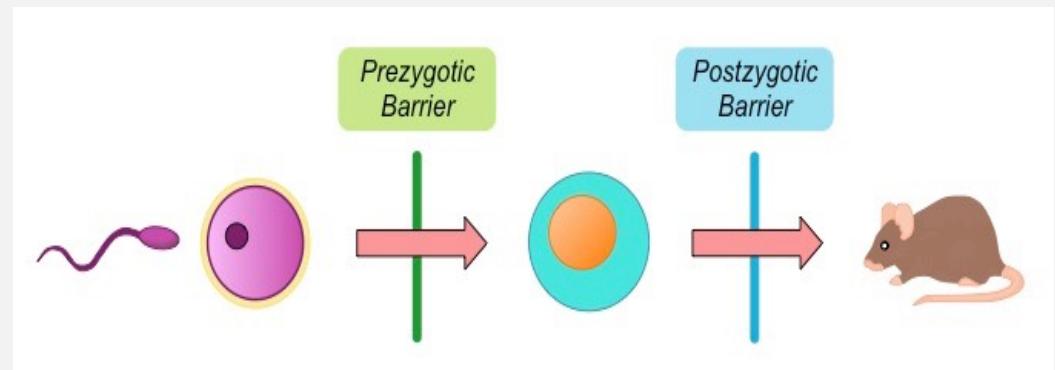


# Geographical modes speciation



# Barriers to gene flow

1. Premating/prezygotic: Features that impede transfer of gametes to members of other species
2. Postmating and prezygotic: Mating or gamete transfer occurs, but zygotes are not formed
3. Postzygotic: zygotes are formed but have reduced fitness



<http://ib.bioninja.com.au>

## I.Premating/prezygotic

### A) Ecological isolation:

➤ Potential mates in the same area do not meet

- Habitat: spatially segregated

Ladybird beetles in Japan exclusively mate on its own host plant

*Epilachna niponica* and *E. yasutomii*



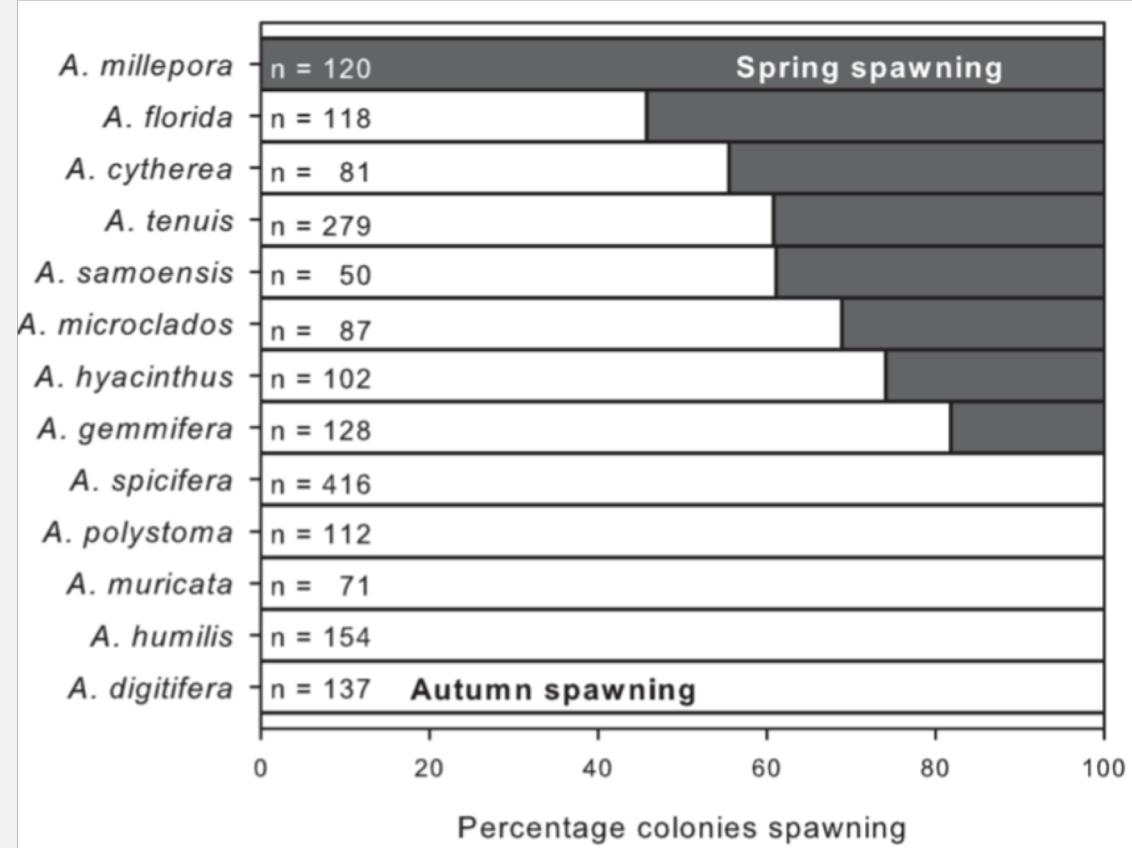
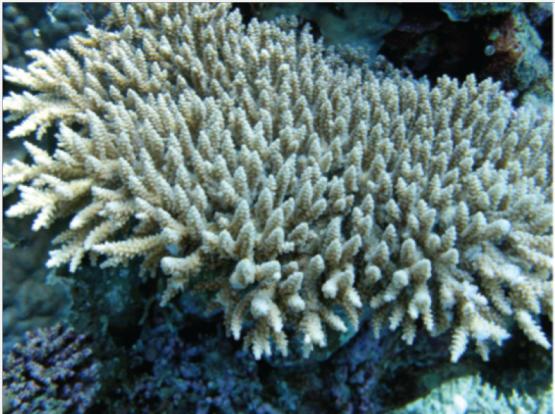
Kakatura (1997), Zoological Science



# I.Premating/prezygotic

## Ecological isolation:

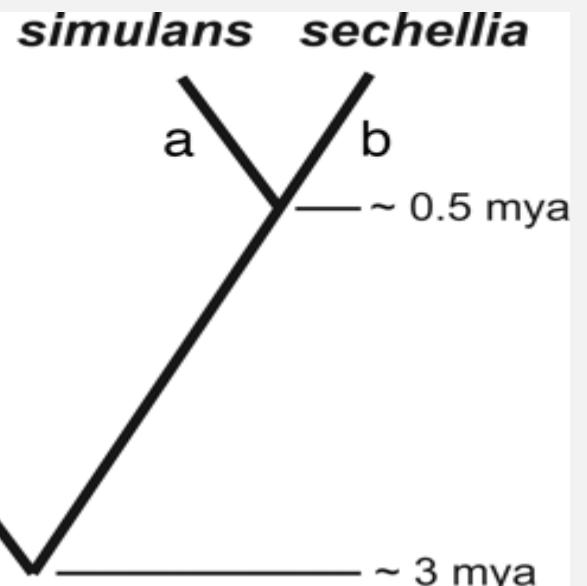
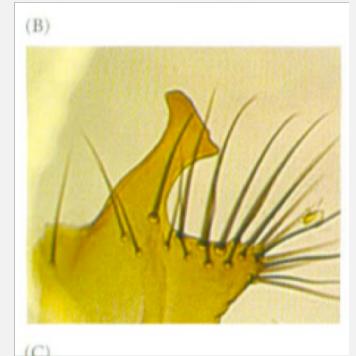
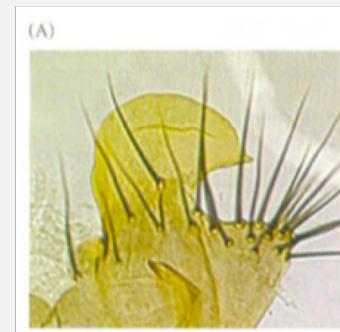
- Temporal: populations breed at different seasons or times of day
- *Acropora spp.* Indopacific



Gilmoure et al.(2016), PLOS ONE

## I.Premating/prezygotic

**B) Copulatory isolation:**  
behavior/structure prevents copulation  
“key-lock” genitalia in *Drosophila simulans* and *D. sechellia*.



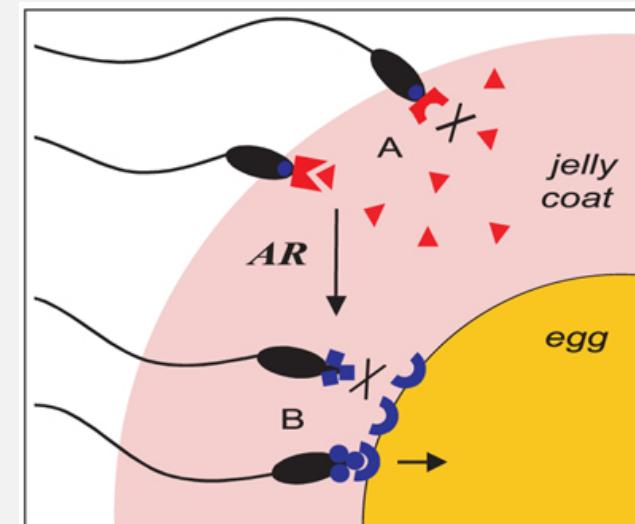
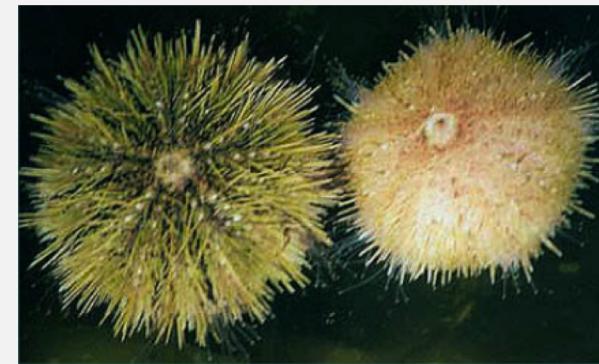
McBride (2007), PNAS

## I.Premating/prezygotic

### C) Gametic isolation:

Released/transferred gametes cannot effect fertilization

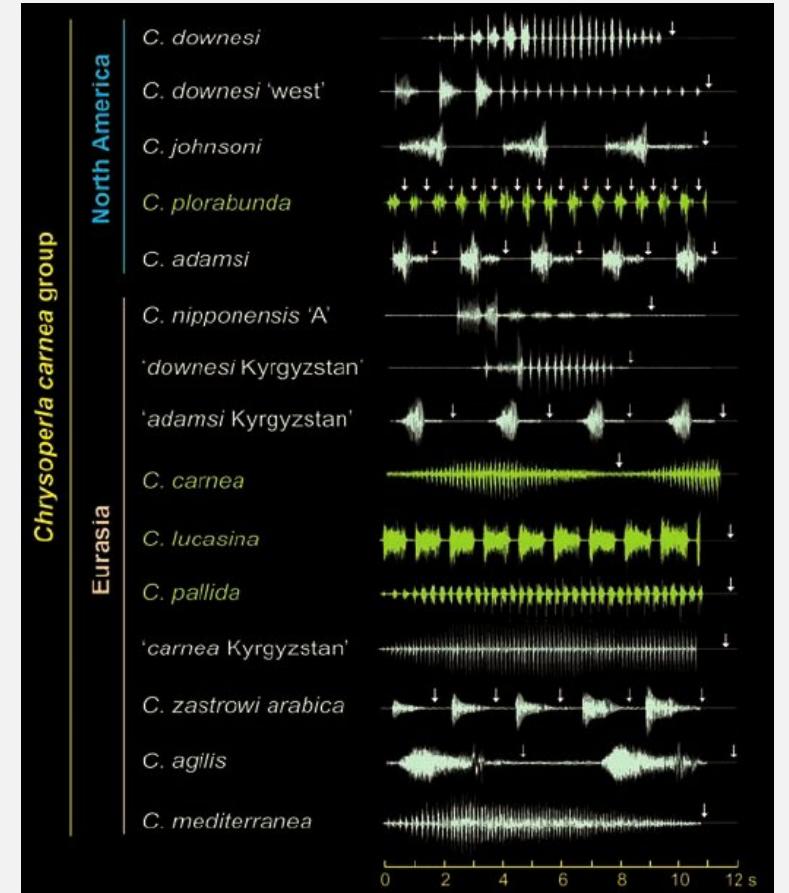
- *Strongylocentrotus droebachiensis* and *S. pallidus*: sperm-egg recognition proteins (bindin and bindin receptor)



## I.Premating/prezygotic

- Potential mates meet but do not mate
- D) Behavioral isolation:** the lack of cross attraction between members of different species.

