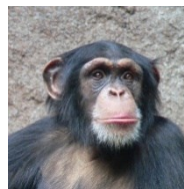
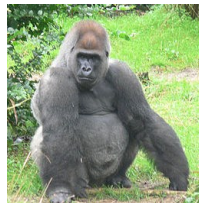
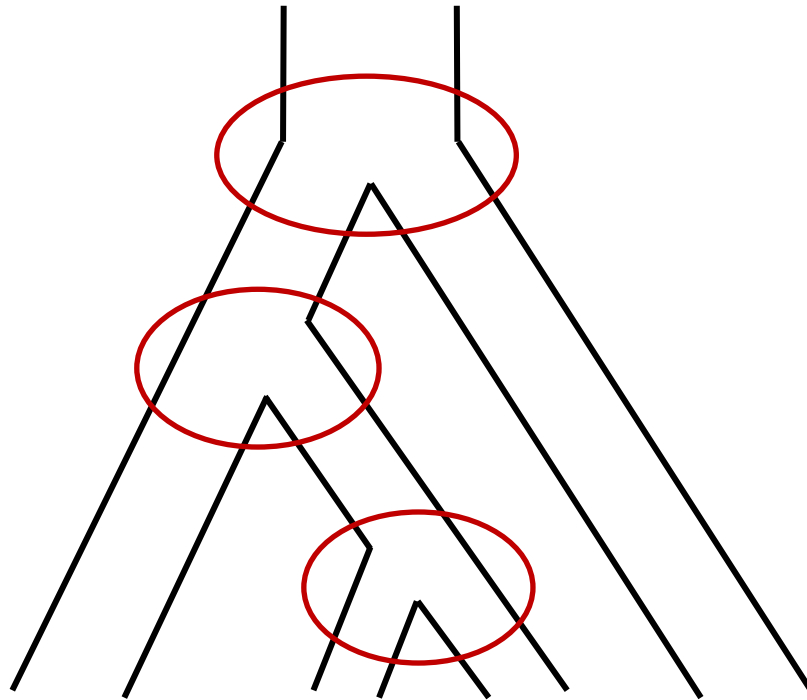
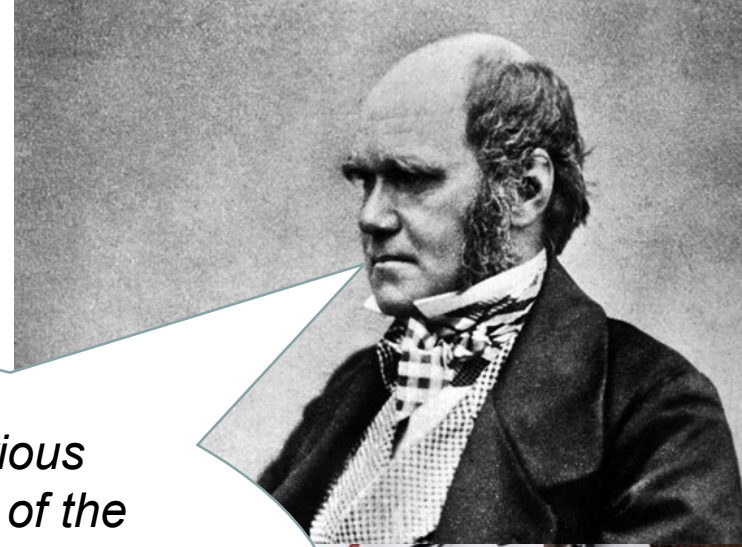


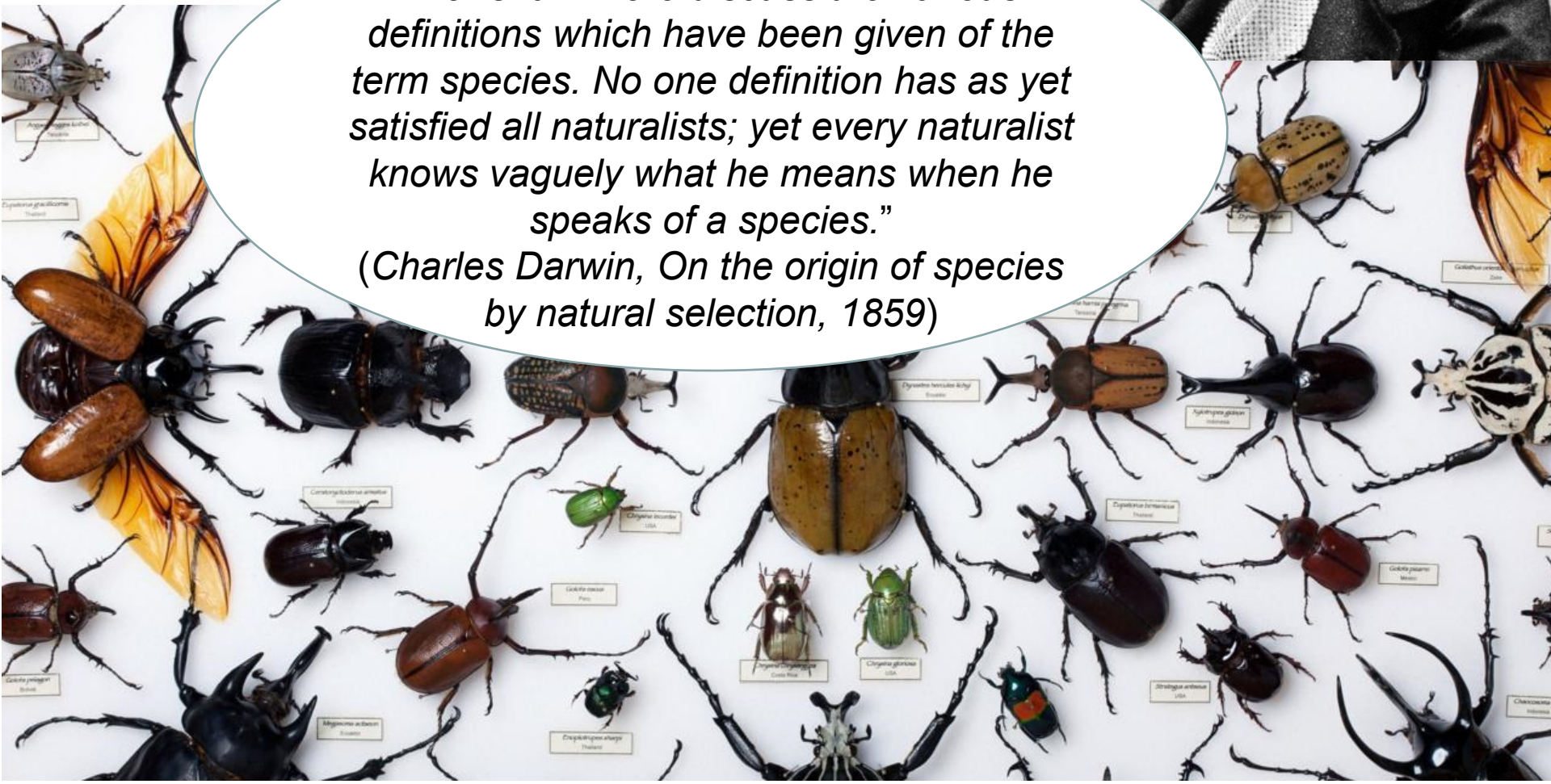
Speciation



Biological diversity is discontinuous.
Species objectively exist.
But how to define them?



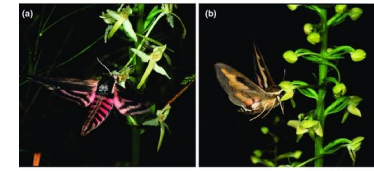
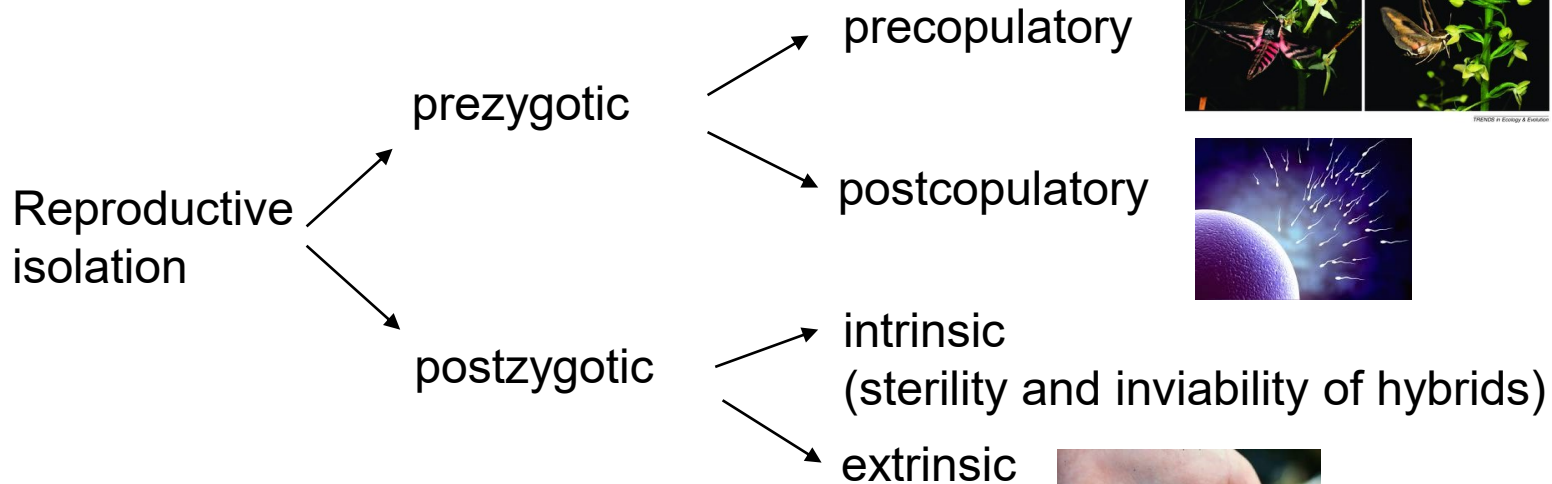
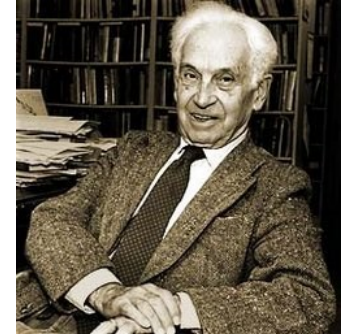
"Nor shall I here discuss the various definitions which have been given of the term species. No one definition has as yet satisfied all naturalists; yet every naturalist knows vaguely what he means when he speaks of a species."
(Charles Darwin, *On the origin of species by natural selection*, 1859)



What is species?

Biological species concept (Ernst Mayr, 1942)

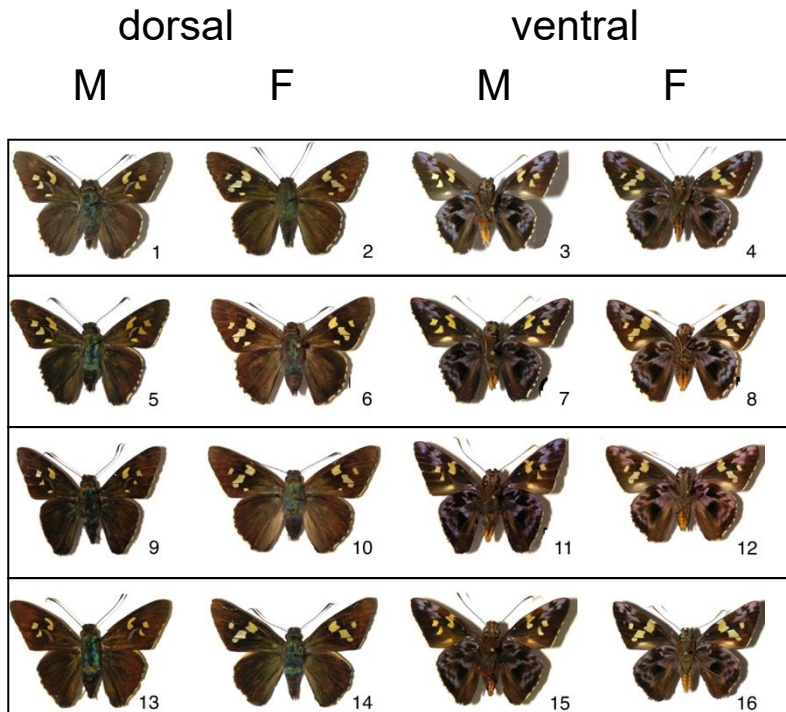
- A group of organisms that can successfully interbreed and produce fertile offspring. A species' integrity is maintained by interbreeding within a species as well as by reproductive barriers between organisms in different species.



Morphological species concept

Groups of individuals that are morphologically similar to one another and are morphologically distinct from other such groups..

Cryptic species



Four cryptic species of *Perichares* butterflies

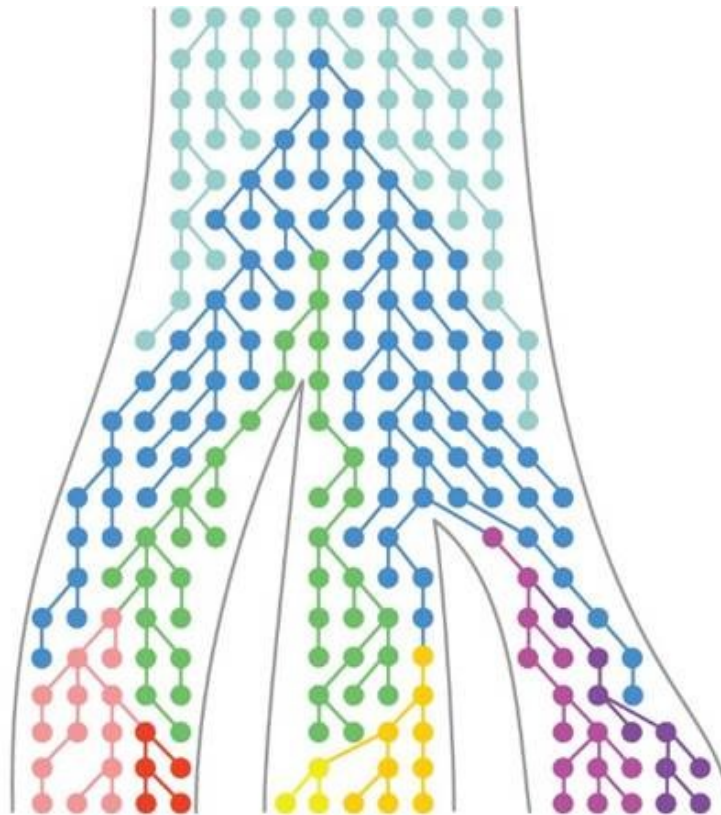
Polytypic species



The western yellow wagtail (*Motacilla flava*)

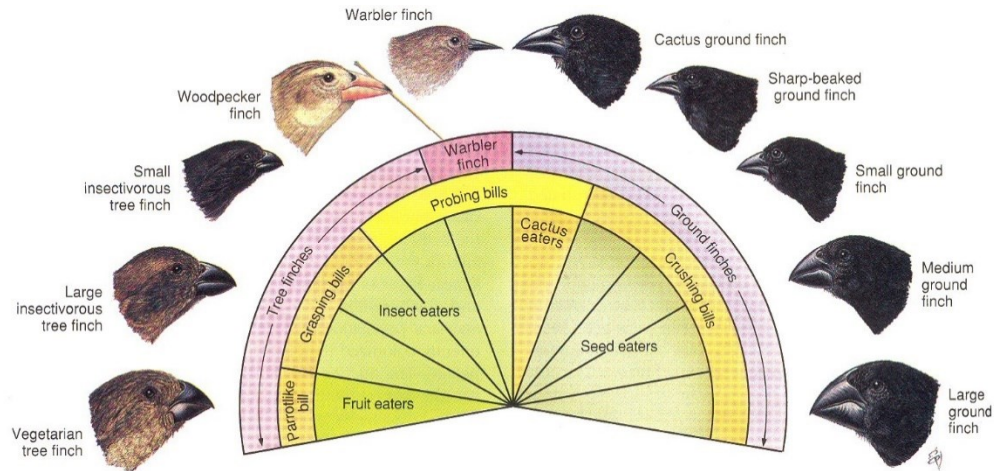
Genealogical/phylogenetic species concept

A group of organisms whose members are all more closely related to each other than they are to any organisms outside the group and share a unique common ancestor.



Ecological species concept

A group of organisms that share the ecological niche.