Lacewing  
(*Chrysoperla spp.*)

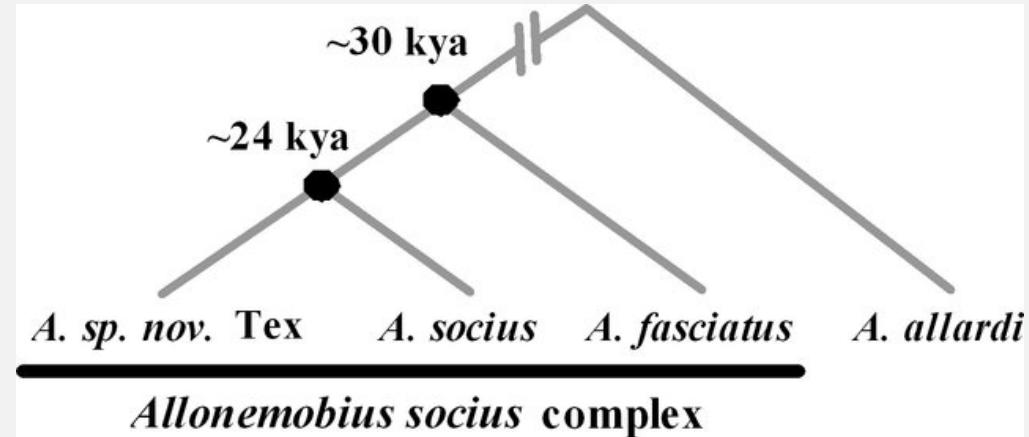


## 2. Postmating and prezygotic

**Conspecific sperm precedence:** females are exposed to sperm from males of multiple species, but preferentially use sperm of a conspecific.



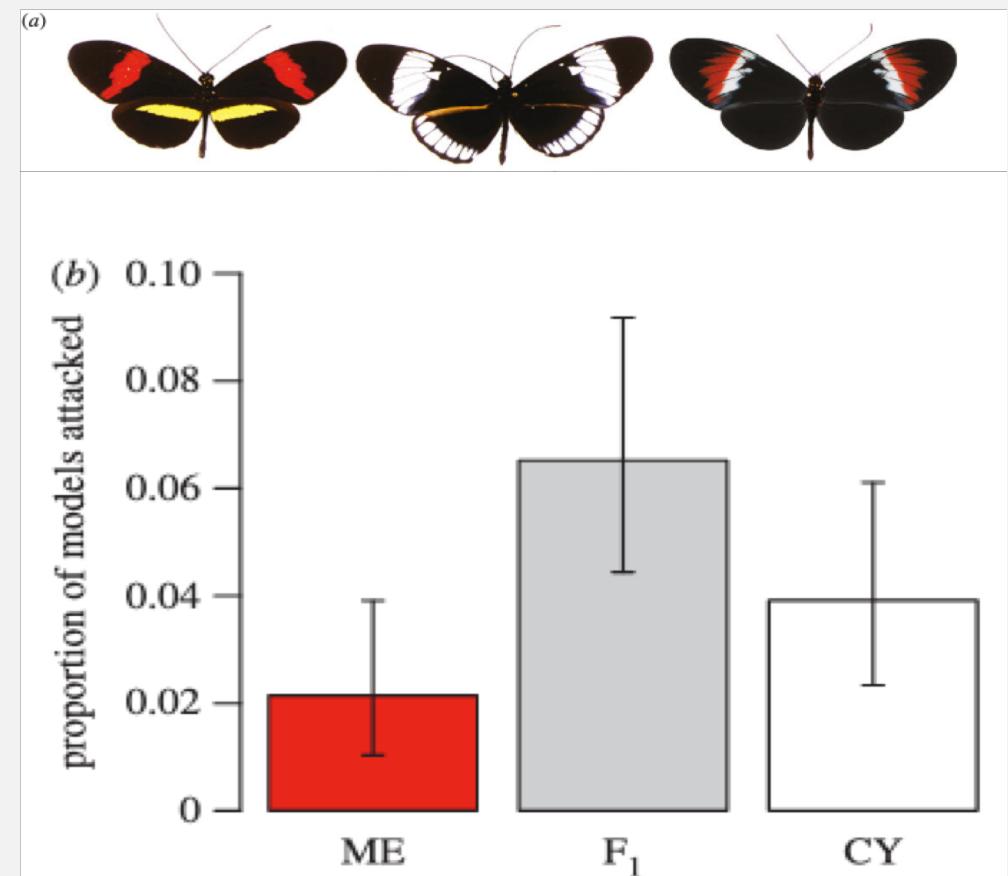
Copyright © 2006 Bill Claff



### 3. Postzygotic

**A) Extrinsic:** environment-dependent or lack of appropriate niche

*Heliconius melpomene* × *H. cydno* = hybrid



### 3. Postzygotic

#### B) Intrinsic: genomic conflict

- Hybrid inviability

*D. melanogaster* × *D. simulans* = inviable zygotes



*melanogaster*

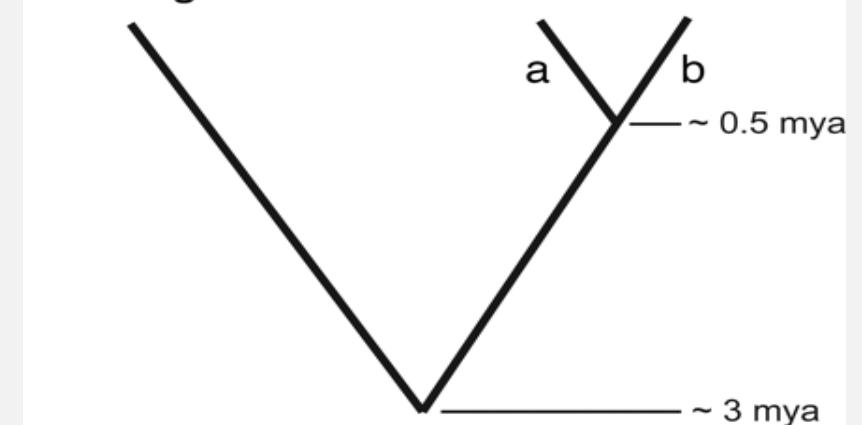


*simulans*

*sechellia*

- Hybrid sterility

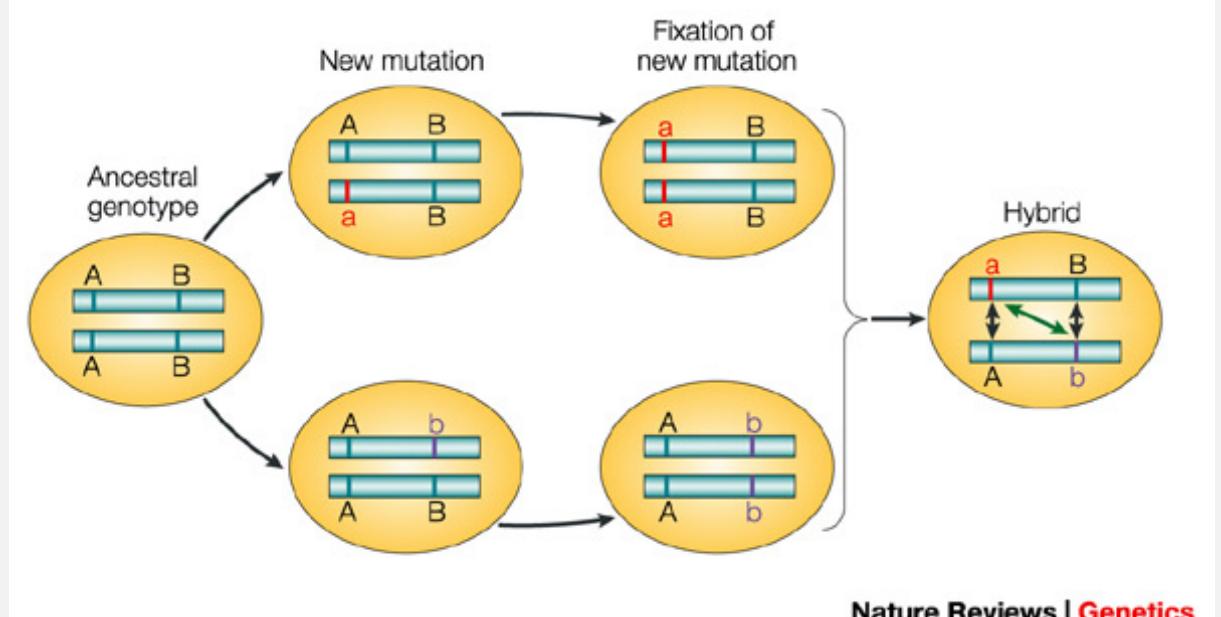
Mule



McBride (2007), PNAS

### 3. Postzygotic

- Dobzhansky-Muller model of hybrid incompatibility

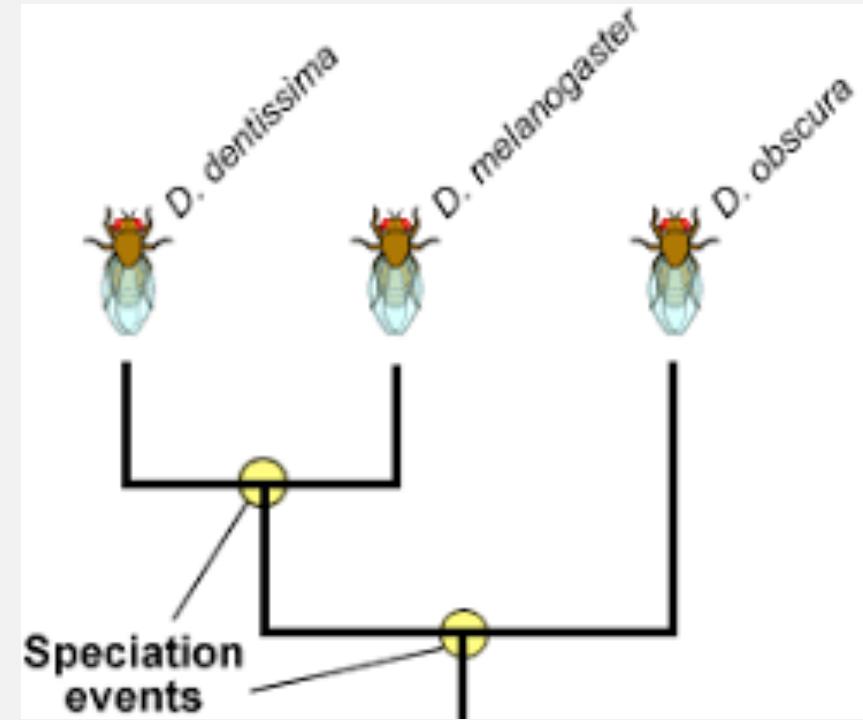


Nature Reviews | Genetics

Wu & Ting (2004)

# The genomics of speciation

- The speciation process leads to an increase in the genomic differences between divergent populations.
  - Main factors affecting speciation:
    - Genetic drift: neutral evolution
    - Migration: homogenizing effect
    - Selection: adaptive evolution



# Simulations

- **Drift, migration and selection**
  - A biallelic locus evolving in two populations
  - Equal population size and number of generations

Pop 1:  $f(A) = 1$   
 $f(a) = 0$

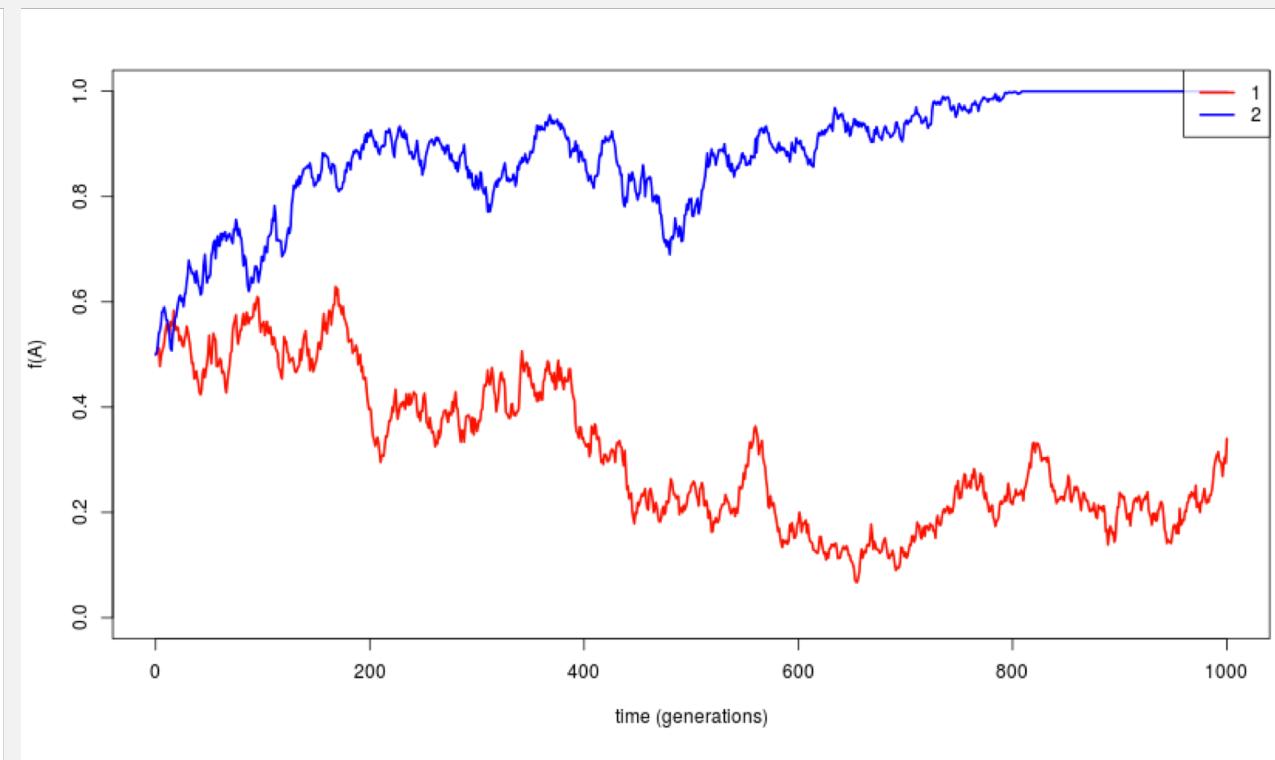
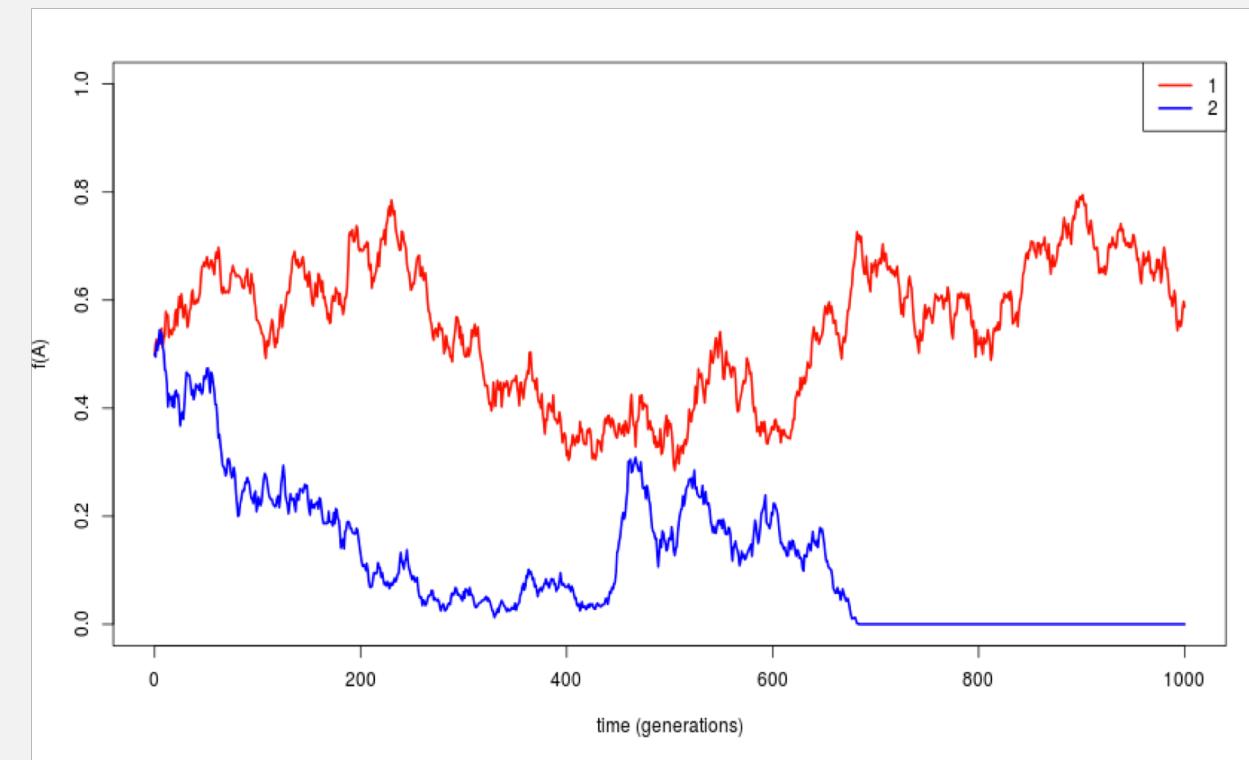
Pop 2:  $f(A) = 0$   
 $f(a) = 1$

# Genetic drift

No selection or migration

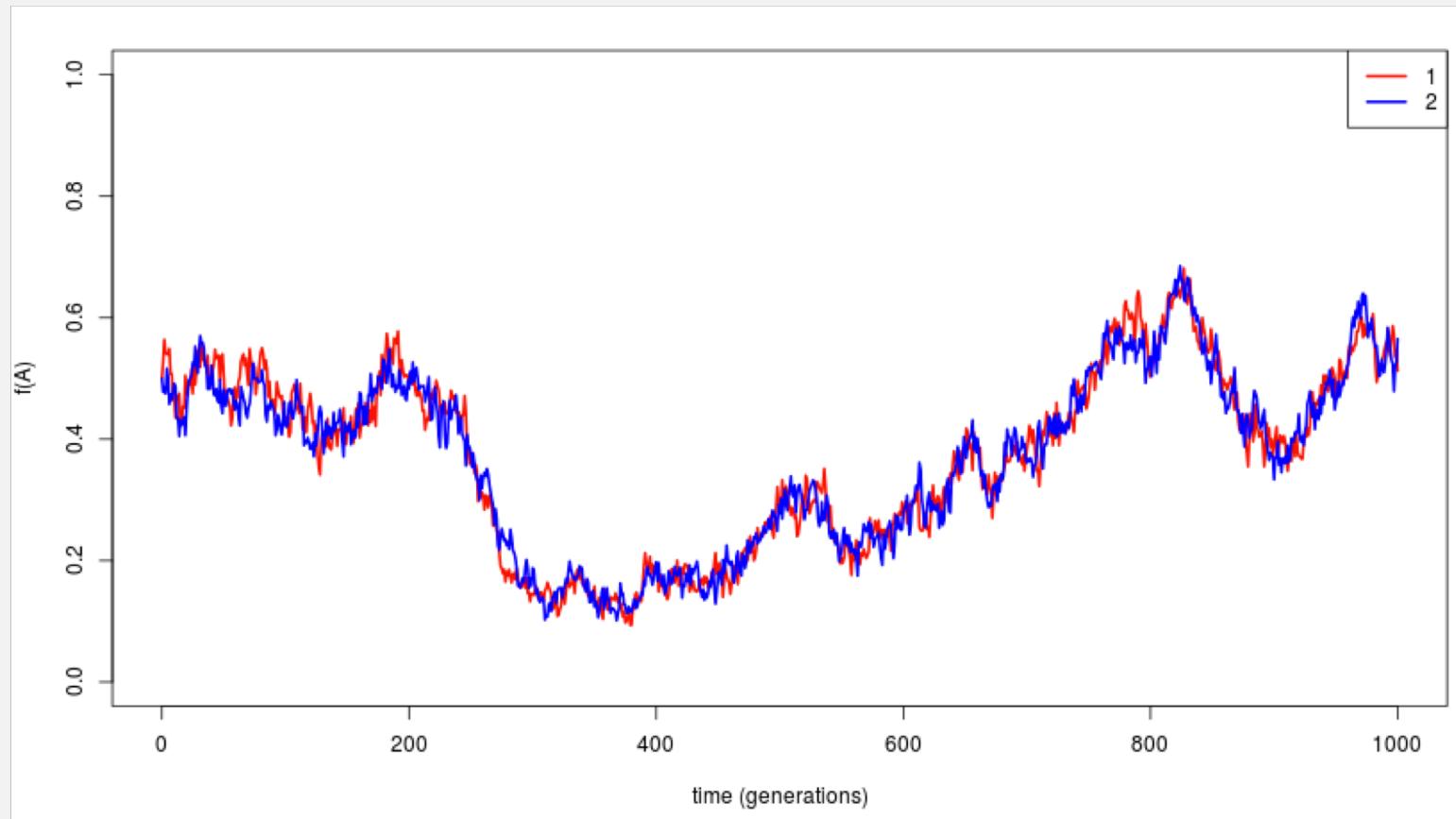
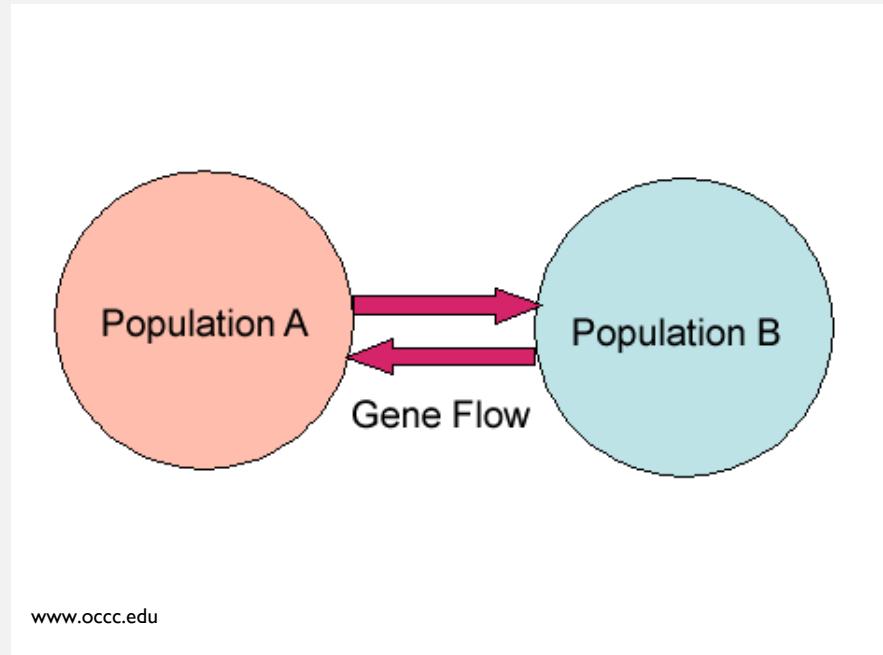
Pop 1:  $f(A) = 0.5$      $f(a) = 0.5$