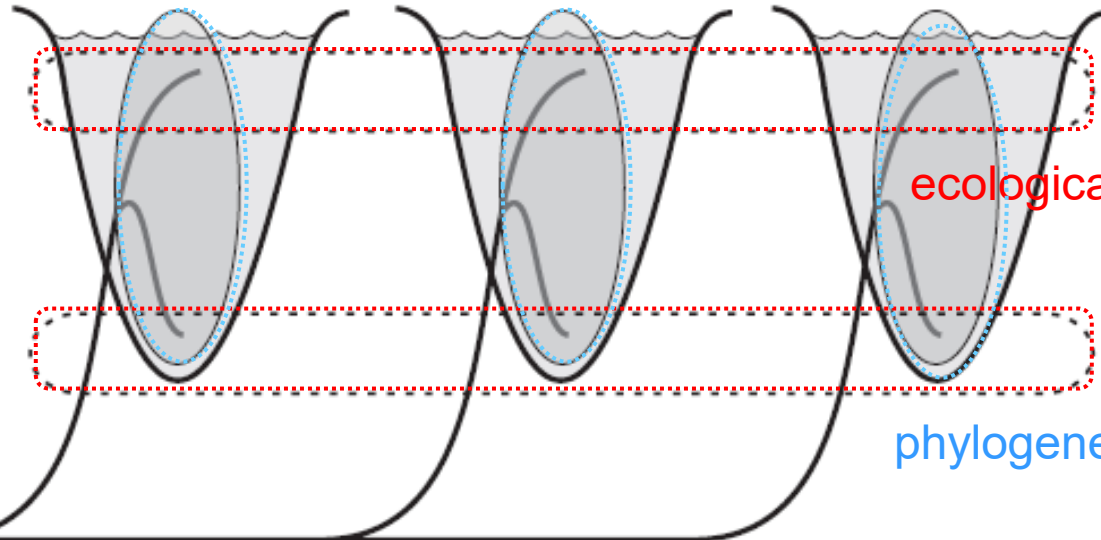


Three-spined stickleback

pelagic form



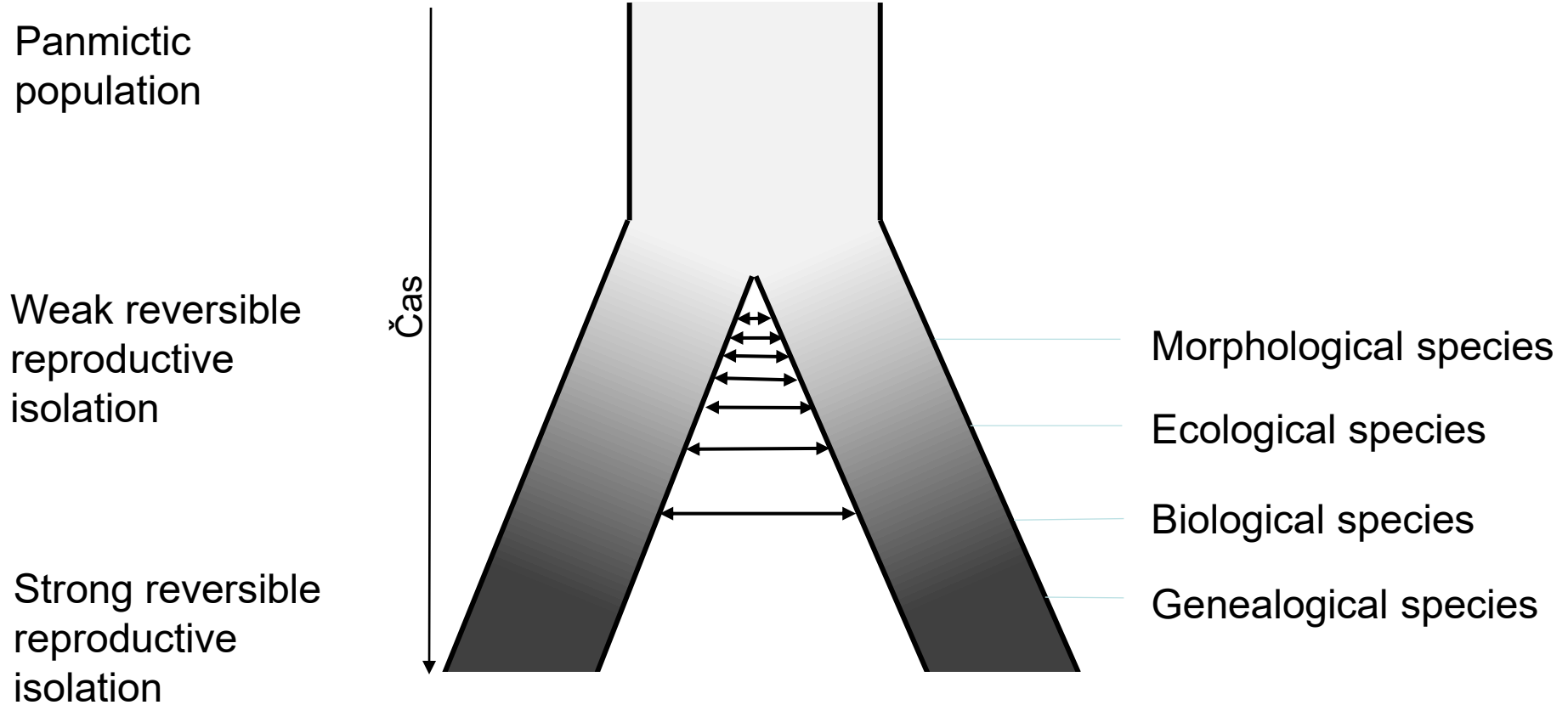
benthic form



ecological species

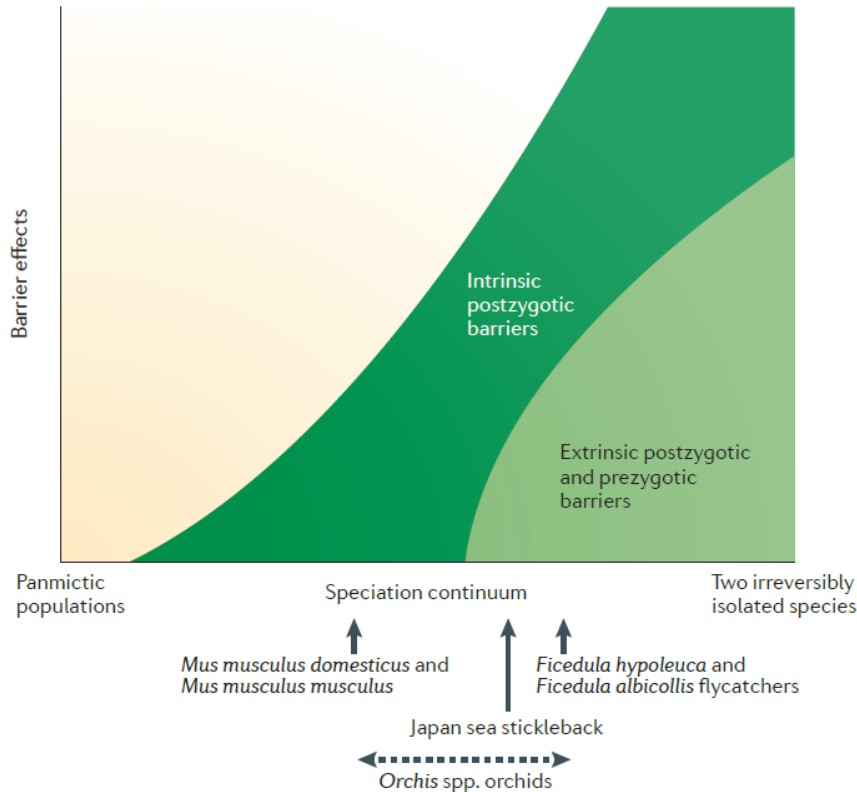
phylogenetic species

Speciation continuum

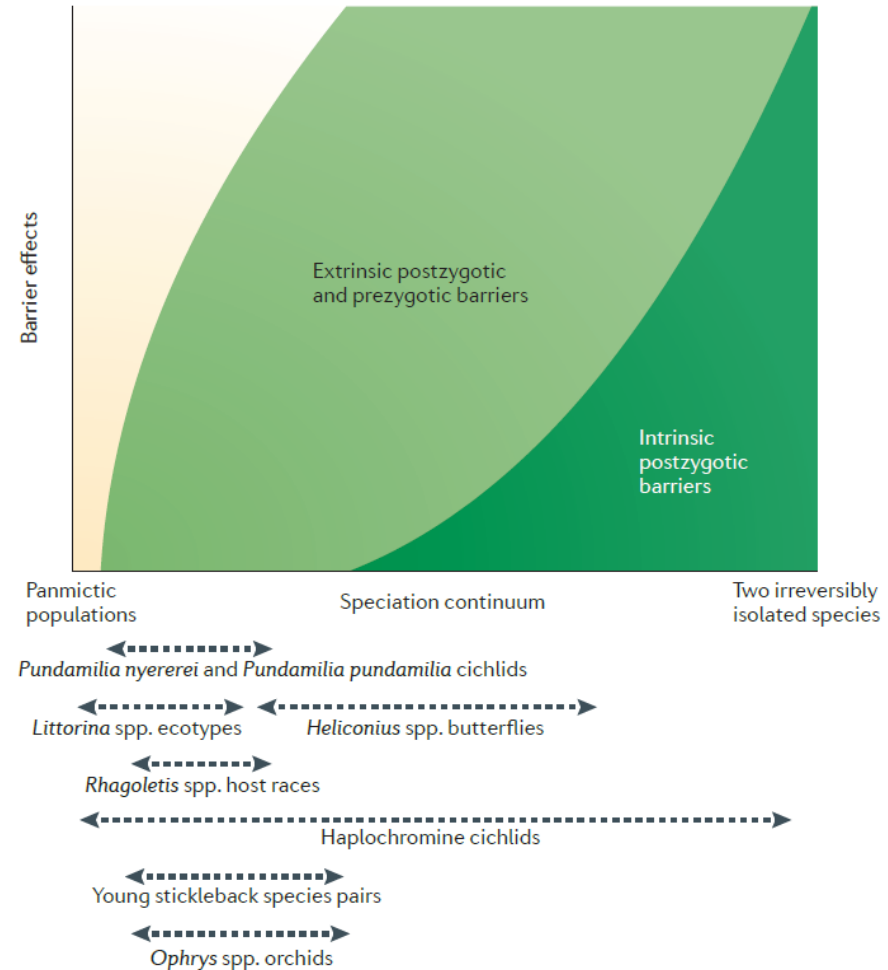


Speciation driven by intrinsic barriers vs. speciation driven by extrinsic postzygotic and prezygotic barriers (ecological speciation and speciation by sexual selection)

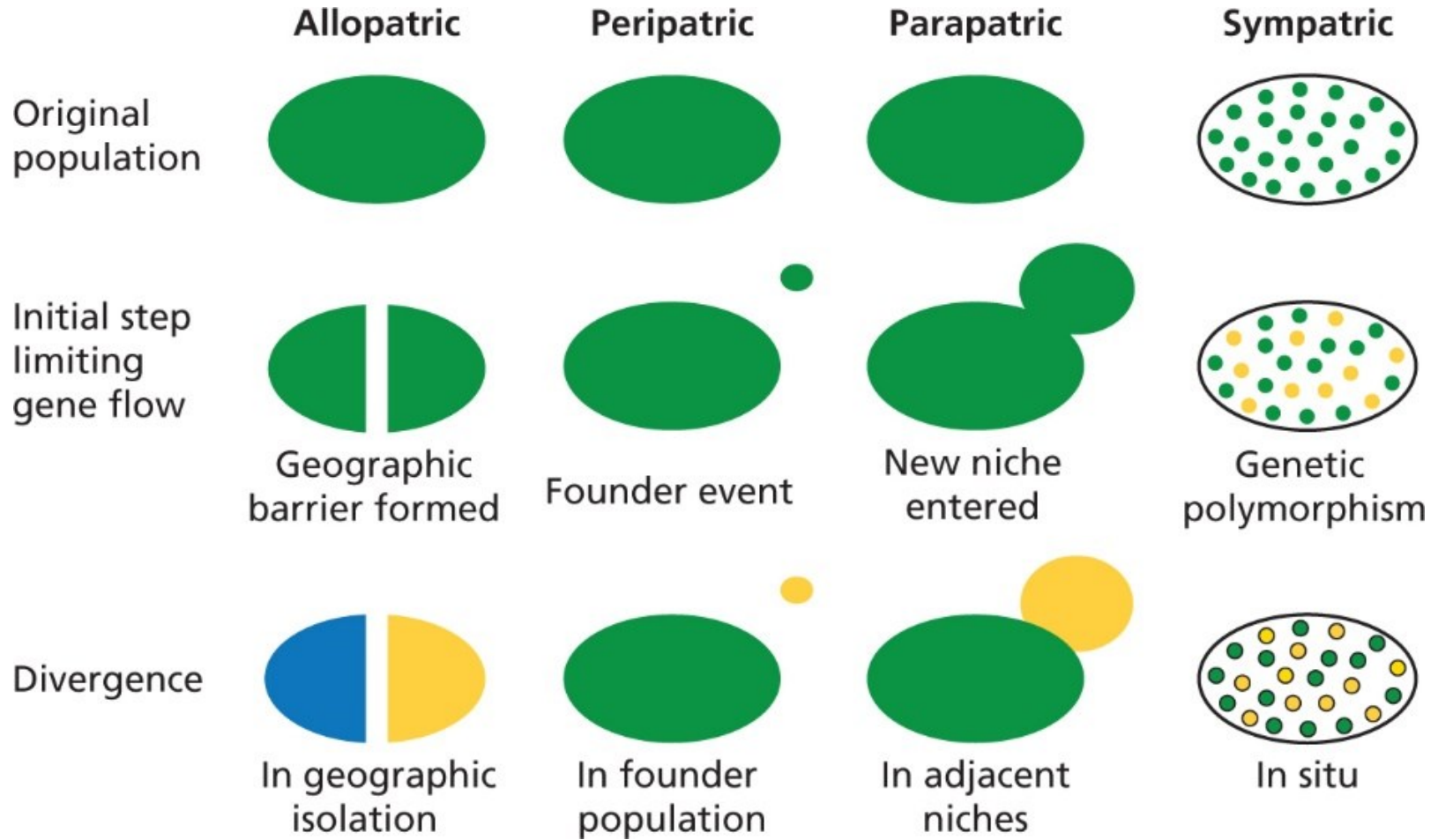
b Speciation driven by intrinsic barriers



a Speciation driven by divergent selection



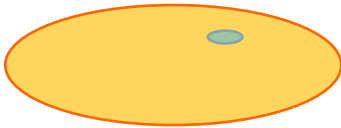
Speciation



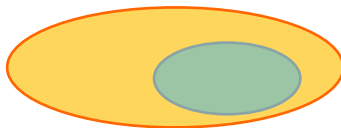
Sympatric speciation



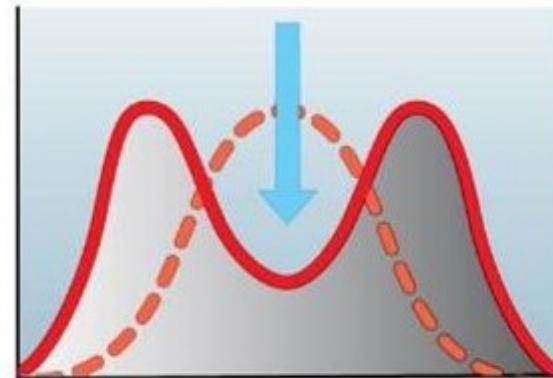
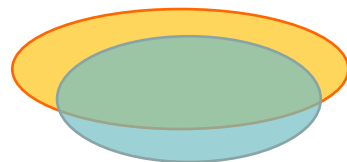
Charles Darwin



Genetic polymorphism



Reproductive isolation arise in the face of gene flow in sympatry

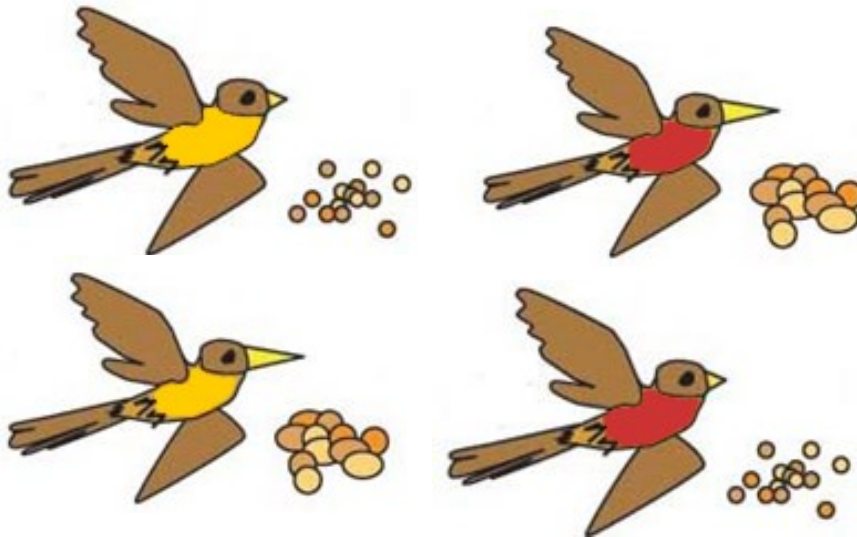


Disruptive selection
Assortative mating

Problems with sympatric speciation

Recombination between genes for:

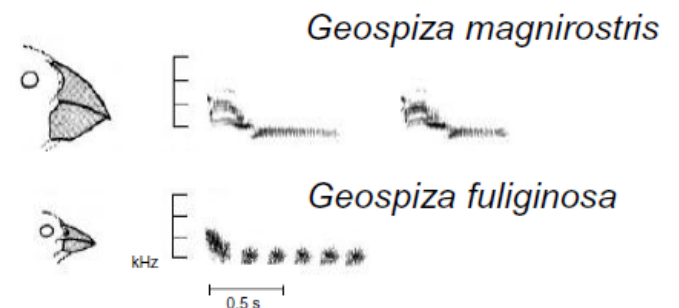
- (1) ecological traits under disruptive selection,
- (2) female preferences



Magic traits



vrtnice (*Rhagoletis pomonella*)



Sympatric speciation



Two palm species on the island of Lord Howe in Pacific.