Pop 2:  $f(A) = 0.5$      $f(a) = 0.5$



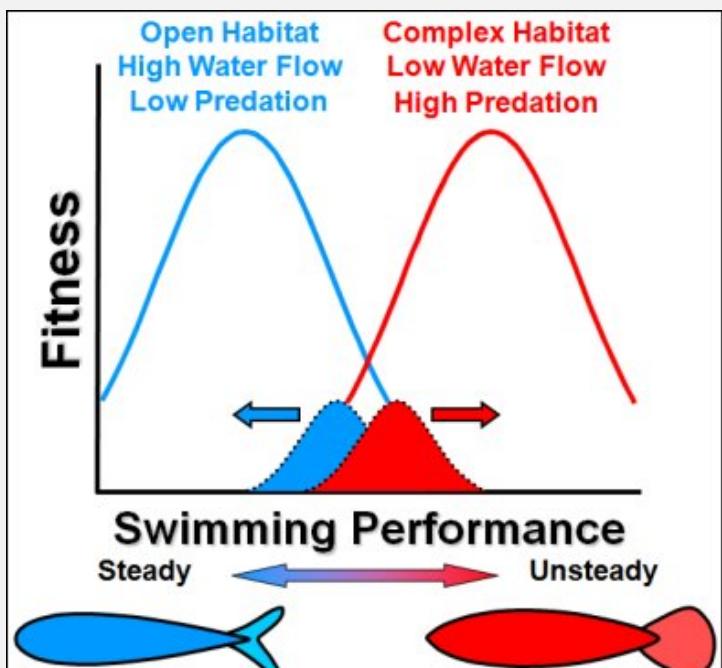
# Migration

Equal migration and no selection

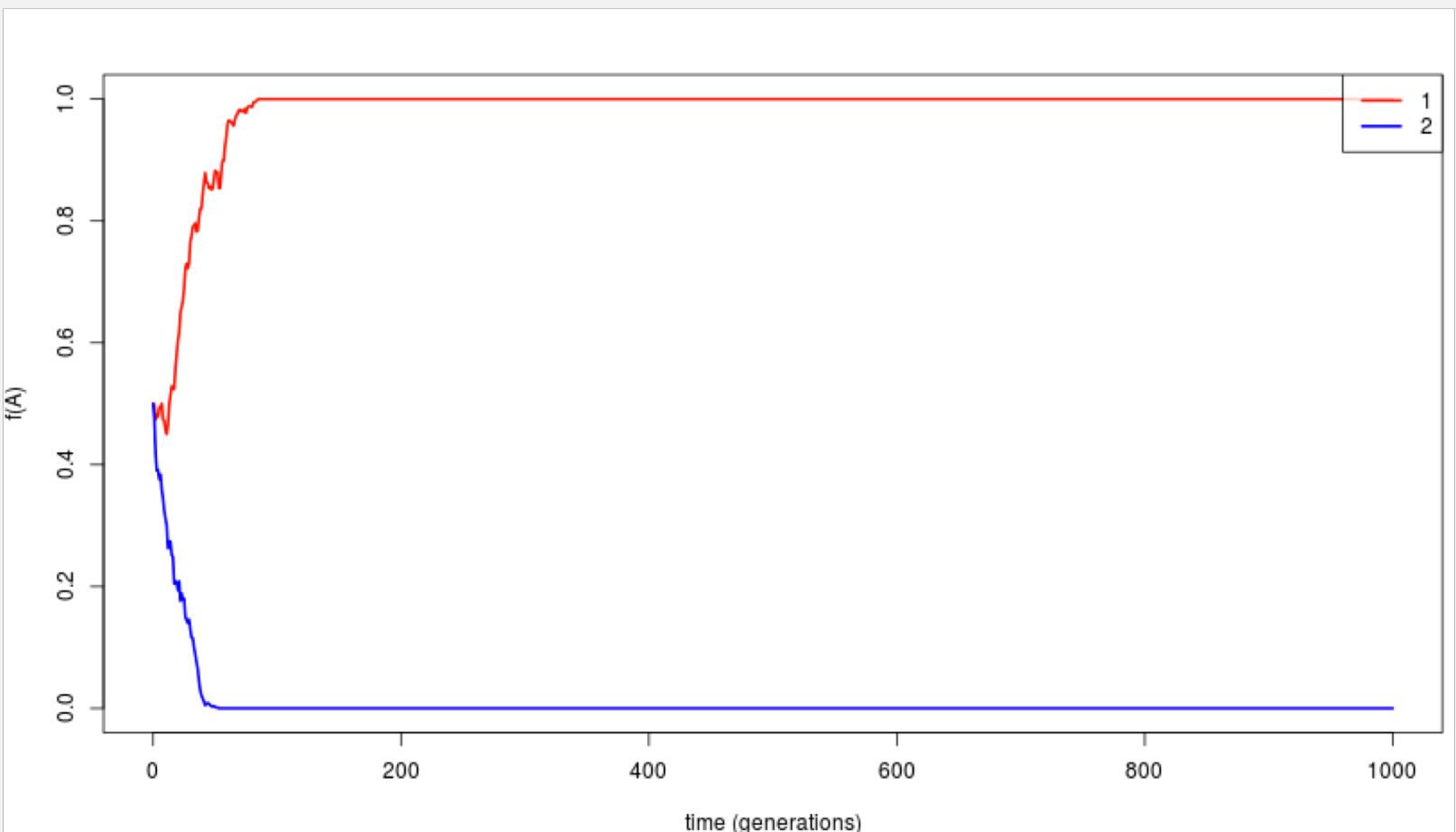


# Selection

Divergent selection without migration



<http://gambusia.zo.ncsu.edu/locomotion.html>

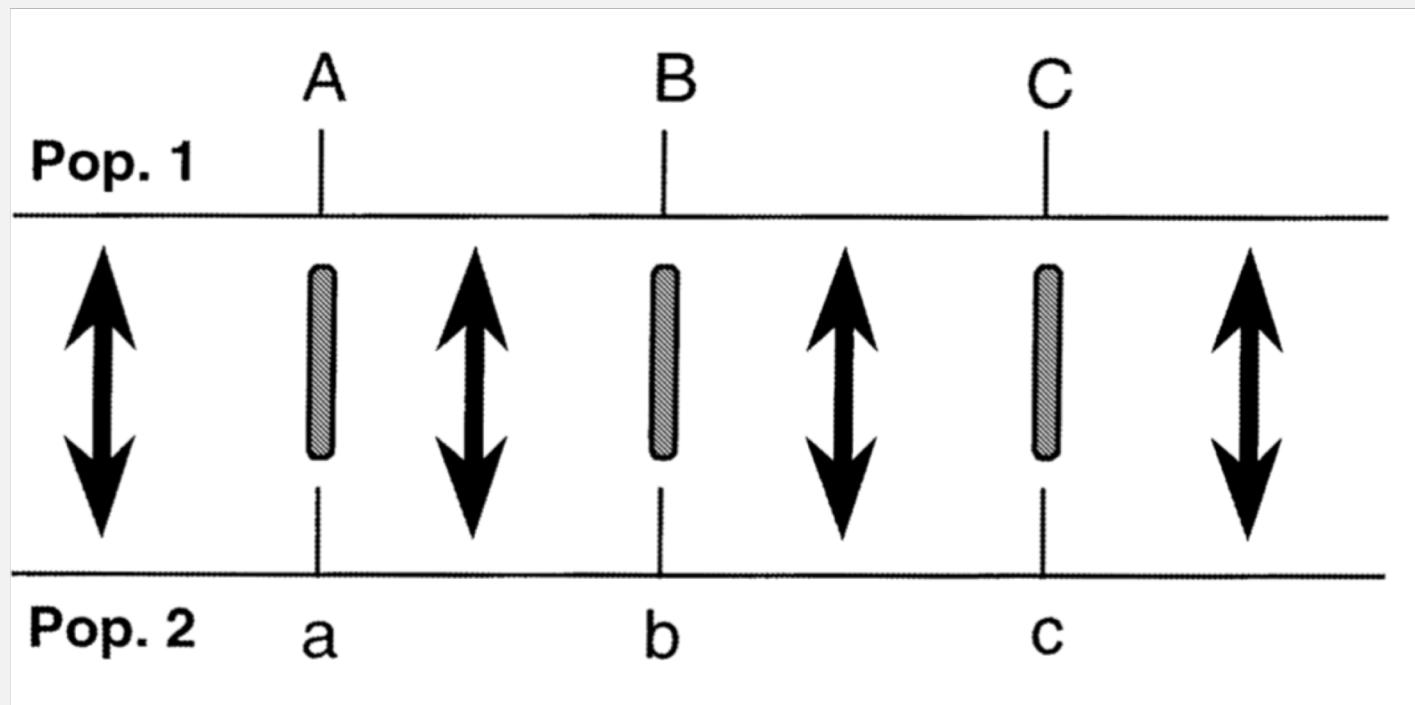


<https://phytools.shinyapps.io/migration-selection-drift/>

# Genomics of speciation

## Four-Phase model:

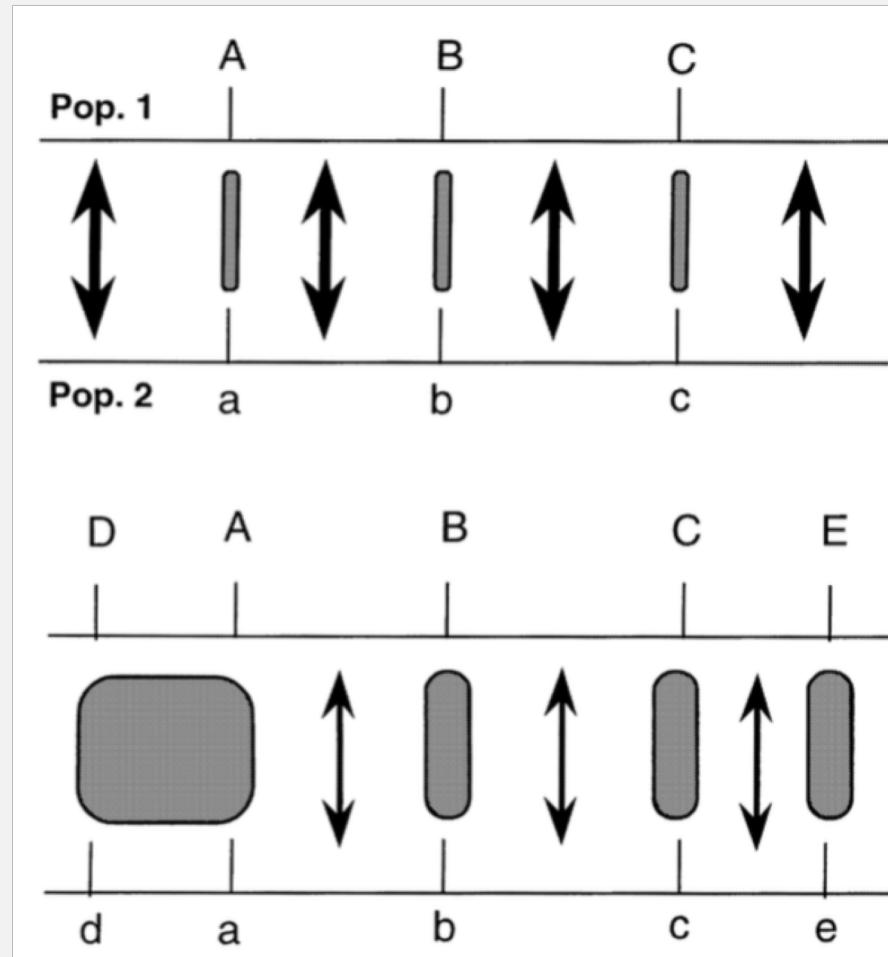
- I. Direct divergent selection (DS)



# Genomics of speciation

## Four-Phase model:

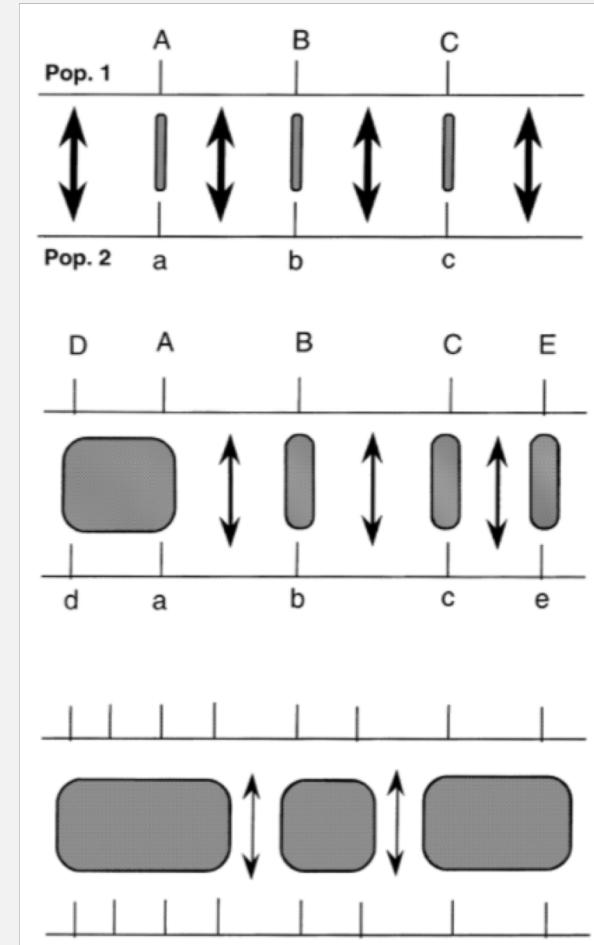
1. Direct divergent selection (DS)
2. Divergence hitchhiking (DH)



# Genomics of speciation

## Four-Phase model:

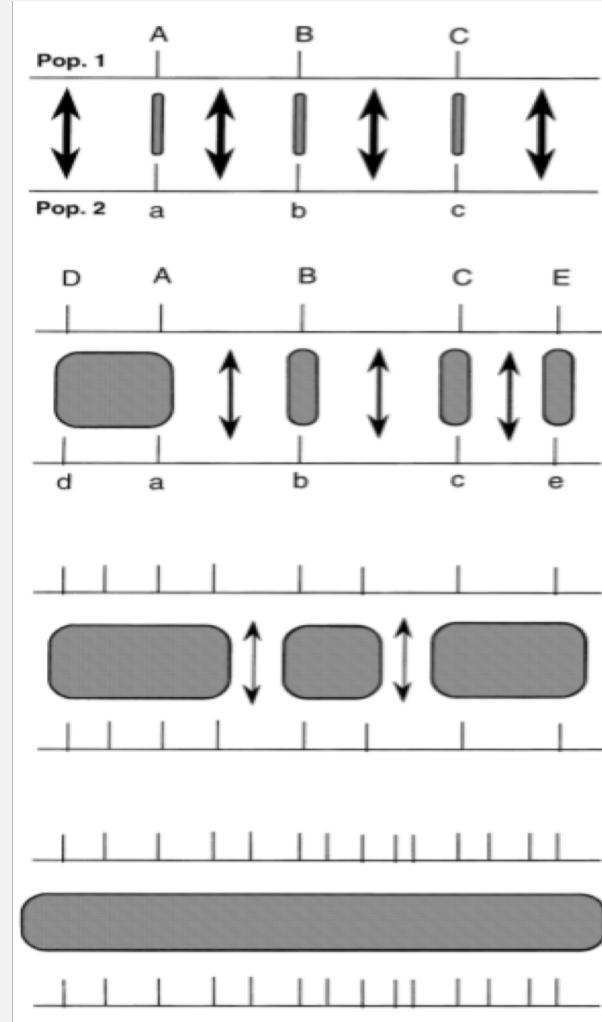
1. Direct divergent selection (DS)
2. Divergence hitchhiking (DH)
3. Genome hitchhiking (GH)