Savolainen et al. (2006), Nature.



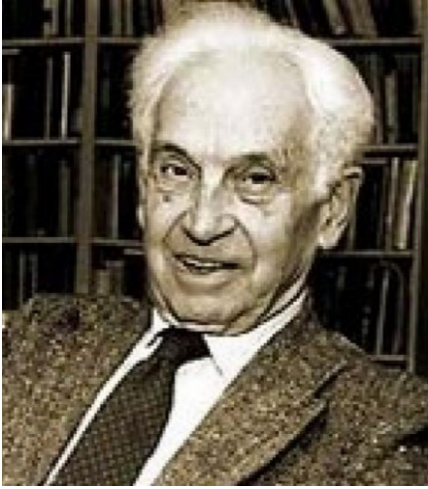
● *Amphilophus citrinellus*



■ *Amphilophus zalius*

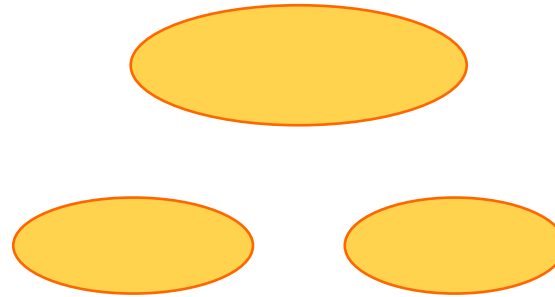
Cichlids in crater lakes

Barluenga et al. (2006), Nature.



Ernst Mayr

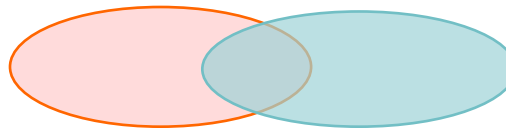
Allopatric speciation



Geographic barrier

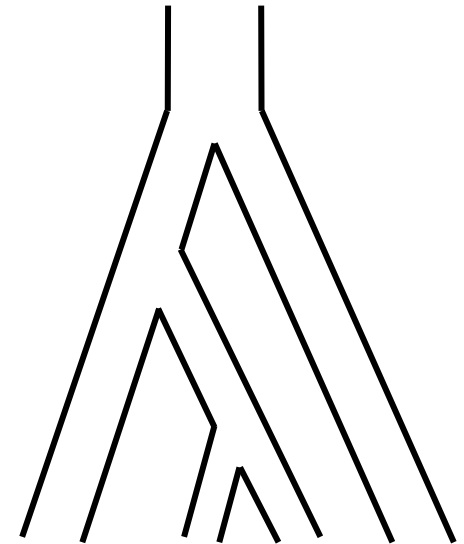


Reproductive barrier arise
without gene flow in
allopatry



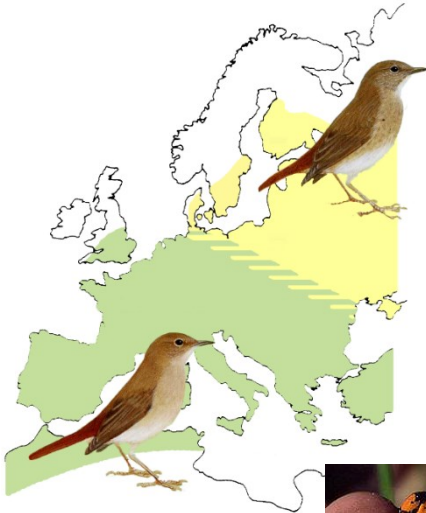
After secondary
contact species do not
interbreed or produce
sterile or inviable
hybrid progeny

Divergence
without gene flow

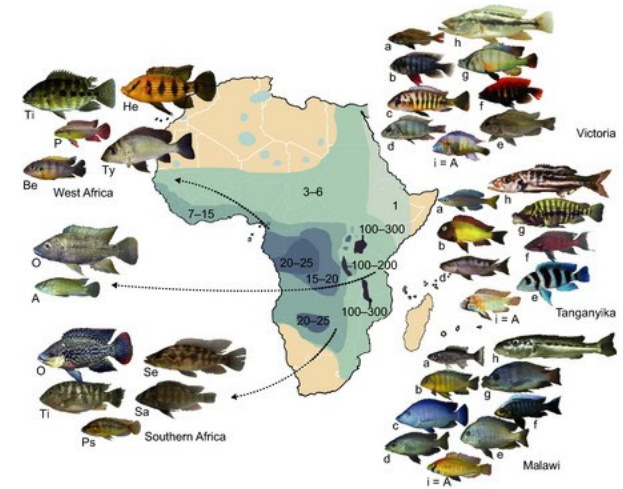


Interspecific hybridization after secondary contact

Luscinia megarhynchos
Luscinia luscinia

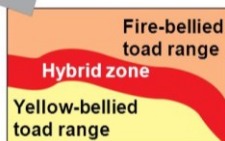


Mus musculus domesticus
Mus musculus musculus



Cichlids

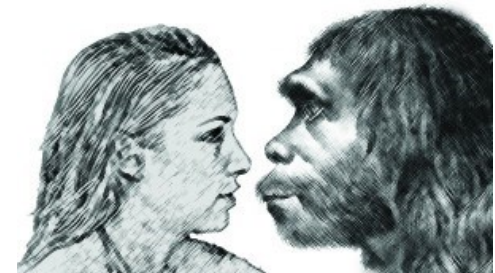
Bombina bombina
Bombina variegata



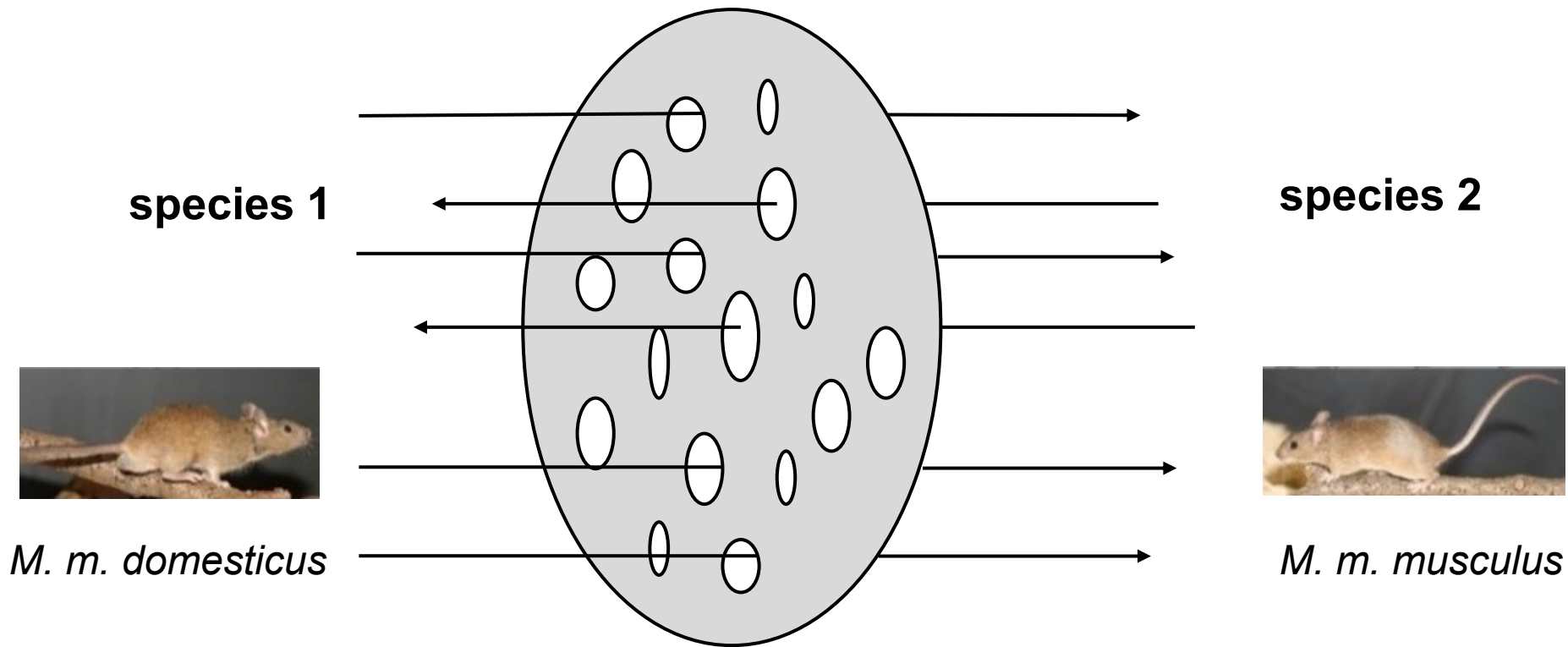
Darwin Finches



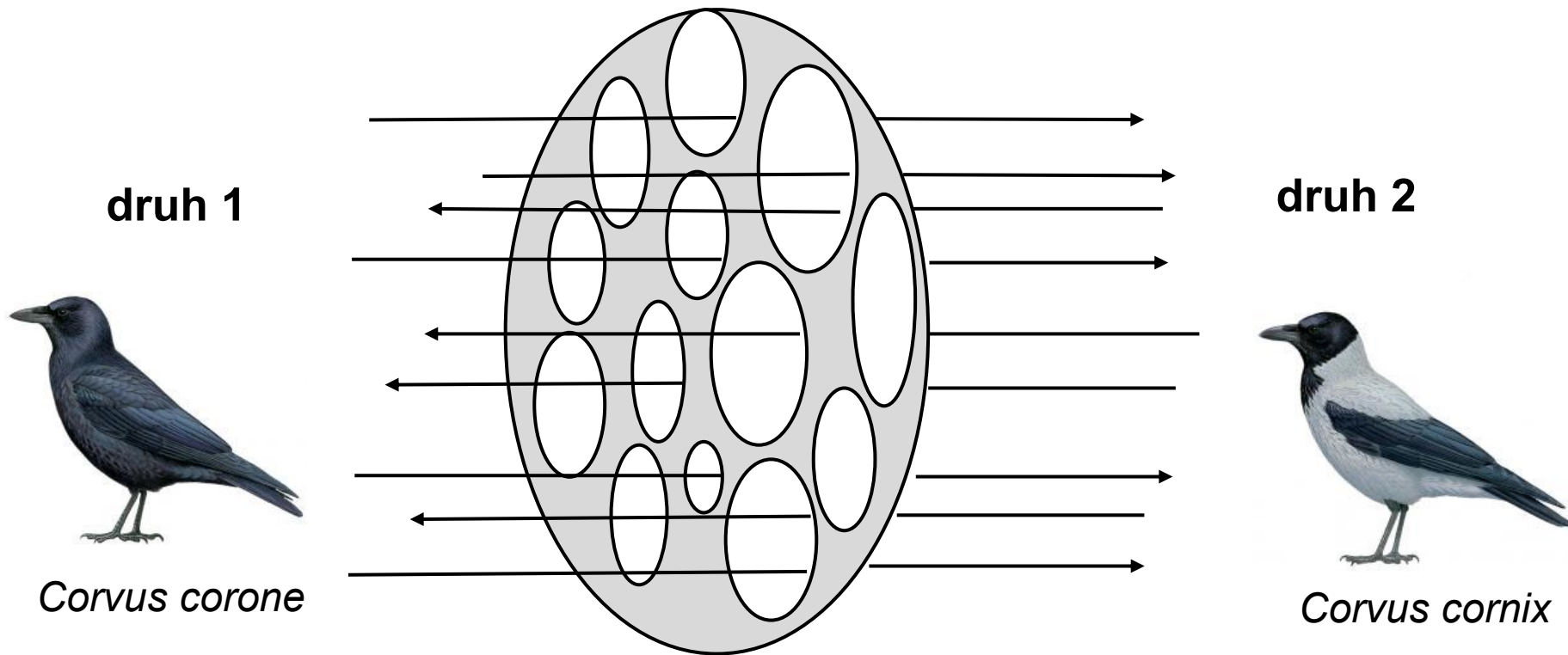
Homo sapiens
Homo neanderthalensis



Semi-permeable species boundary



Semi-permeable species boundary

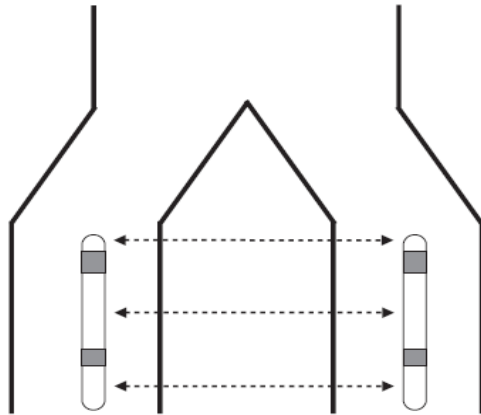


Genic species concept (Chung-I Wu, 2001)

- Reproductive isolation can be underlined by small part of the genome.
- Species as a coadapted gene complexes.

Speciation islands

Increase in F_{ST} a D_{XY}

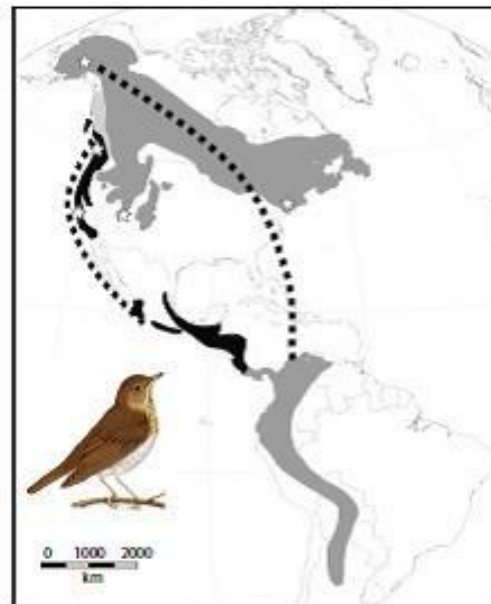


Speciation-with-gene-flow

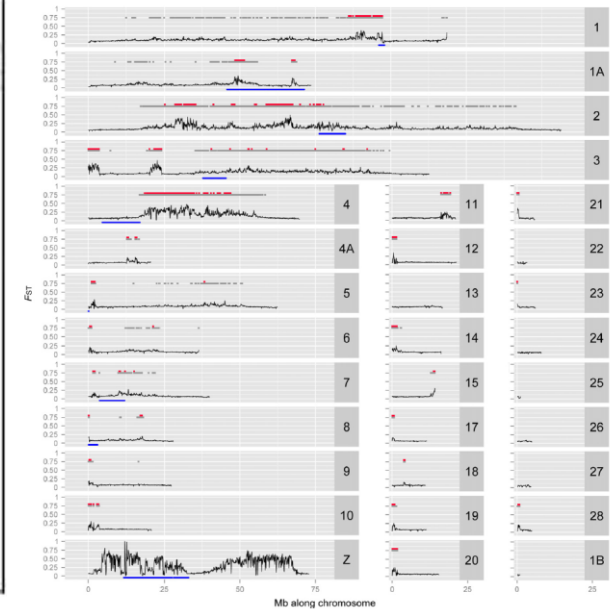
Migration homogenizes much of the genome



Swainson's Thrush

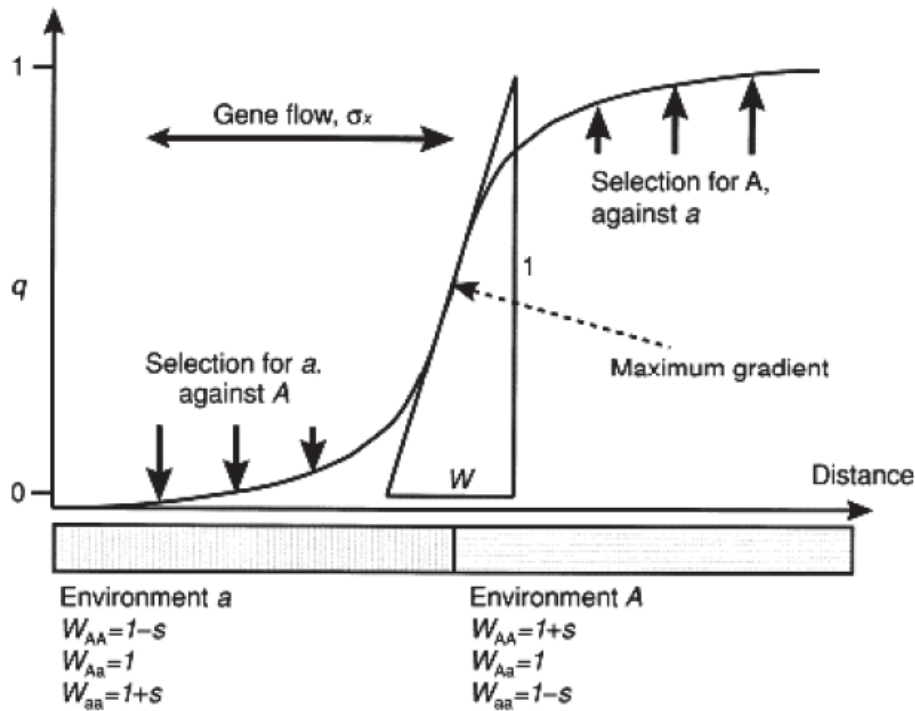


1880 K. E. DELMORE ET AL.



Delmore et al. 2015. Mol.Ecol.

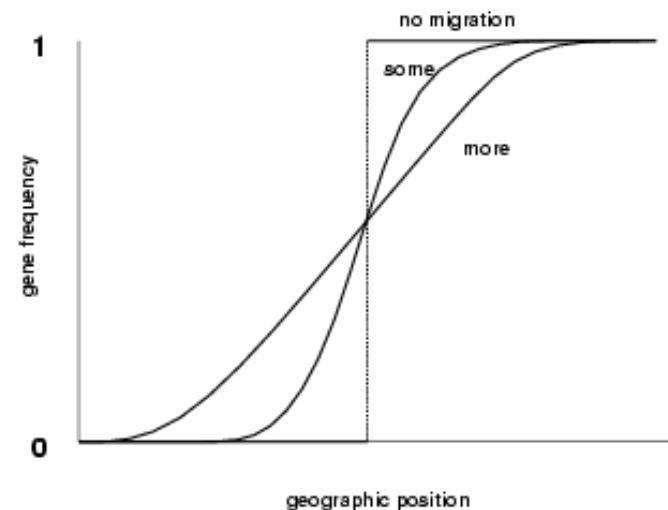
Geographic clines



- measure the movement of alleles across hybrid zones.
- Enable to compare levels of introgression for different markers.

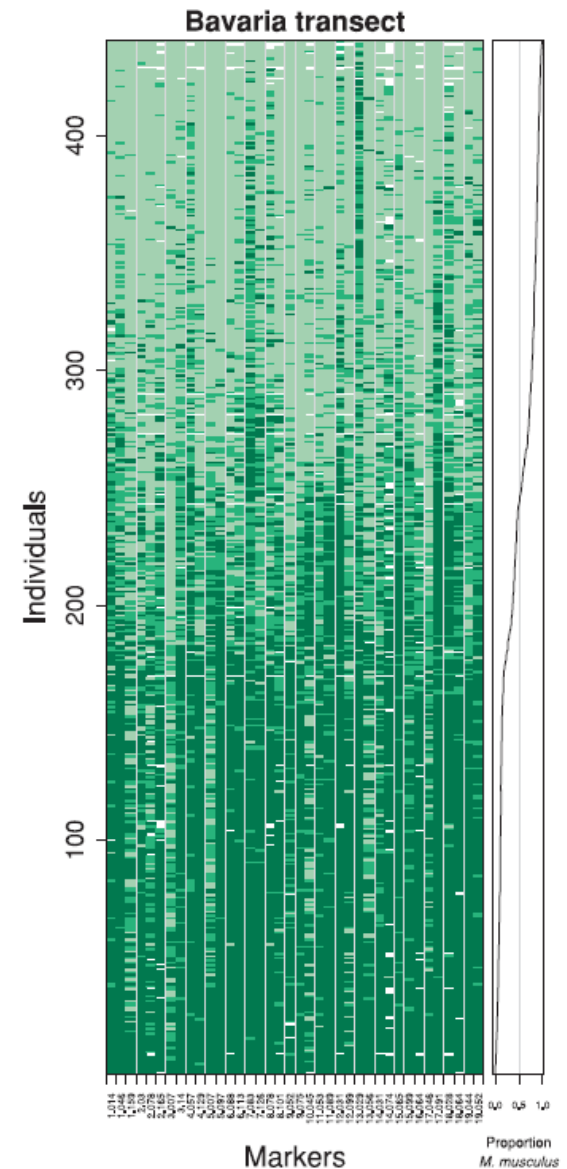
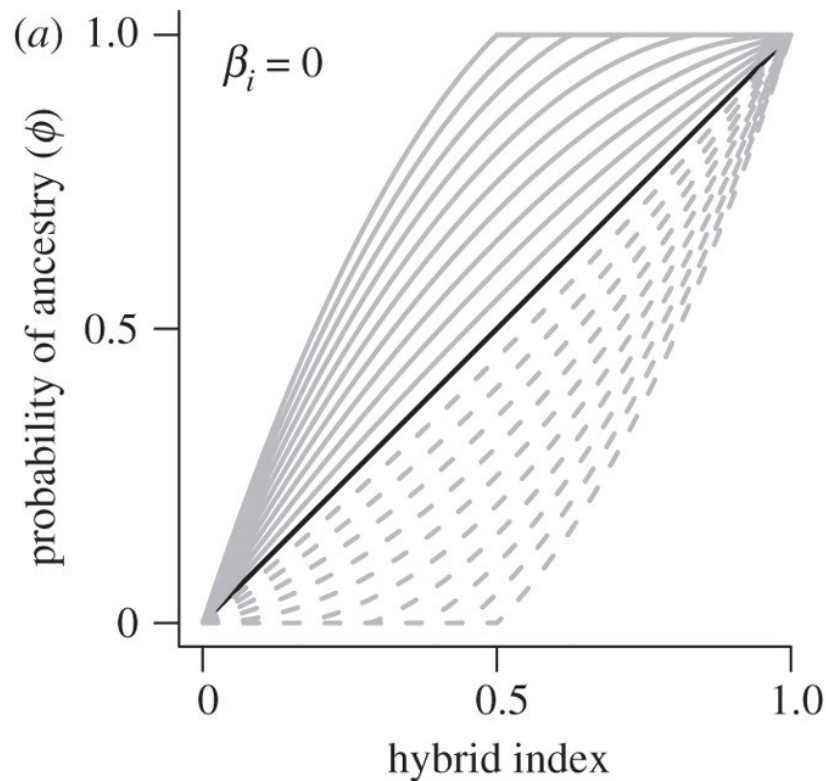


A cline



Genomic clines

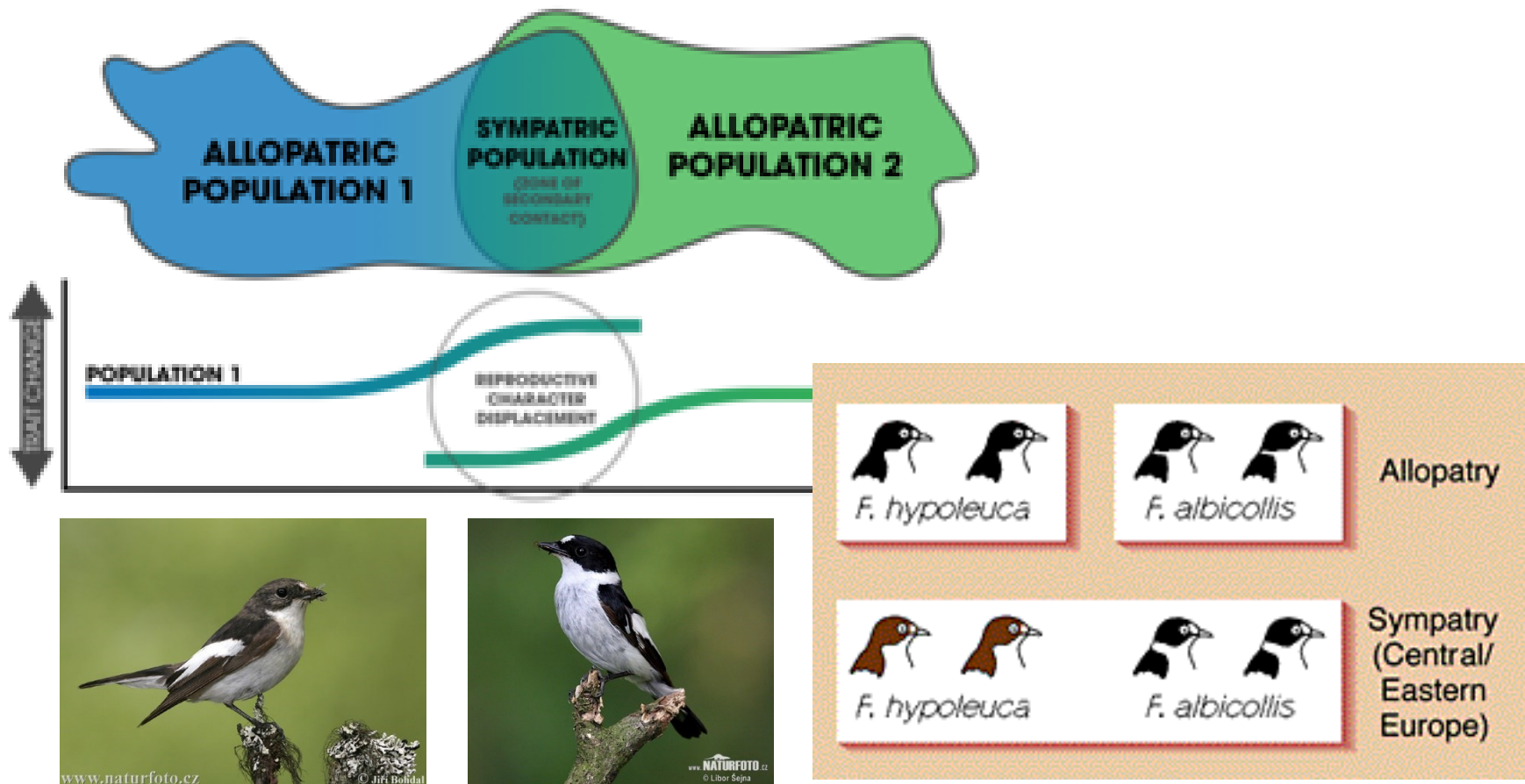
- Measure the movement of alleles into different genomic backgrounds.



How to complete speciation after secondary contact

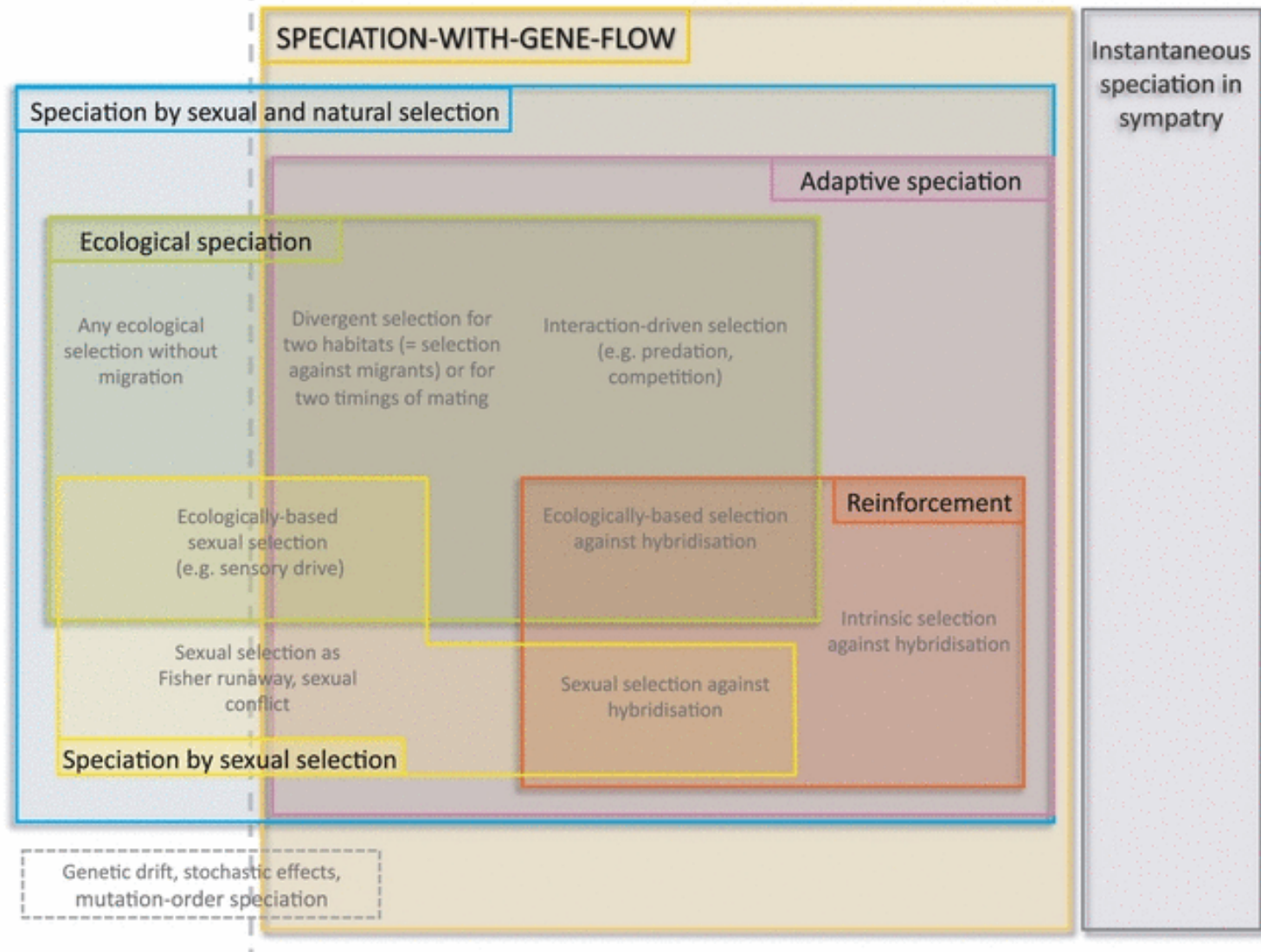
Reinforcement

- Origin of prezygotic barrier due to selection against hybridization

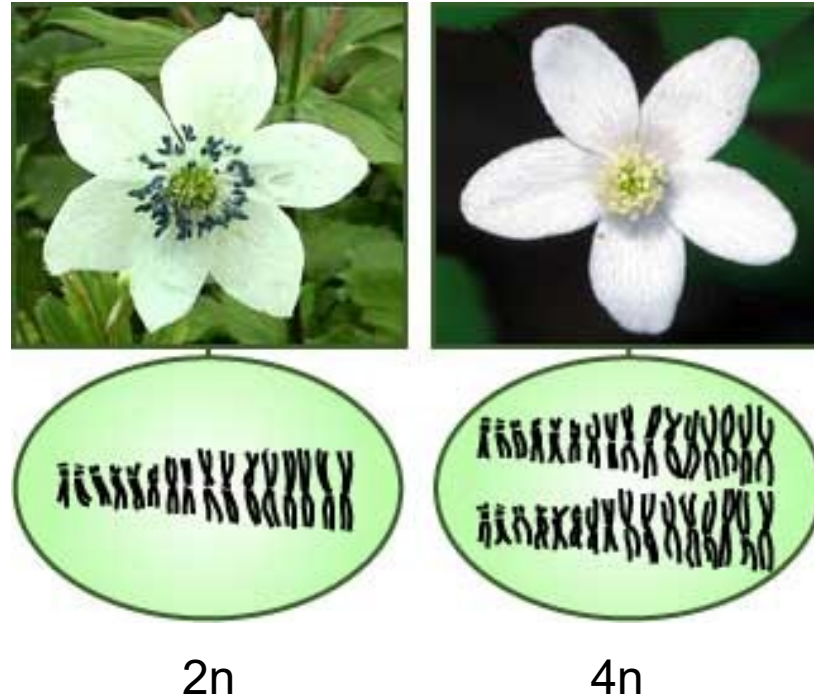


No contact (allopatry)

Geographical/Ecological contact (sympatry- parapatry; primary-secondary contact)



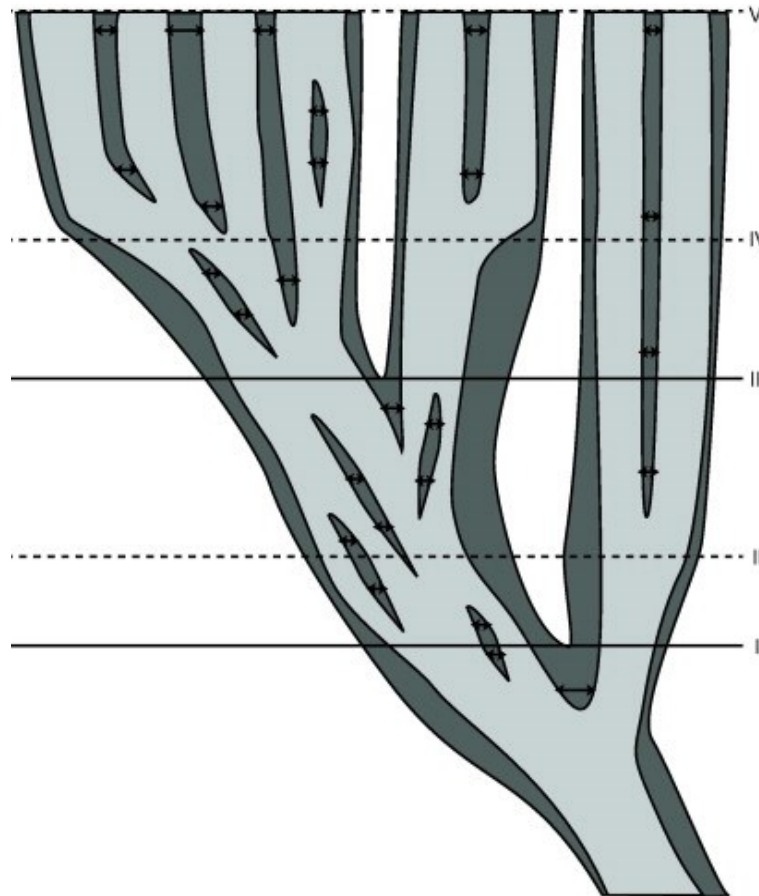
Instantaneous sympatric speciation by polyploidization



- Whole genome duplication will instantaneously create reproductive isolation between $2n$ and $4n$ plants as $3n$ hybrids will be sterile.