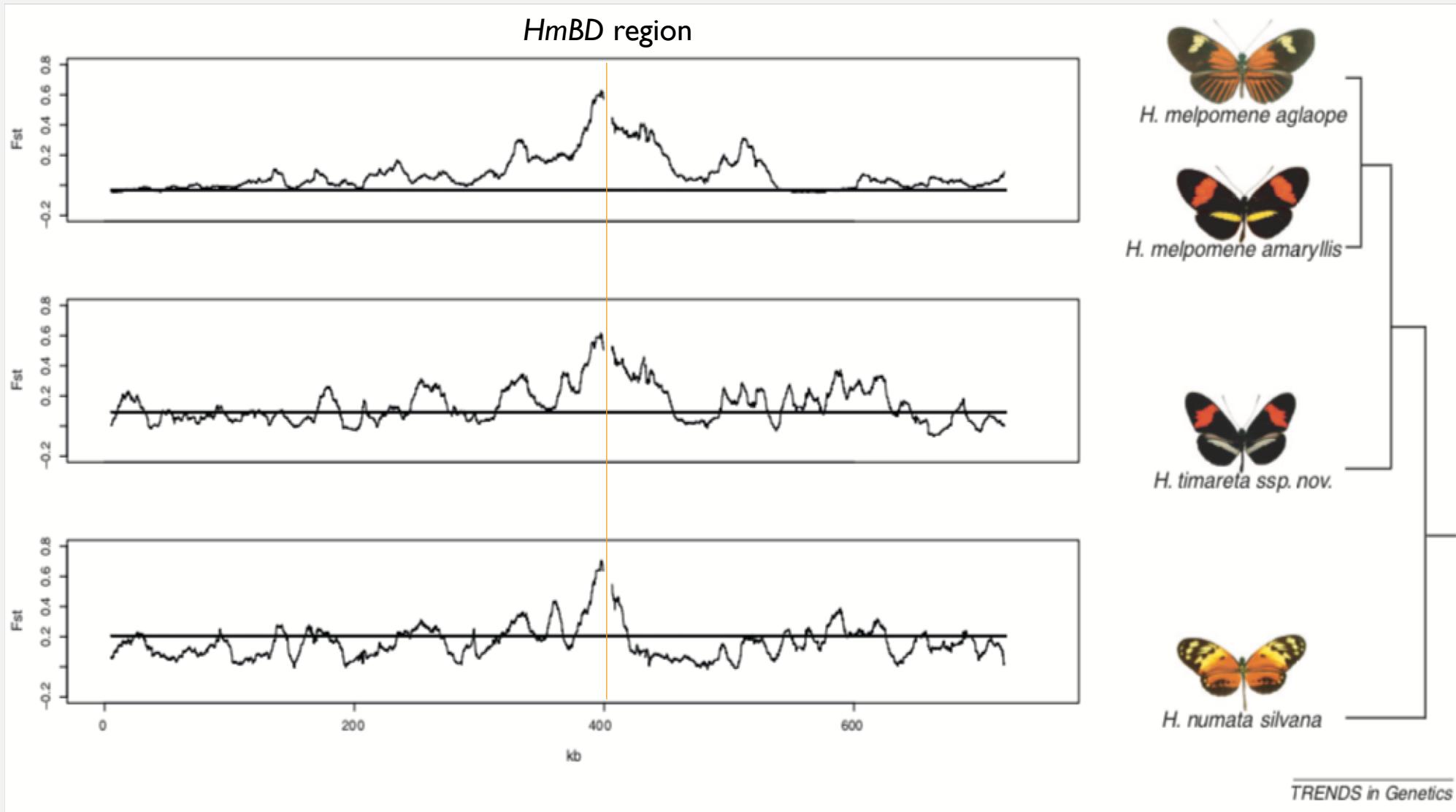
# Genomics of speciation

## Four-Phase model:

1. Direct divergent selection (DS)
2. Divergence hitchhiking (DH)
3. Genome hitchhiking (GH)
4. Speciation divergence

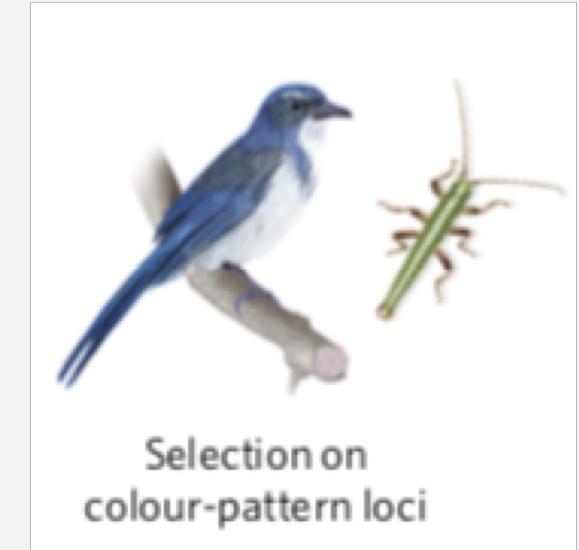
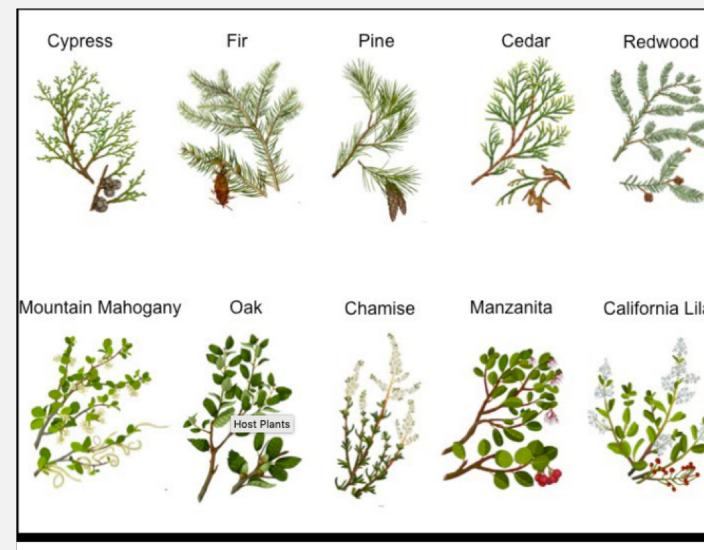
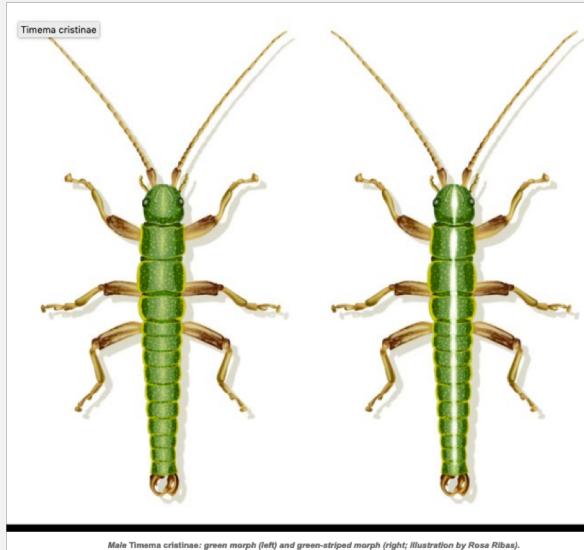


# Genomic divergence

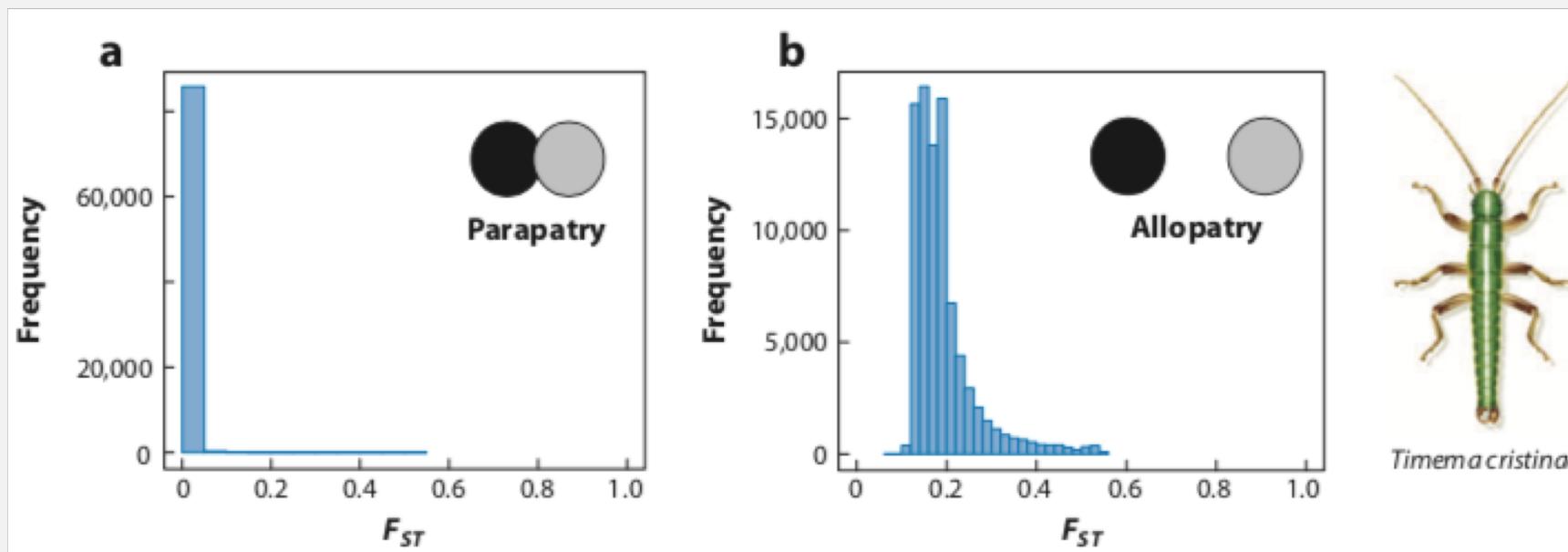


# Genomic divergence

## Genus *Timema*

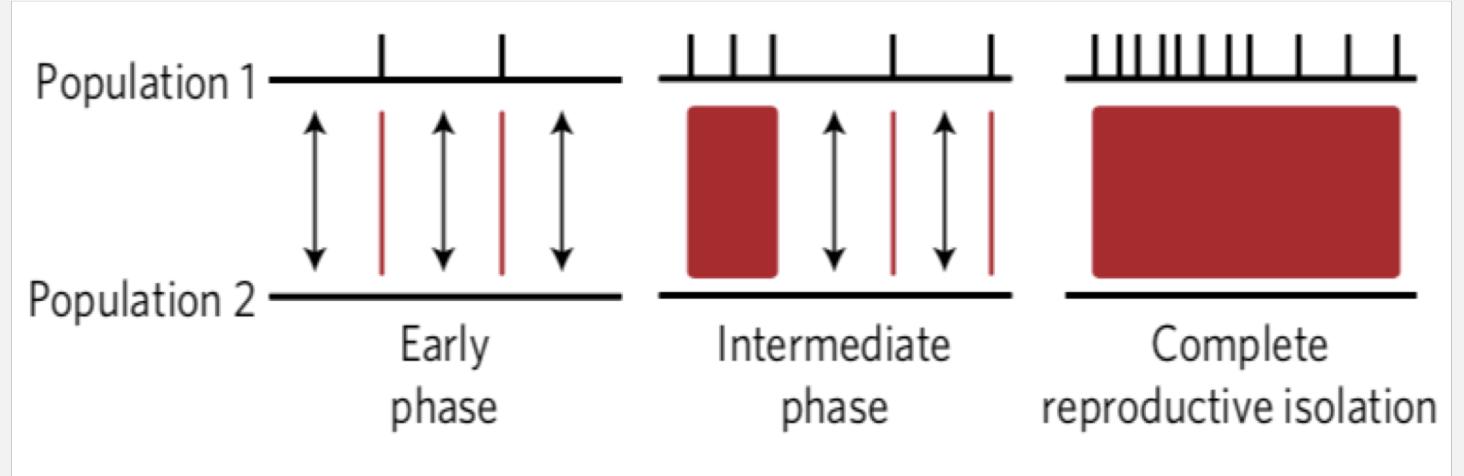