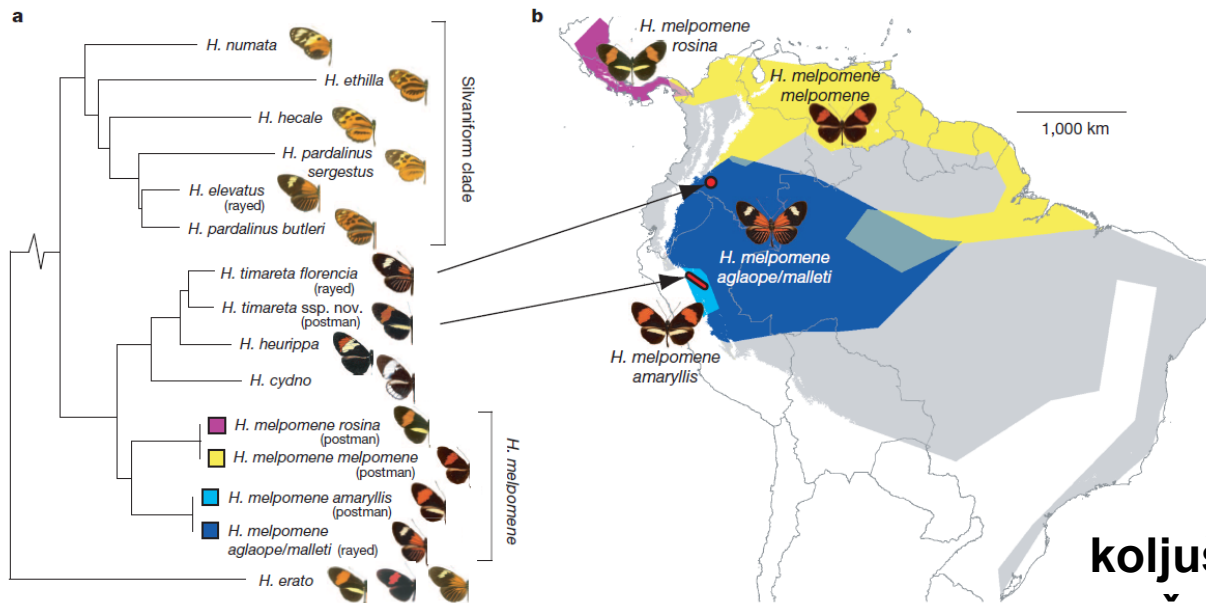
Divergence with gene flow

- Evolution of complete reproductive isolation is often a slow process.
- Many closely related species are not fully reproductively isolated.
- Species in various stages of origin



Evolutionary consequences of interspecific hybridization

Important source of genetic variation, speed up adaptive evolution
Parallel evolution



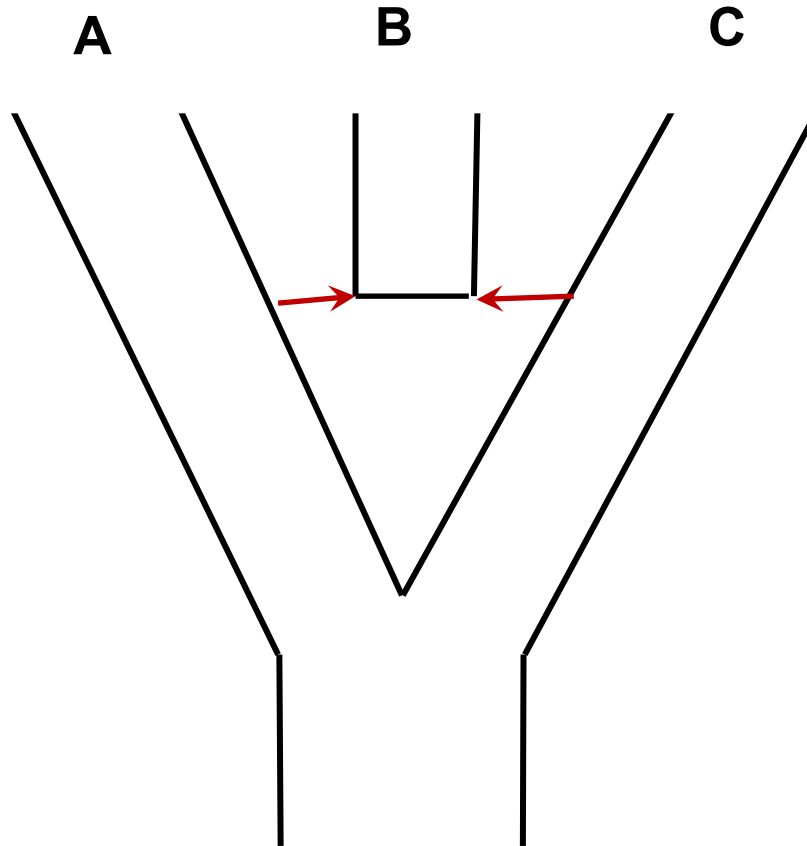
koljuška tříostná, forma
mořská (nahore) a říční (dole)



- Mixing up genetic variation



Hybridization can lead to the origin of new hybrid species

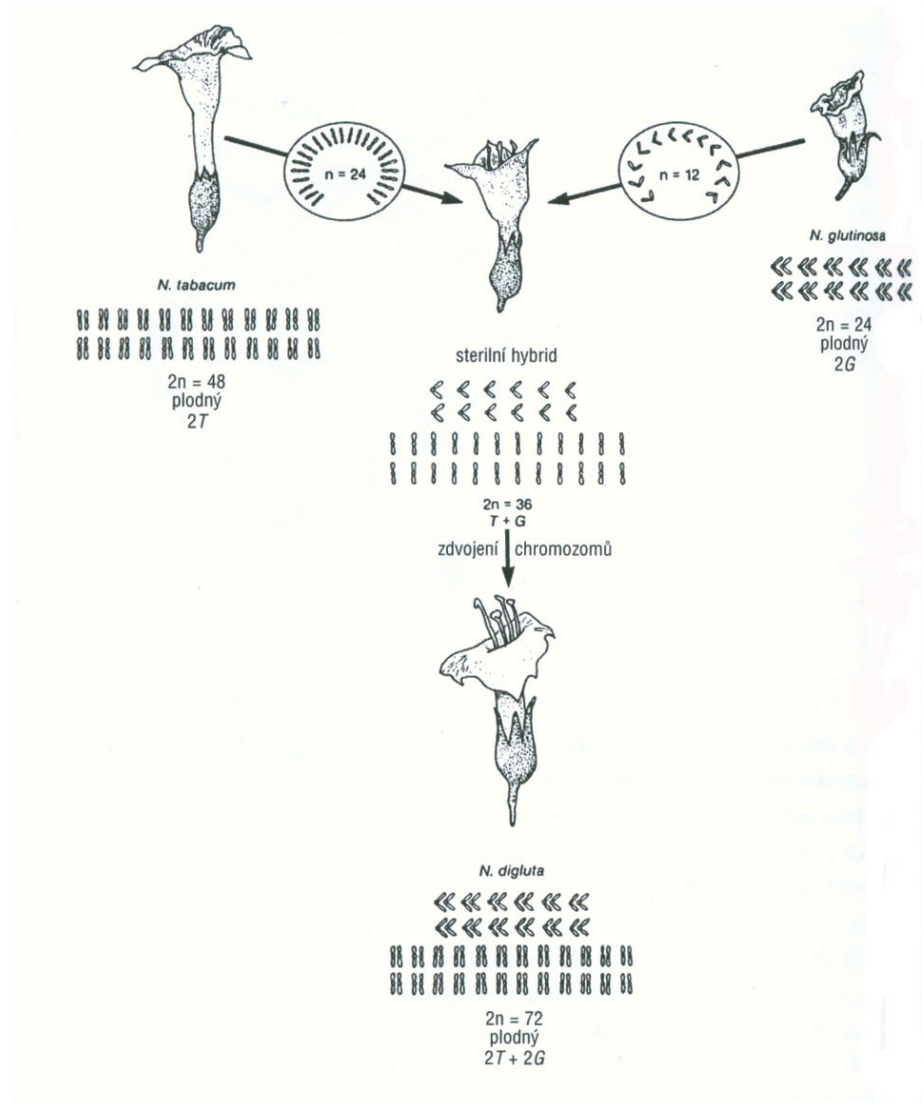


Polyploid hybrid speciation

- Whole genome duplication in hybrid.
- Will restore the fertility of hybrids.

At the same time will create reproductive barrier between hybrid and parental species.

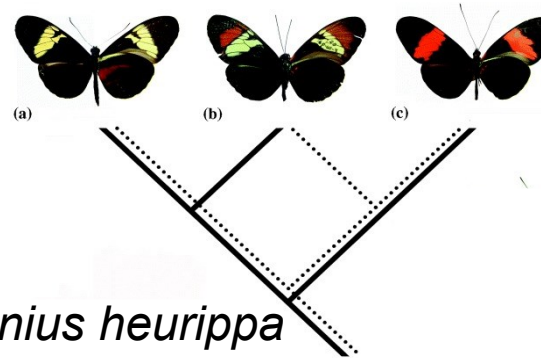
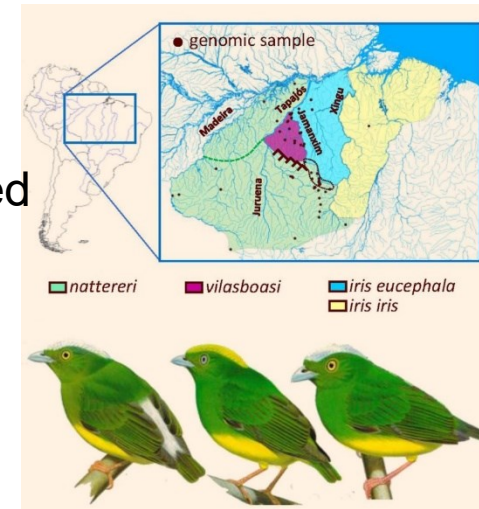
- ~ 15% of angiosperm plants originated by polyploid hybrid speciation.



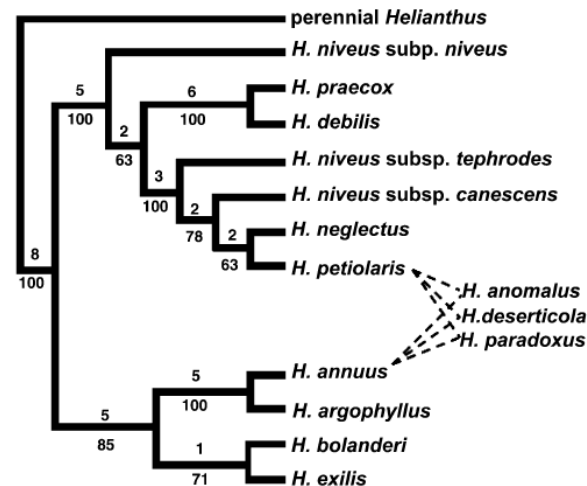
Homoploid hybrid speciation

- Without changes in ploidy.
- Reproductive isolation between parental species and hybrid species can arise if hybrids have different phenotype, occupy different ecological niche or are geographically isolated.

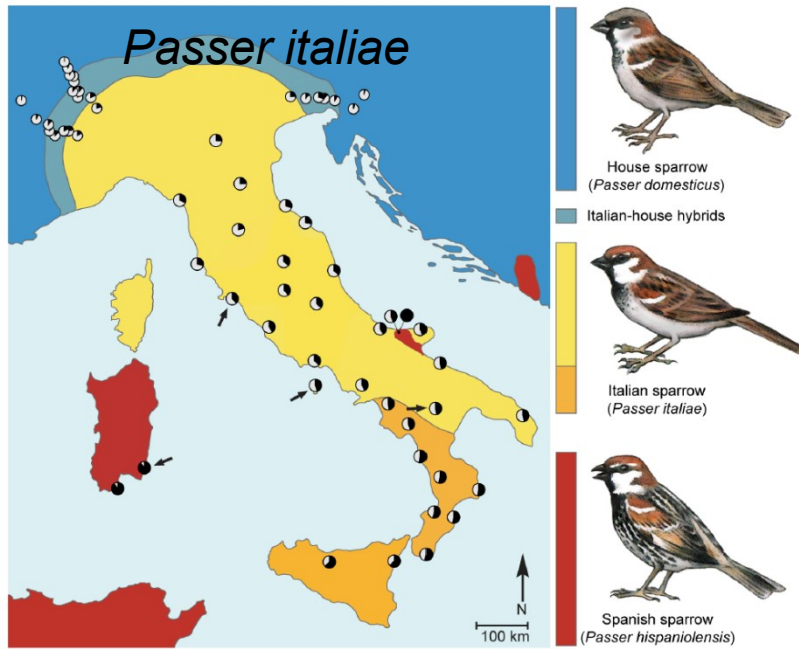
Golden crowned manakin



Heliconius heurippa



(sand dune)
(desert floor)
(salt marsh)

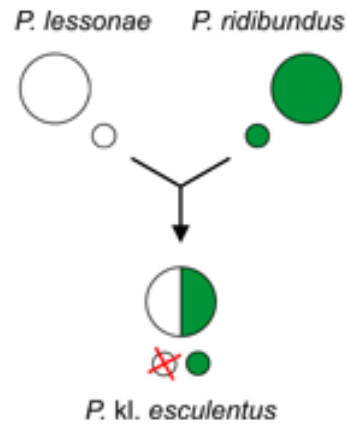


Hybridogenesis

Pelophylax ridibundus

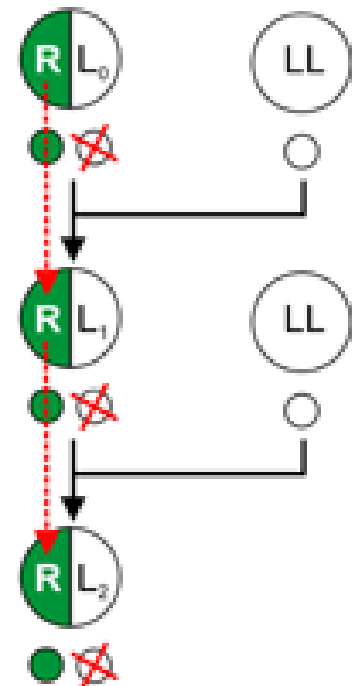


Pelophylax lessonae



Pelophylax esculentus

P. kl. esculentus *P. lessonae*



Intrinsic postzygotic isolation

Hybrid sterility



Oryza sativa
indica hybrid japonica

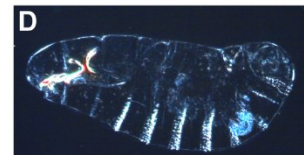
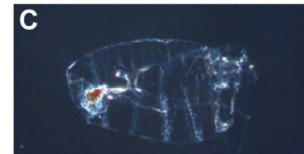


Hybrid inviability

♀ *D. melanogaster*
x ♂ *D. santomea*

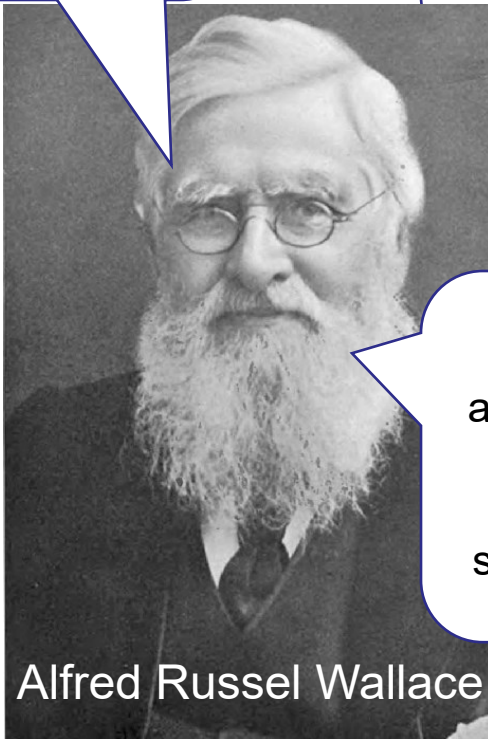


♀ *D. melanogaster* C(1)RM
x ♂ *D. santomea*



Origin of intrinsic postzygotic isolation

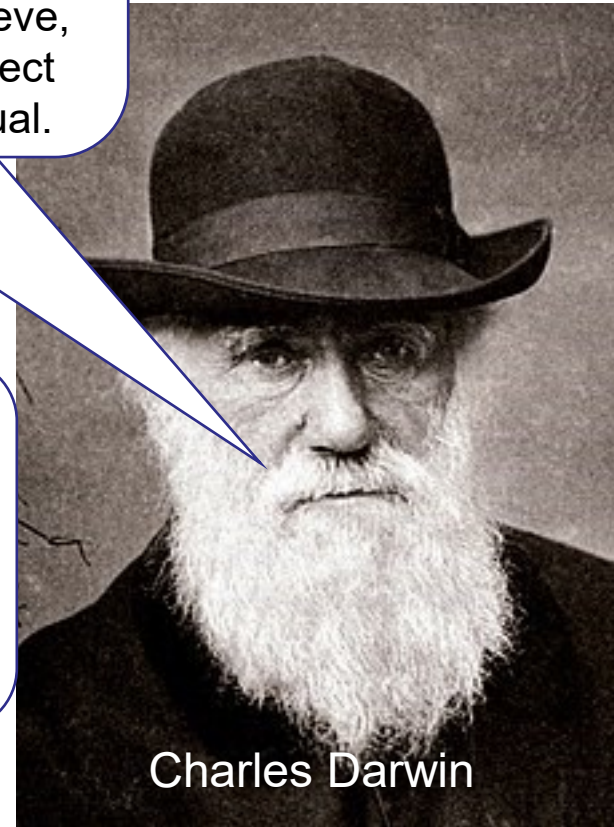
Could you imagine that hybrid sterility arose by natural selection?



Alfred Russel Wallace

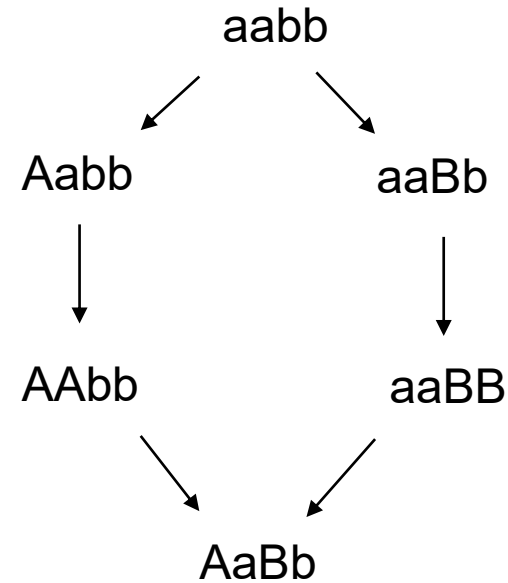
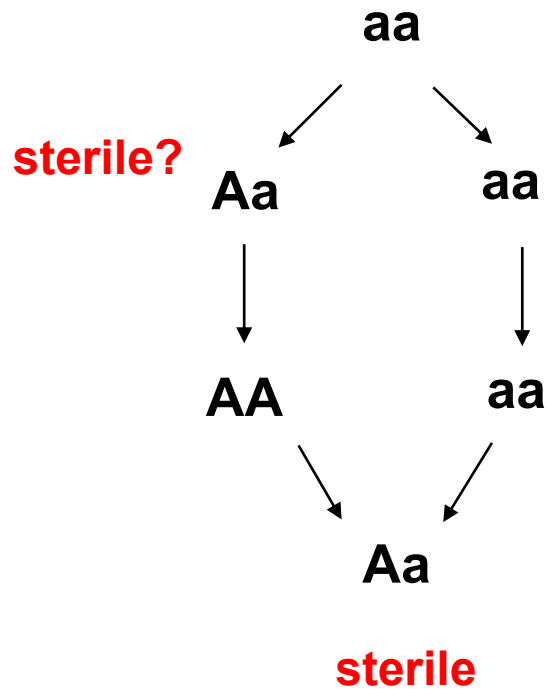
No man could have more earnestly wished for the success of natural selection in regard to sterility, than I did.... I always felt sure it could be worked out, but always failed in detail. The cause being as I believe, that natural selection cannot effect what is not good for the individual.

I am sorry you should have given yourself the trouble to answer my ideas on sterility... I now think there is about an even chance that natural selection may or not be able to accumulate sterility.

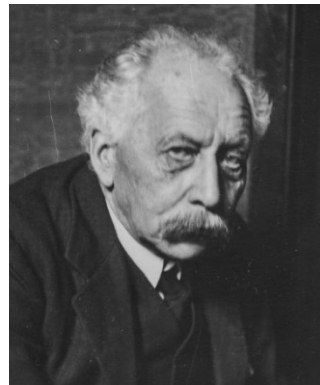


Charles Darwin

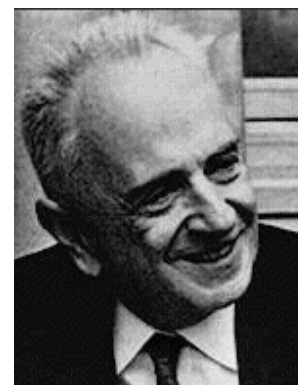
(Bateson)-Dobzhansky-Muller incompatibilities



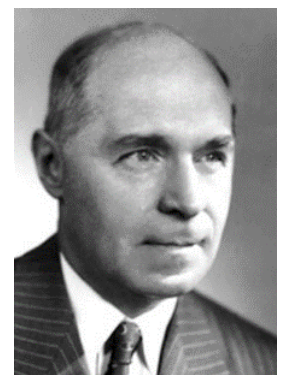
- Intrinsic postzygotic isolation does not arise adaptively, but as a consequence of species genetic divergence after isolation.



William
Bateson

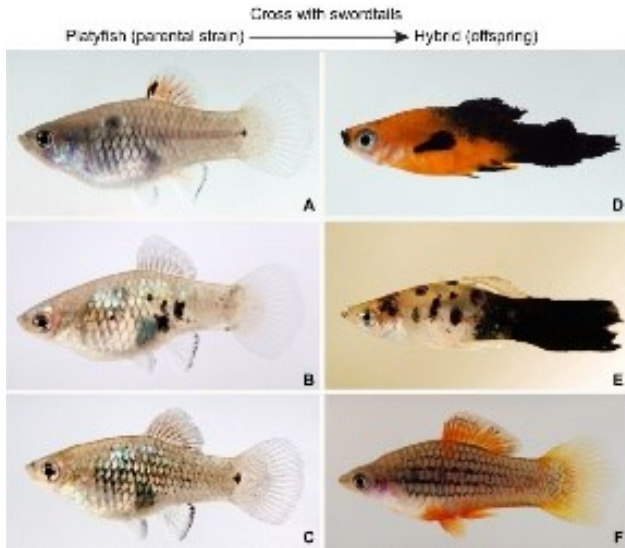


Theodosius
Dobzhansky



Herman J.
Muller

Speciation genes



Xiphophorus

Xmrk-2 (hybrid inviability)

Malign melanomas in hybrids



Drosophila

Overdrive (hybrid sterility)

OdsH (hybrid sterility)

Nup96 (hybrid inviability)

Hmr (hybrid inviability)

.....



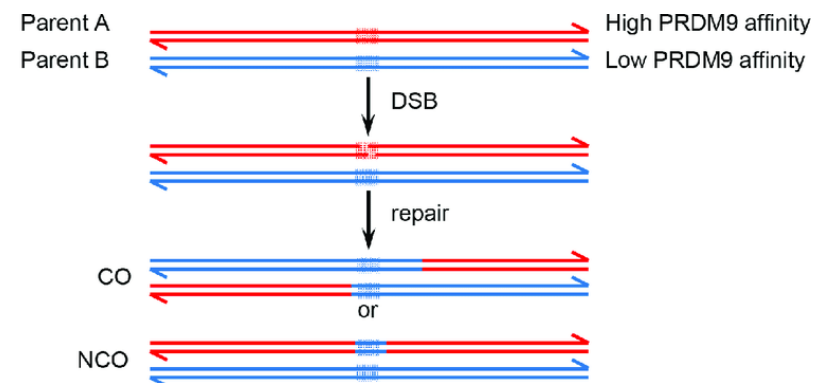
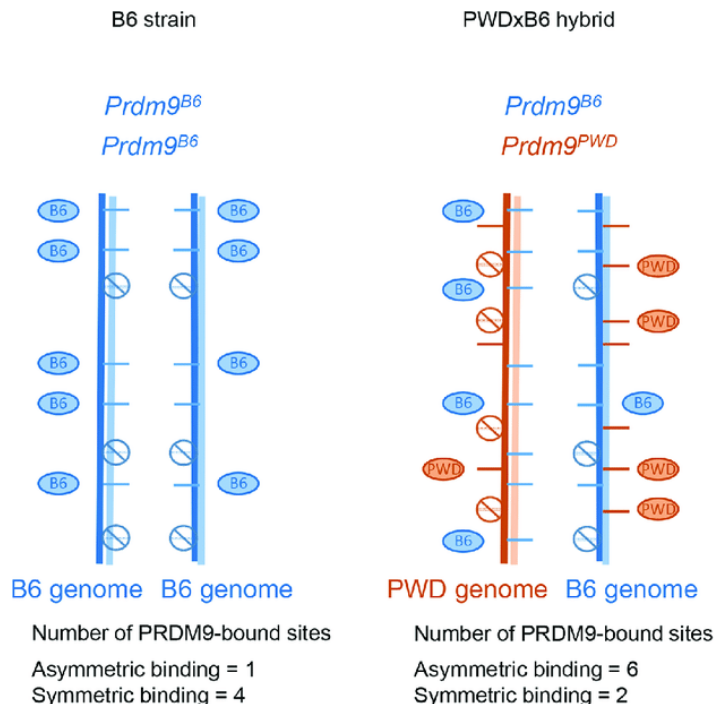
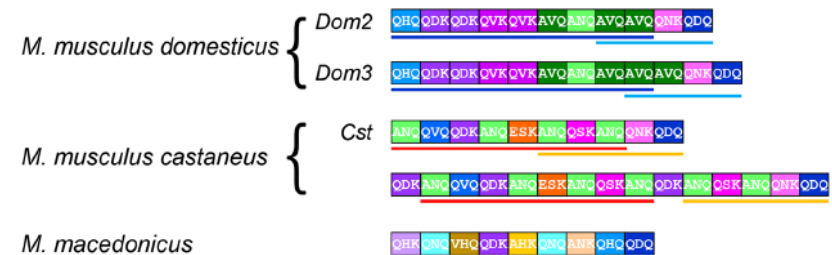
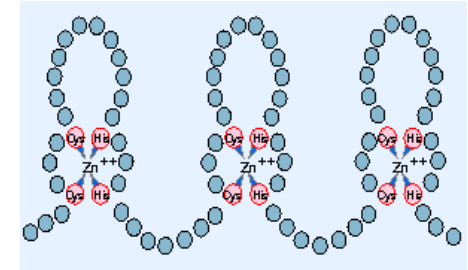
House mouse

Prmd9 (hybrid sterility)

- Fast molecular evolution usually due to positive selection

Prdm9

- Hybrid male sterility gene in house mouse.
- DNA binding protein (zinc-finger domain).
Determines the position of double strand breaks in meiosis (hotspots of recombination).
- Fast molecular evolution due to gene conversions.



Asymmetrical gene conversion.

Parental conflict, genomic imprinting and hybrid inviability in mammals

- Hybrid inviability arise relatively quickly in mammals compared to other vertebrates.



- Hybrid placental dysplasia in hybrids**

M. musculus x *M. spretus*

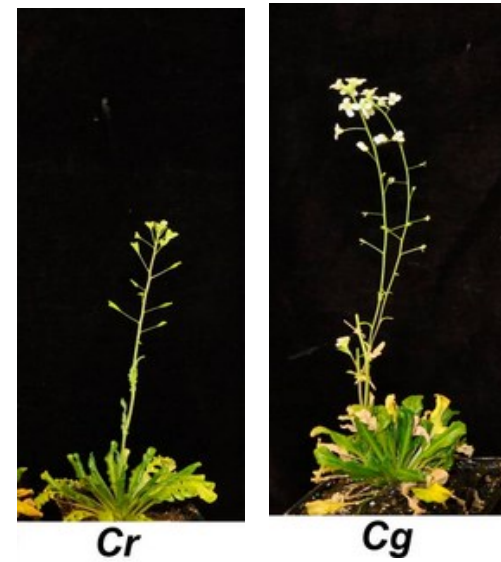
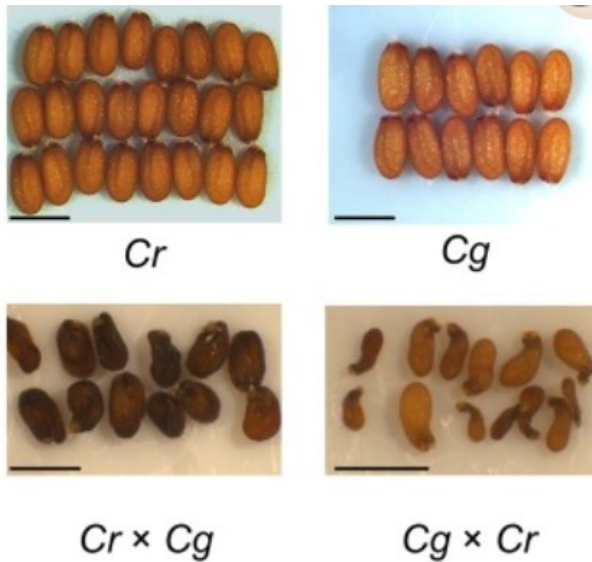
- smaller placenta

M. spretus x *M. musculus*

- larger placenta



Abnormal endosperm development in agiosperm hybrids

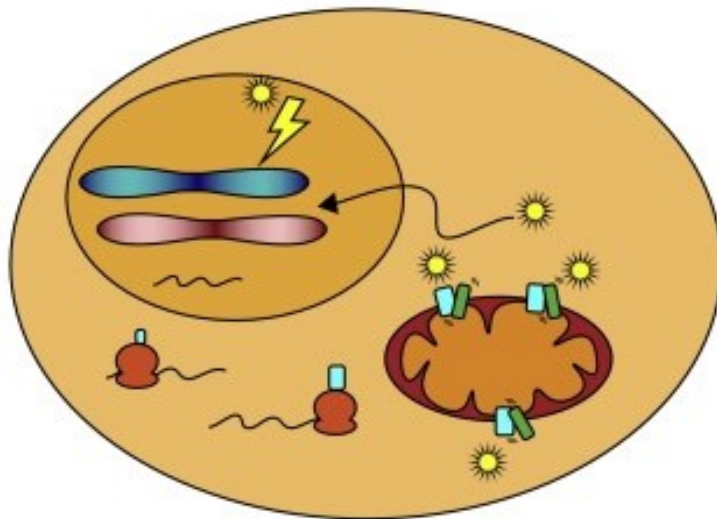


Capsella rubella a *C. grandiflora*

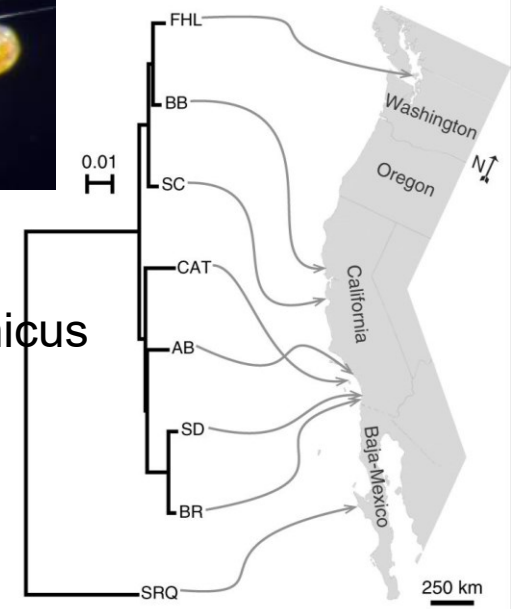
Reberning et al. 2015
Lafon-Placette and Kohler, 2016

Mito-nuclear inkompatibilities

- Proteins important for oxidative phosphorylation are partially coded in mtDNA and partially in nucleus.
- Relatively fast molecular evolution of mtDNA in mammals.
- Incompatibilities between mtDNA and nuclear genes can lead to suboptimal electron transfer and higher production of oxygen radicals.