# Genomic divergence



# Phylogenetics

- Allele frequencies: within species
- New character states (mutations): between species



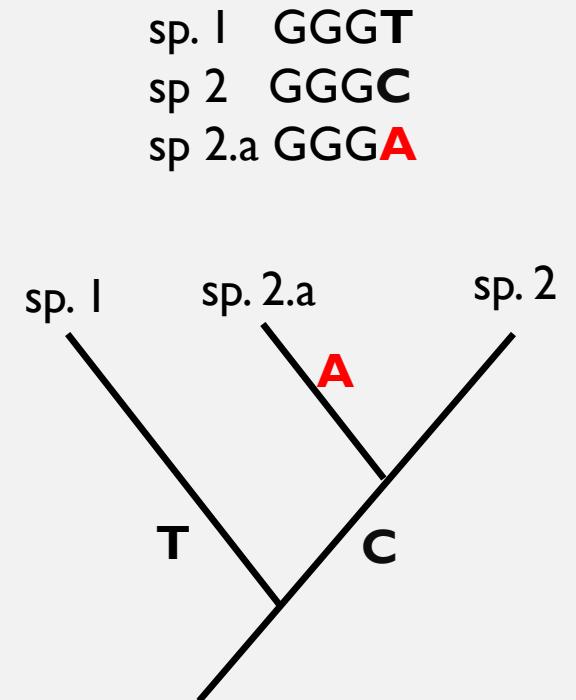
Riesch (2017) *Nature*

# Phylogenetics

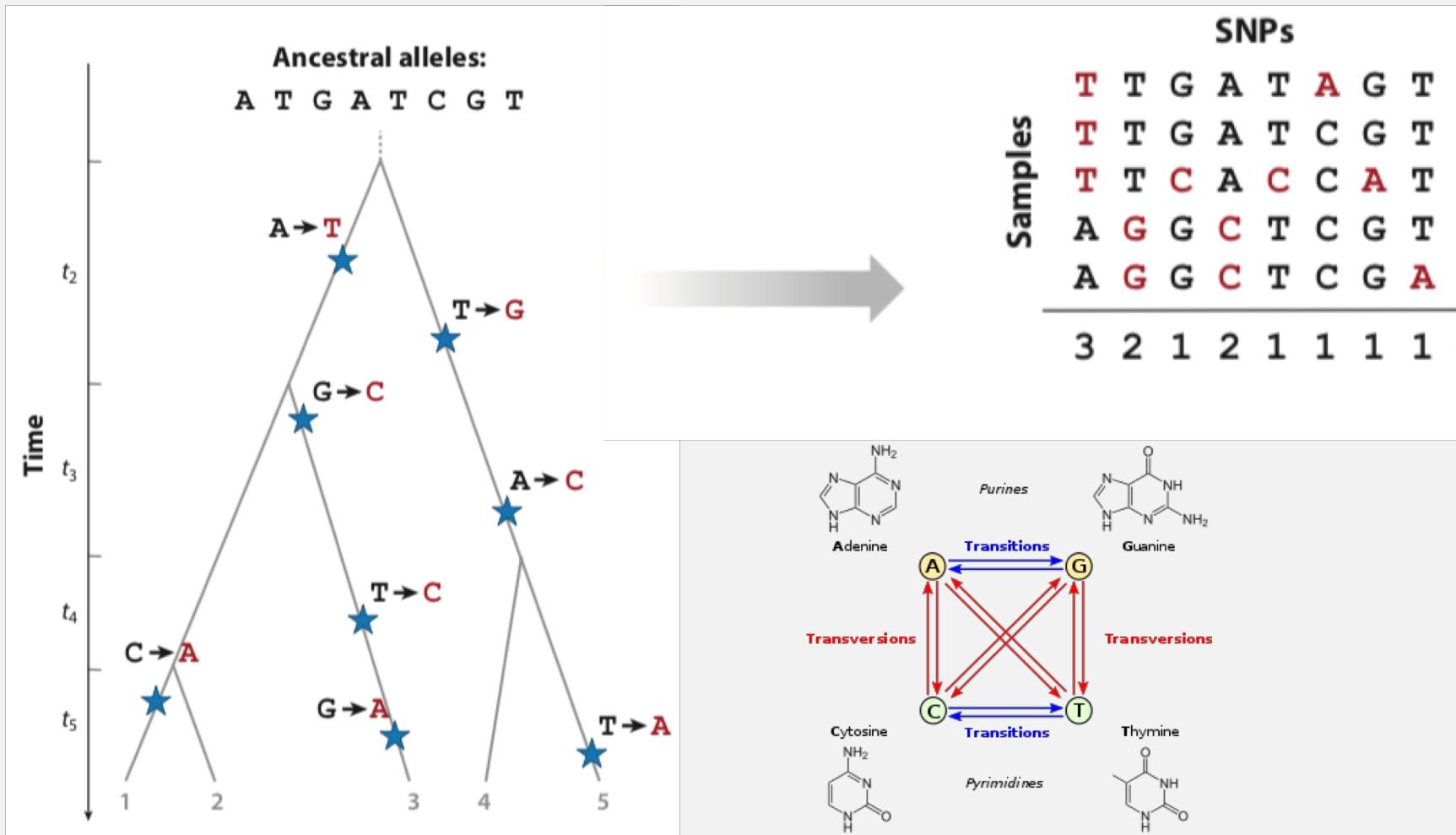
- New sp. I
  - I. GGGT
  - 2. GGGT
  - 3. GGGT
  - 4. GGGT
- Ancestral Pop
  - I. GGGT
  - 2. GGGT
  - 3. GGGC
  - 4. GGGC
- New sp. 2
  - I. GGGC
  - 2. GGGC
  - 3. GGGC
  - 4. GGGC