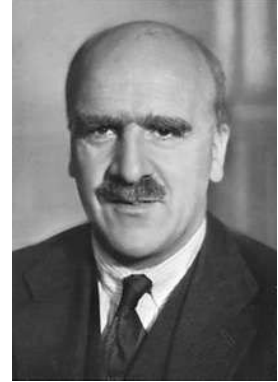
marine cope-pod
Tigriopus californicus



Two rules of speciation

Haldane's rule

When in the F1 offspring of two different species one sex is absent, rare, or sterile, that sex is the heterogametic sex (1922).



John B. S. Haldane

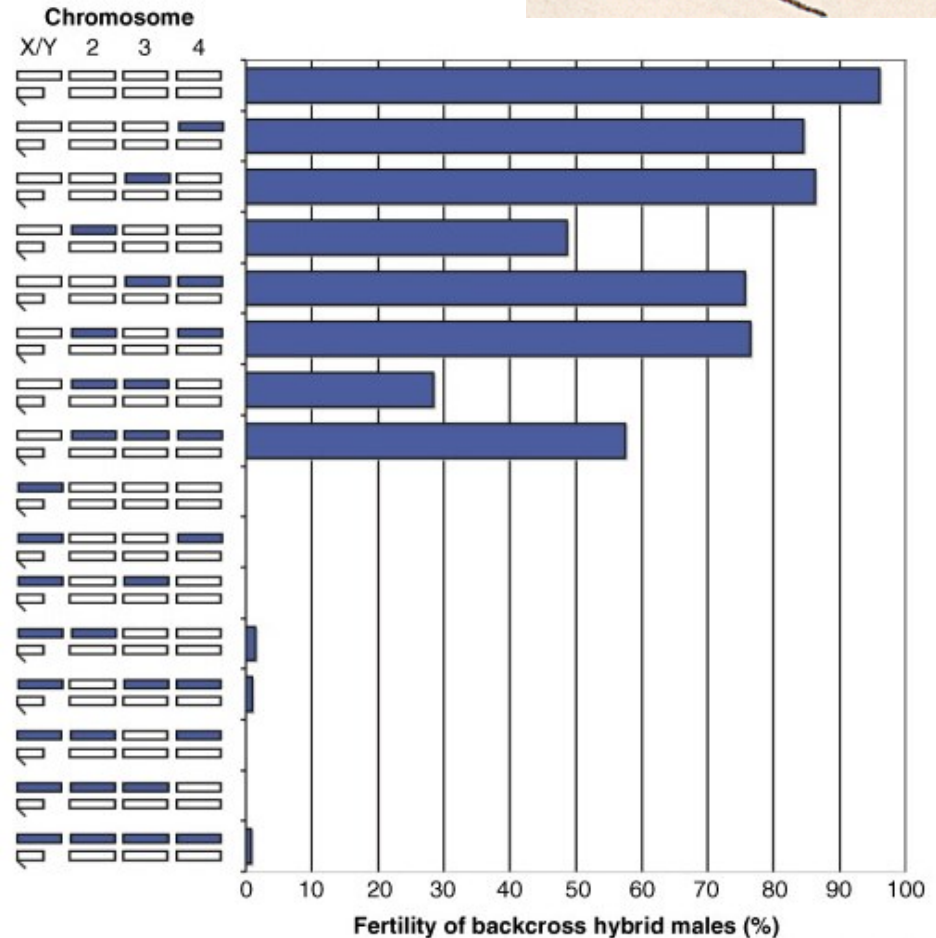
Two rules of speciation

Large X effect

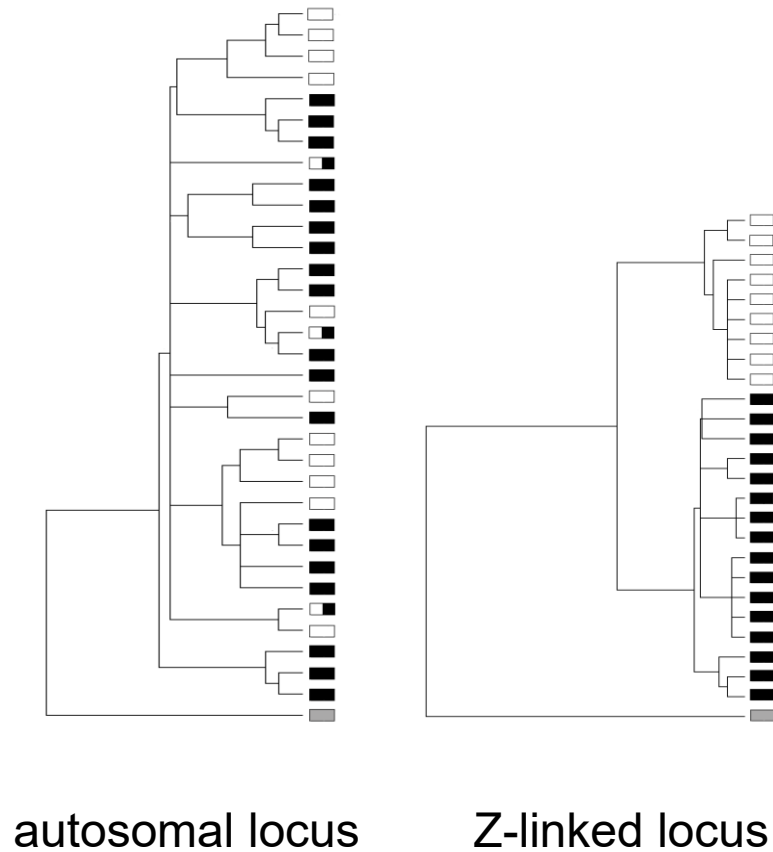
- X chromosome have a disproportionately large effect on intrinsic postzygotic isolation (sterility) compared to autosomes.



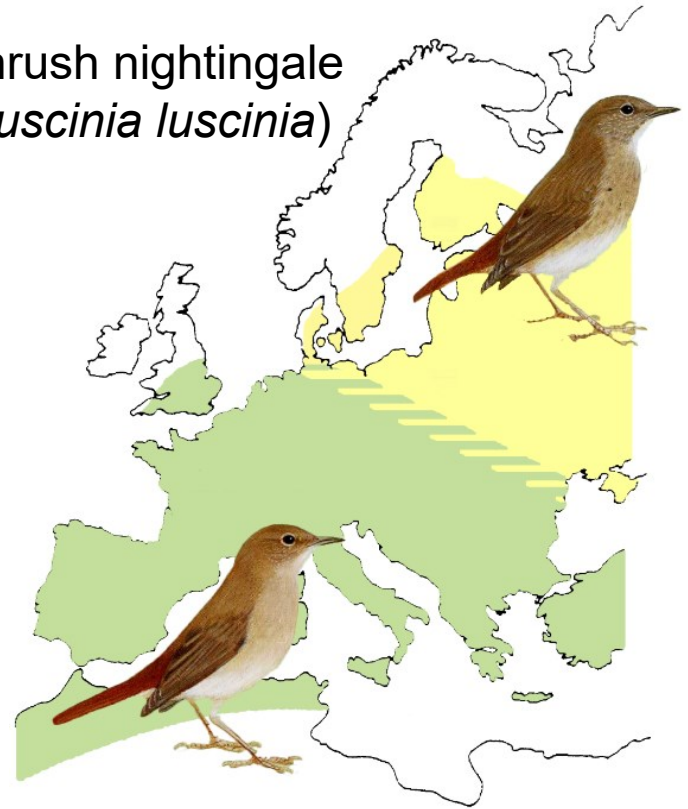
H. Allen Orr



The large Z effect



Thrush nightingale
(*Luscinia luscinia*)

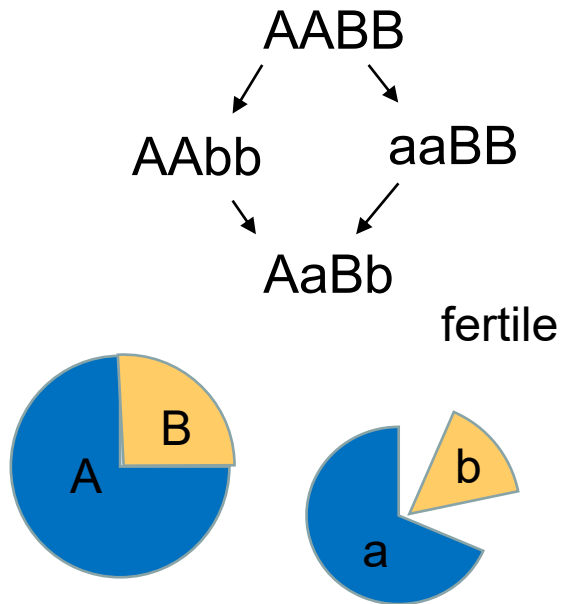


Common nightingale
(*Luscinia megarhynchos*)

Storchová et al. 2010

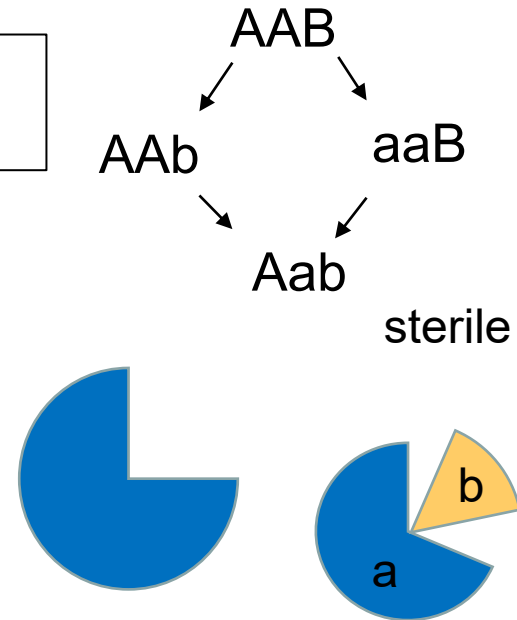
Dominance theory

Homogametic sex



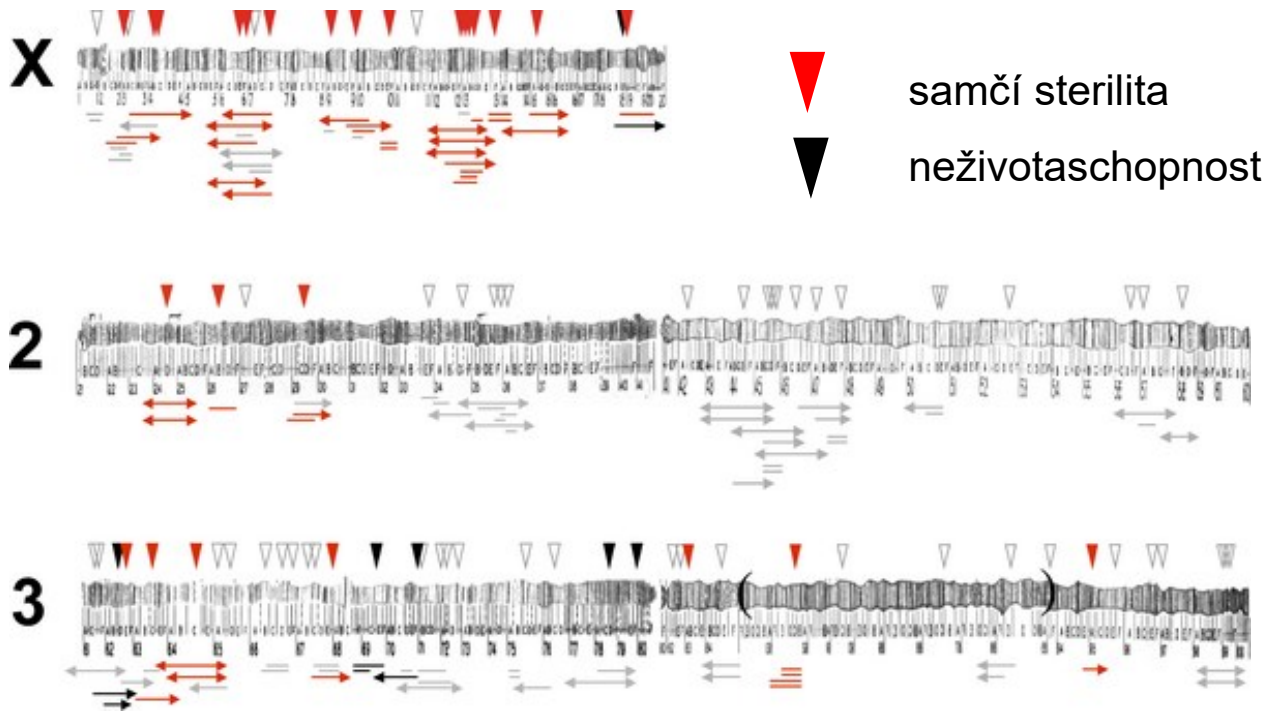
A,a... autosom
B,b....X chrom.

Heterogametic sex



Genes for hybrid sterility are more common on the X even if recessive autosomal incompatibilities are counted.

Homozygotní introgrese z *D. mauritiana* do *D. sechellia*

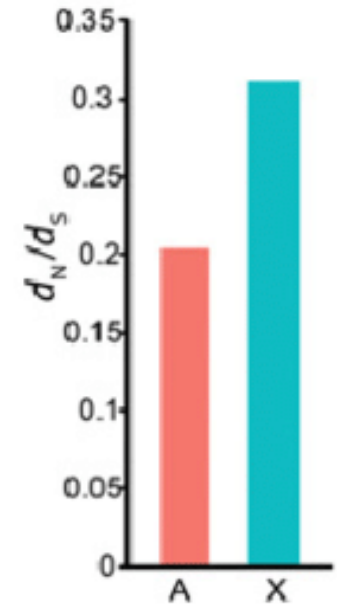


Masly and Presgraves, 2007

Another causes of the Haldane's rule and the large X/Z effect

Faster molecular evolution on the X/Z chromosomes

- i. Faster fixation of recessive advantageous mutations on the hemizygous X/Z.
- ii. Lower N_e and stronger genetic drift cause faster fixation of mildly disadvantageous mutations.



Another causes of the Haldane's rule and the large X/Z effect

Meiotic drive

Fast co-evolution between meiotic drivers on sex chromosomes and their suppressors. In hybrids can cause sterility.

- Hybrid male sterility.
- Caused by gene **Overdrive**. Normally causes meiotic drive on the X. Fast evolution due to recurrent positive selection (high dN/dS).

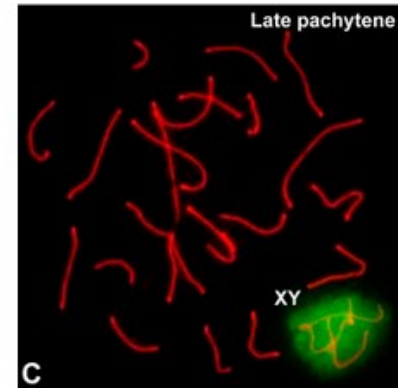


Drosophila pseudoobscura
Subspecies from Colombia
(Bogotá) and USA

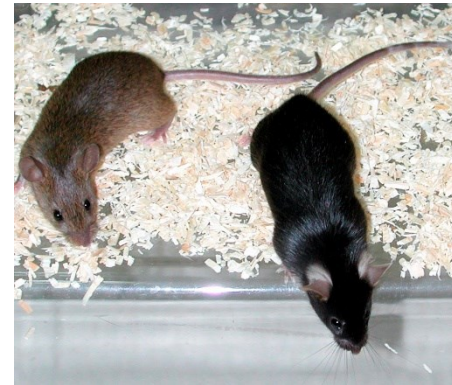
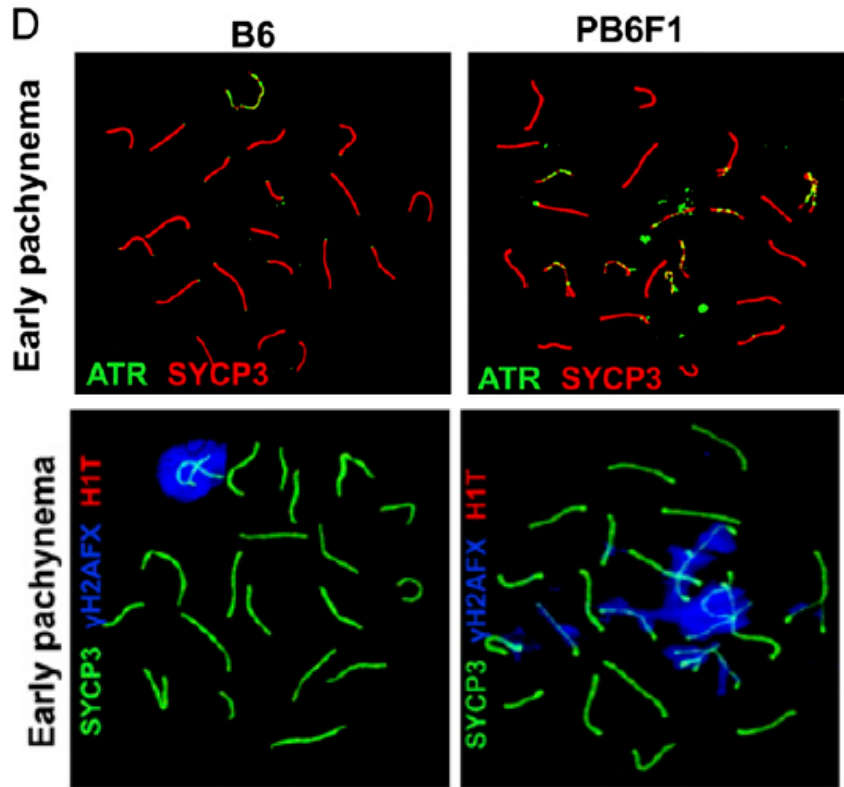
Phadnis and Orr, 2009

Another causes of the Haldane's rule and the large X/Z effect

Failure of sex chromosome inactivation during meiosis



sex body



PWD/Ph
(*M.m.musculus*)

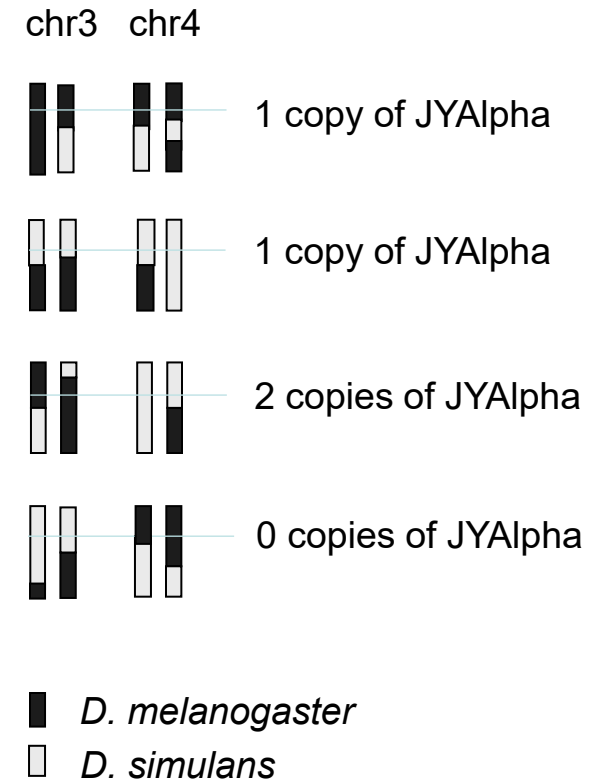
C57BL/6J
(*M.m.domesticus*)

- Hybrid males sterile.
- Asynapsis of homologous chromosomes in pachytene. In males at the same time failure of sex chromosome inactivation.

Bhattacharyya et al. (2013) PNAS

Translocation of genes as a cause of intrinsic postzygotic isolation

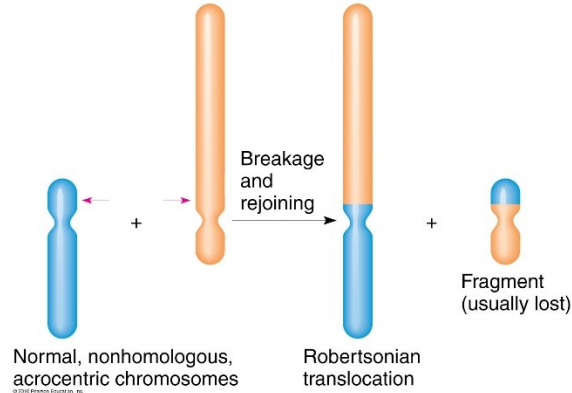
- JYAlpha gene
- On chr 3 in *Drosophila melanogaster*
- On chr 4 in *Drosophila simulans*
- Part of F2 hybrids lack JYAlpha -> sterility



Chromosomal speciation

- Changes in karyotypes (chromosomal rearrangements, fusions, fissions) cause problems with chromosome pairing and segregation during meiosis in hybrids (underdominance).
- Underdominant rearrangements are, however, unlikely to be fixed in the population unless they arise in very small populations inbreeding populations.
- Sterility can be caused by interaction of more rearrangements

Robertsonian translocations in mice

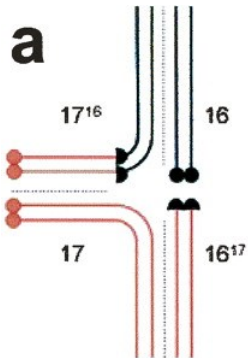


Mus musculus domesticus has more than 40 distinct local chromosomal races, characterized by about 100 types of Rb chromosomes with different arm combinations.

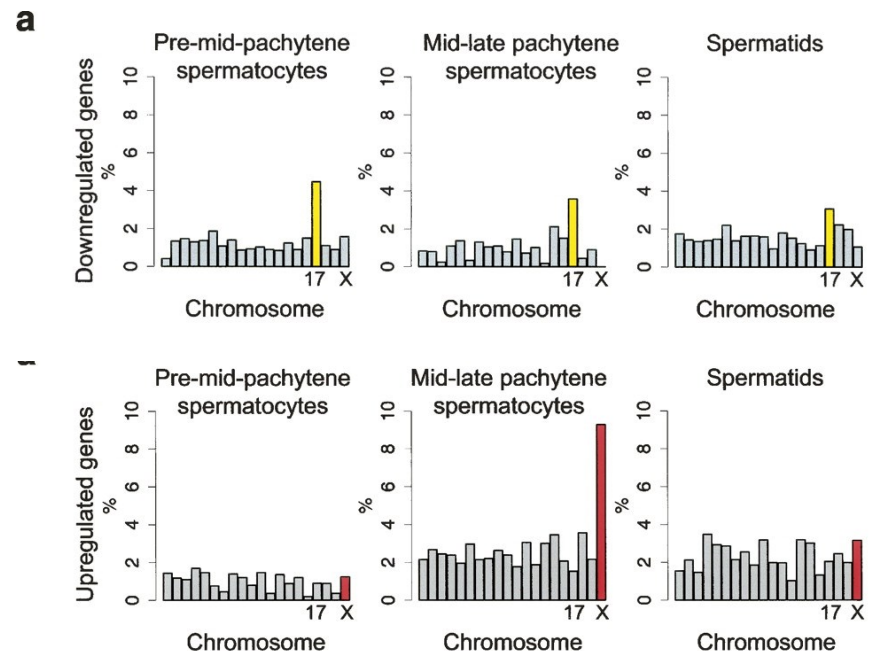
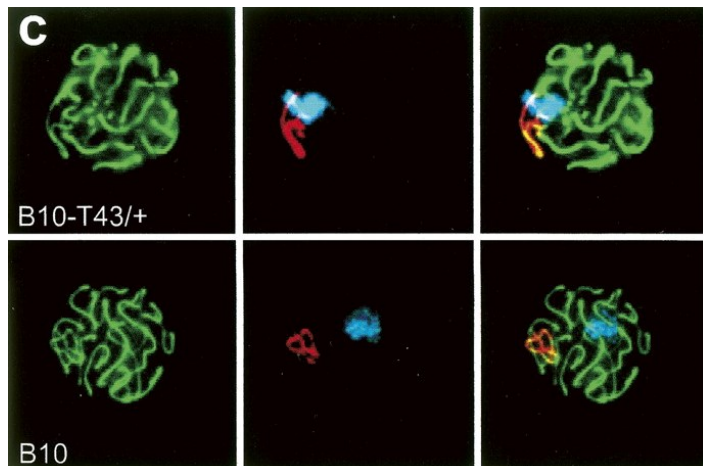
- Hybrids between species differing in single Robertsonian translocation may be fertile, but doubly heterozygous for two Robertsonian translocations involving the same chromosome are often sterile.

Chromosomal rearrangements and Haldane's rule

Many Robertsonian translocations, reciprocal autosome translocations and X-autosome translocations causes sterility of hybrid males, but not females (Haldane's rule).

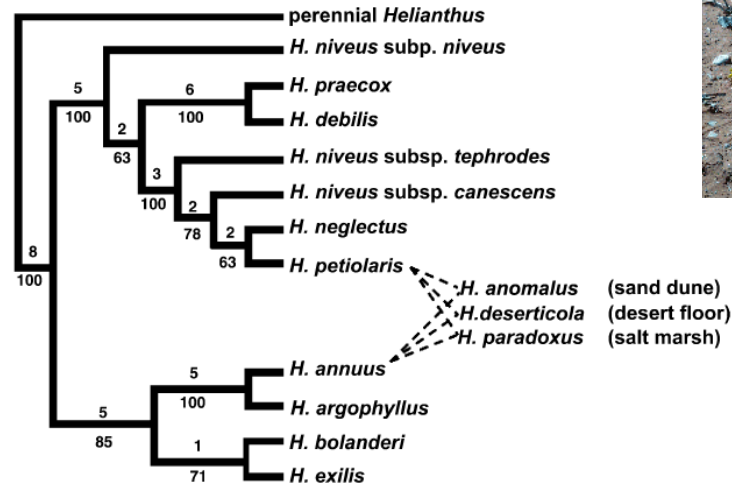
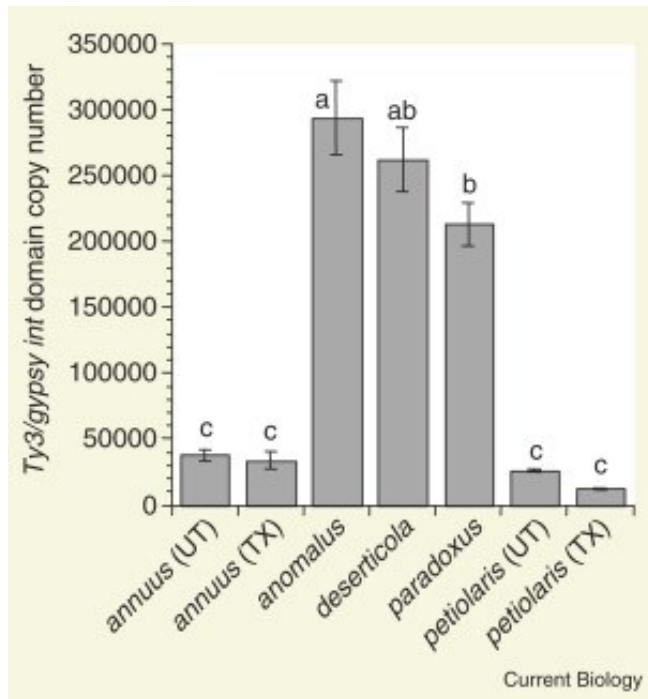


Sterile males heterozygous for chromosomal rearrangements show incomplete meiotic synapsis, meiotic silencing of unsynapsed chromosomes and disturbance of X chromosome inactivation.



Transposon derepression in interspecific hybrids

- Transposable elements are repressed by small RNAs (piRNAs) in the germline.
- Mismatch between paternally transmitted transposon and maternally inherited piRNAs can lead to transposon activation.



Diverse transposable elements are mobilized in hybrid dysgenesis in *Drosophila virilis*

(regulation of transposition)

DMITRI A. PETROV, JENNIFER L. SCHUTZMAN, DANIEL L. HARTL, AND ELENA R. LOZOVSKAYA

Department of Organismic and Evolutionary Biology, Harvard University, 16 Divinity Avenue, Cambridge, MA 02138

Communicated by Matthew Meselson, Harvard University, Cambridge, MA, May 25, 1995 (received for review April 4, 1995)

ABSTRACT We describe a system of hybrid dysgenesis in *Drosophila virilis* in which at least four unrelated transposable elements are all mobilized following a dysgenic cross. The data are largely consistent with the superposition of at least three different systems of hybrid dysgenesis, each repressing a different transposable element, which break down following the hybrid cross, possibly because they share a common pathway in the host. The data are also consistent with a mechanism in which mobilization of a single element triggers that of others, perhaps through chromosome breakage. The mobilization of multiple, unrelated elements in hybrid dysgenesis is reminiscent of McClintock's evidence [McClintock, B. (1955) *Brookhaven Symp. Biol.* 8, 58–74] for simultaneous mobilization of different transposable elements in maize.



viral element not detectable in either parent species. These results, taken with the observation of deficient methylation and *de novo* chromosome change in other mammalian hybrids, indicate that the failure of DNA methylation and subsequent mobile-element activity in hybrids could facilitate rapid karyotypic evolution.

Undermethylation associated with retroelement activation and chromosome remodelling in an interspecific mammalian hybrid

Rachel J. Waugh O'Neill^{††}, Michael J. O'Neill[‡], & Jennifer A. Marshall Graves^{*}

^{*} Department of Genetics and Human Variation, La Trobe University, Bundoora, Victoria 3083, Australia

[‡] Department of Molecular Biology, Princeton University, Princeton, New Jersey 08544, USA

Genetic models^{1,2} predict that genomic rearrangement in hybrids can facilitate reproductive isolation and the formation of new species by preventing gene flow between the parent species and hybrid (sunflowers are an example³). The mechanism underlying hybridization-induced chromosome remodelling is as yet unknown, although mobile element activity has been shown to be involved in DNA rearrangement in some dysgenic *Drosophila* hybrids^{4,5}. It has been proposed that DNA methylation evolved as a means of repressing the movement of mobile elements (the host defence model^{6,7}). If such a protective mechanism were to fail, mobile elements could be activated, and could cause major and rapid genome alterations^{8,9}. Here we demonstrate the occurrence of genome-wide undermethylation, retroviral element amplification and chromosome remodelling in an interspecific mammalian hybrid (*Macropus eugenii* × *Wallabia bicolor*). Atypically extended centromeres of *Macropus eugenii* derived autosomes in the hybrid were composed primarily of an unmethylated, amplified retro-

^{††} Present address: Department of Ecology and Evolutionary Biology, Princeton University, Princeton, New Jersey 08544, USA.



***Drosophila buzzatii* and *D. koepferae* coexist in the arid zones of Bolivia and the Argentine Northwest.**

- Mobilization of TEs has been described in interspecific hybrids.
- TE deregulation very likely due to fast divergence in piRNA pathway proteins and piRNA (signs of positive selection, differential expression between species).



Article

***Drosophila* Interspecific Hybridization Causes a Deregulation of the piRNA Pathway Genes**

Víctor Gámez-Visairas ^{1,†}, Valèria Romero-Soriano ^{1,2,†}, Joan Martí-Carreras ^{1,3} , Eila Segarra-Carrillo ¹ and Maria Pilar García Guerreiro ^{1,*}

¹ Grup de Genòmica, Bioinformàtica i Biologia Evolutiva, Departament de Genètica i Microbiologia (Edifici C), Universitat Autònoma de Barcelona, 08193 Bellaterra, Spain; victor.gamez@uab.es (V.G.-V.); V.Romero-Soriano@liverpool.ac.uk (V.R.-S.); joan.marti@kuleuven.be (J.M.-C.); eilasegarracarrillo@gmail.com (E.S.-C.)

² Institute of Integrative Biology, University of Liverpool, Liverpool L69 7ZB, UK

³ Laboratory of Clinical Virology, Department of Microbiology, Immunology and Transplantation, Rega Institute, KU Leuven, B3000 Leuven, Belgium

* Correspondence: mariapilar.garcia.guerreiro@uab.es; Tel.: +34-935814703

† Co-first author, these authors contributed equally to this work.

Received: 14 January 2020; Accepted: 14 February 2020; Published: 19 February 2020