Fst = 1

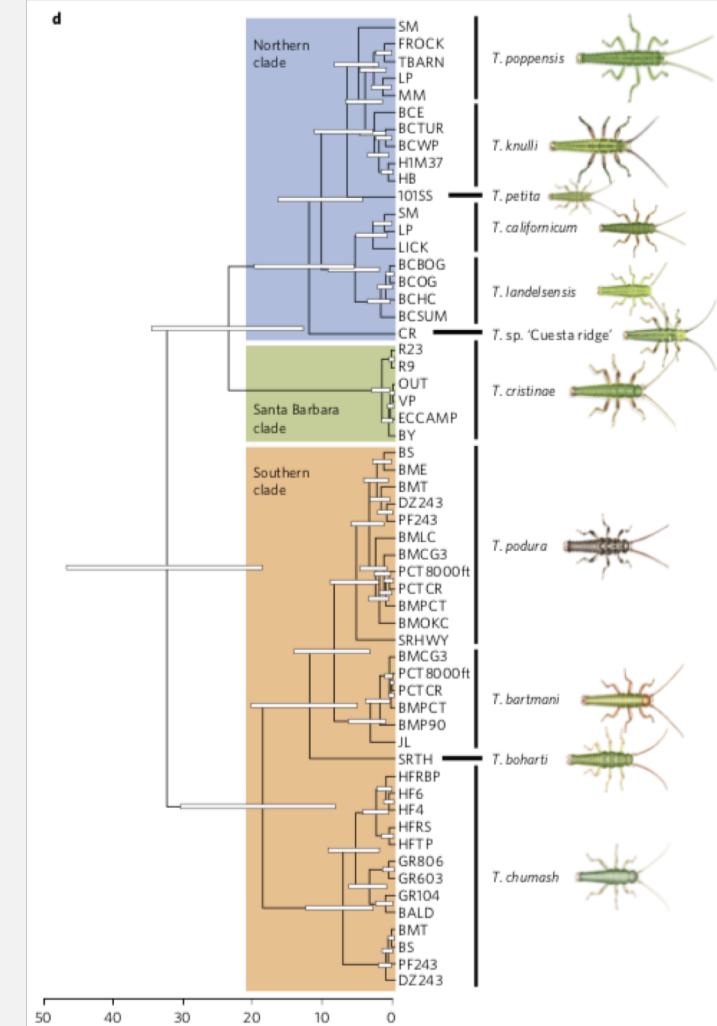
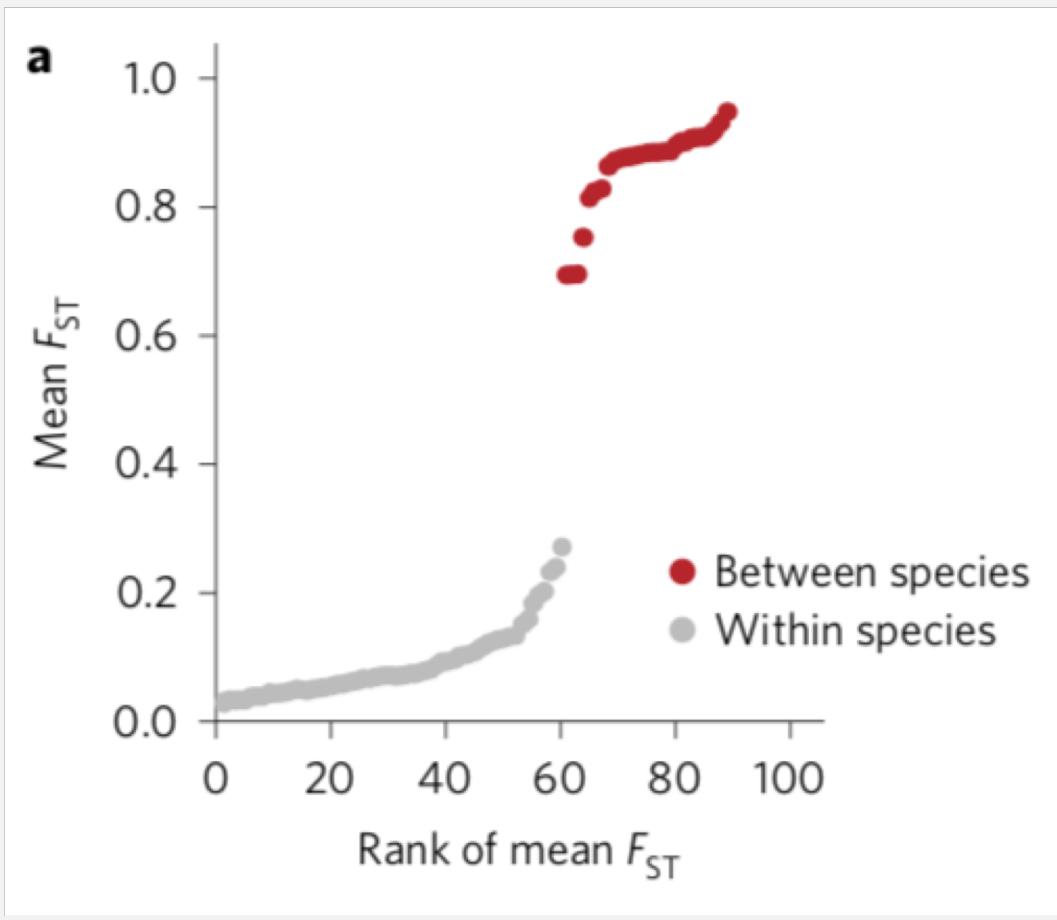
- New sp 2.a
  - I. GGGA
  - 2. GGGA
  - 3. GGGA
  - 4. GGGA
- New pop 2
  - I. GGGA
  - 2. GGGC
  - 3. GGGC
  - 4. GGGC



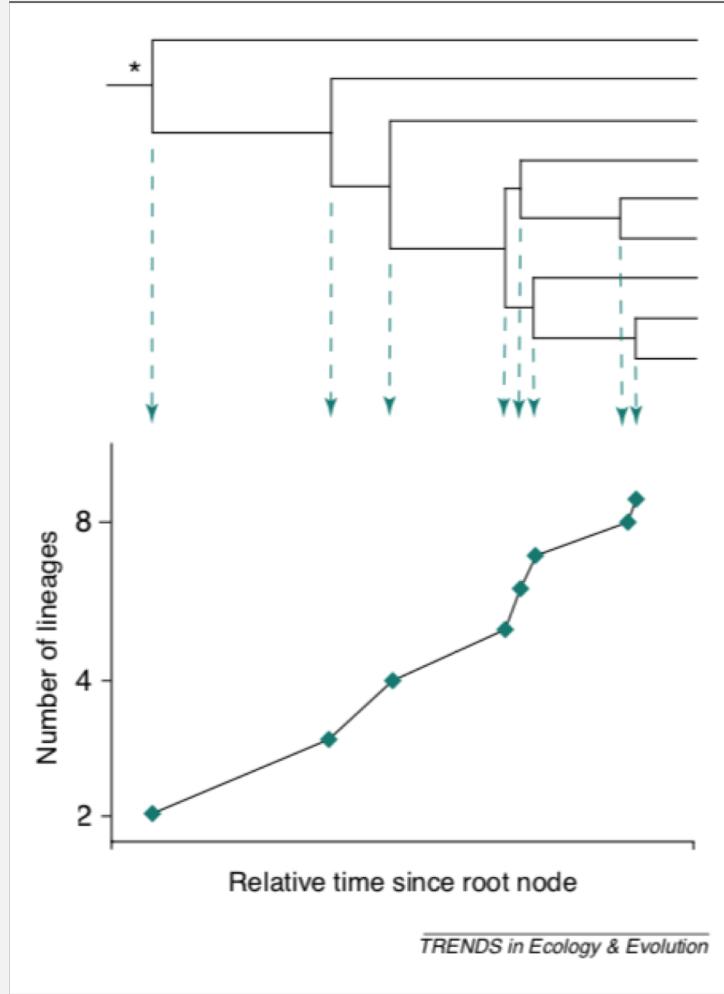
# Phylogenetics



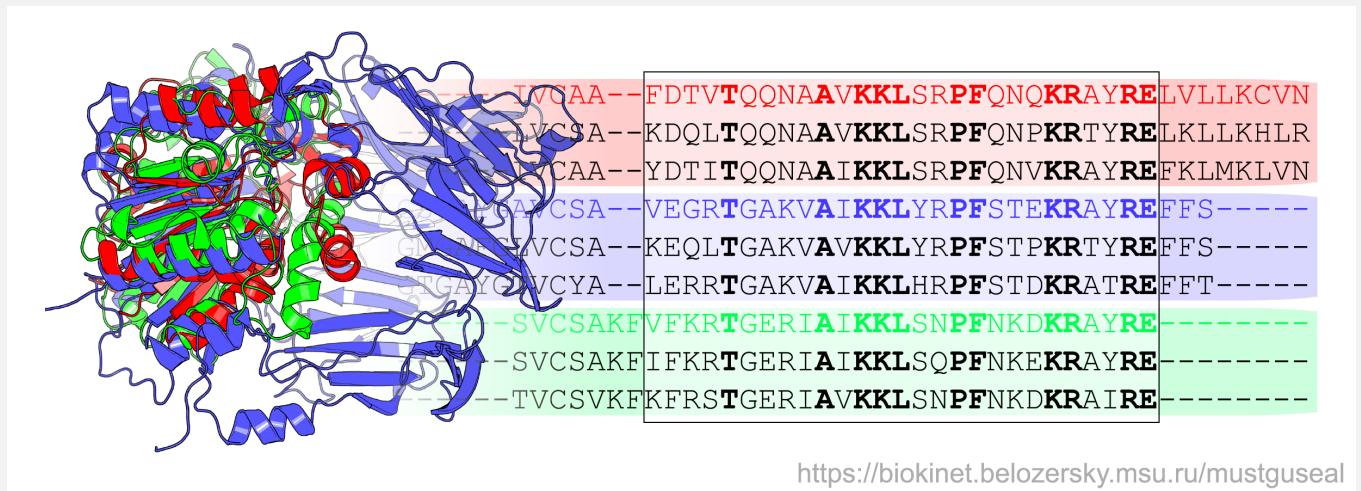
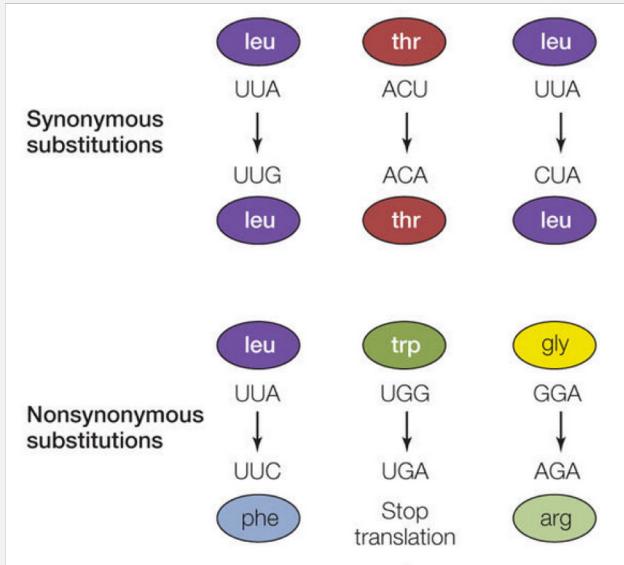
# Phylogenetics



# Phylogenetics



Barraclough & Sean Nee (2001)



<https://biokinет.belozersky.msu.ru/mustguseal>

# Phylogenetics

- **Synonymous vs non-synonymous**
- **dN/dS ratio**

dN = number of non-synonymous changes **per non-synonymous site**

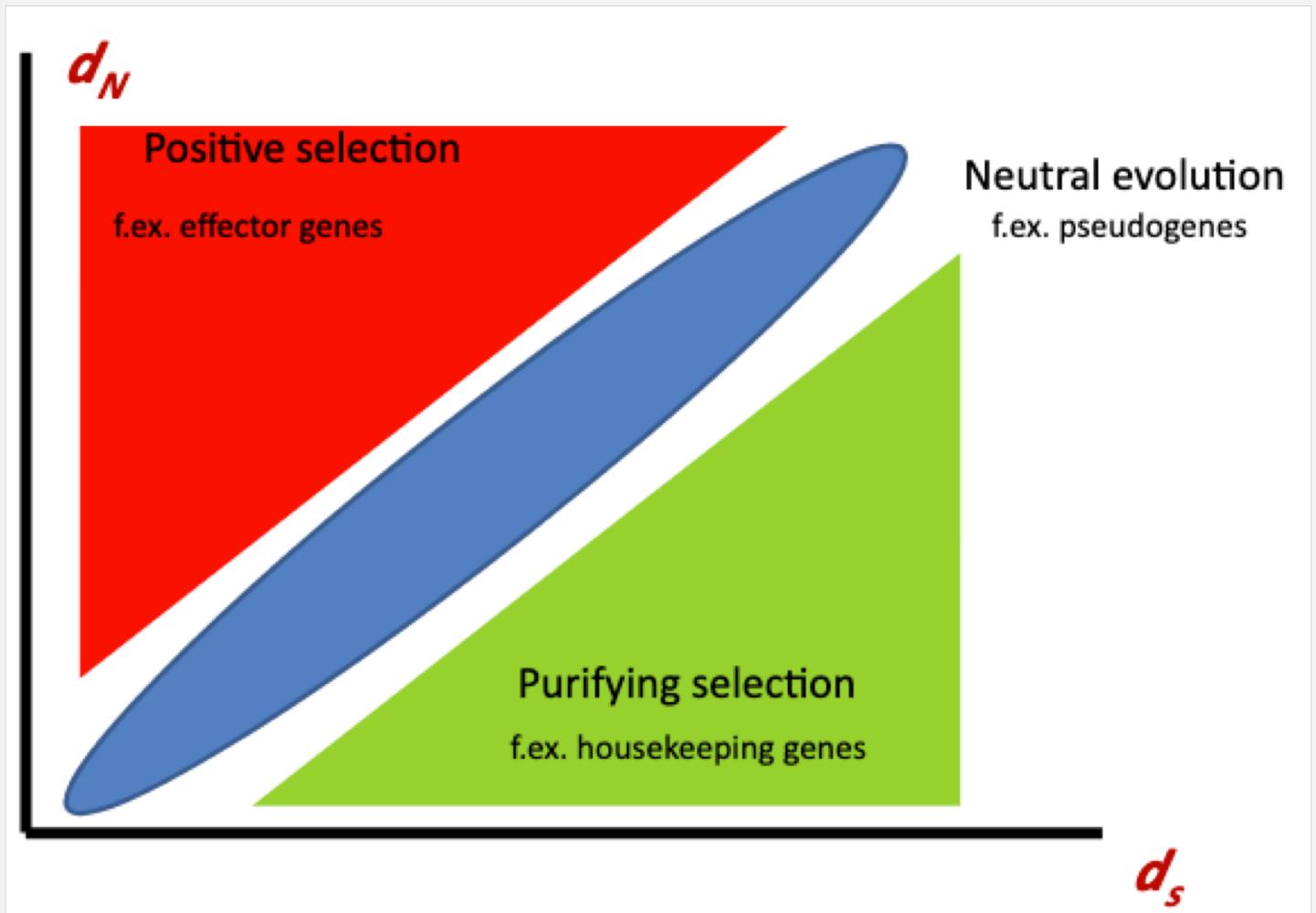
dS = number of synonymous changes **per synonymous site**

		2nd letter in the codon								
		U	C	A	G					
		U	UUU   Phe (F) UUC   UUA   Leu (L) UUG	C	UCU   Ser (S) UCC   UCA   UCG	A	UAU   Tyr (Y) UAC   UAA   STOP UAG   STOP	G	UGU   Cys (C) UGC   UGA   STOP UGG   Trp (W)	U C A G
		C	CUU   CUC   Leu (L) CUA   CUG	CCU   CCC   CCA   CCG	CAU   His (H) CAC   CAA   Gln (Q) CAG	CGU   CGC   CGA   CGG	CGU   Arg (R) CGC   CGA   CGG	U C A G		
		A	AUU   AUC   AUA   AUG   Met (M) START	ACU   Ile (I) ACC   ACA   ACG	AAU   Asn (N) AAC   AAA   AAG	AGU   Ser (S) AGC   AGA   Arg (R) AGG	AGU   Ser (S) AGC   AGA   Arg (R) AGG	U C A G		
		G	GUU   GUC   GUA   GUG	GCU   Val (V) GCC   GCA   GCG	GAU   Asp (D) GAC   GAA   Glu (E) GAG	GGU   GGC   GGA   GGG	Gly (G)	U C A G		

# Phylogenetics

## Synonymous vs non-synonymous

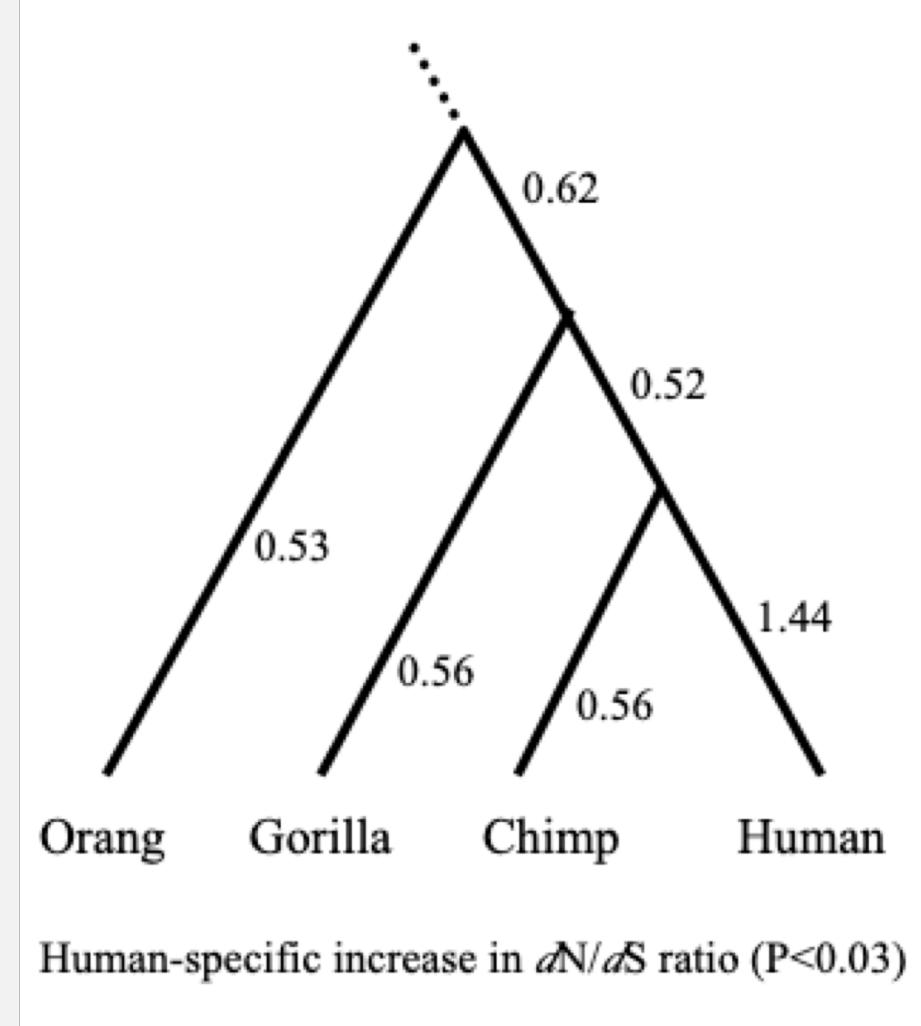
- $dN/dS = 1$   
Neutral selection
- $dN/dS < 1$   
Negative selection
- $dN/dS > 1$   
Positive selection



# Phylogenetics

## Synonymous vs non-synonymous

- *ASPM* gene
- Brain size



# Discussion papers

HYBRID SPECIATION

## Rapid hybrid speciation in Darwin's finches

Sangeet Lamichhaney,<sup>1\*</sup> Fan Han,<sup>1</sup> Matthew T. Webster,<sup>1</sup> Leif Andersson,<sup>1,2,3†</sup>  
B. Rosemary Grant,<sup>4</sup> Peter R. Grant<sup>4</sup>

RESEARCH ARTICLE

## Multispecies Outcomes of Sympatric Speciation after Admixture with the Source Population in Two Radiations of Nicaraguan Crater Lake Cichlids

Andreas F. Kautt, Gonzalo Machado-Schiaffino, Axel Meyer\*

Department of Biology, University of Konstanz, Konstanz, Baden-Württemberg, Germany