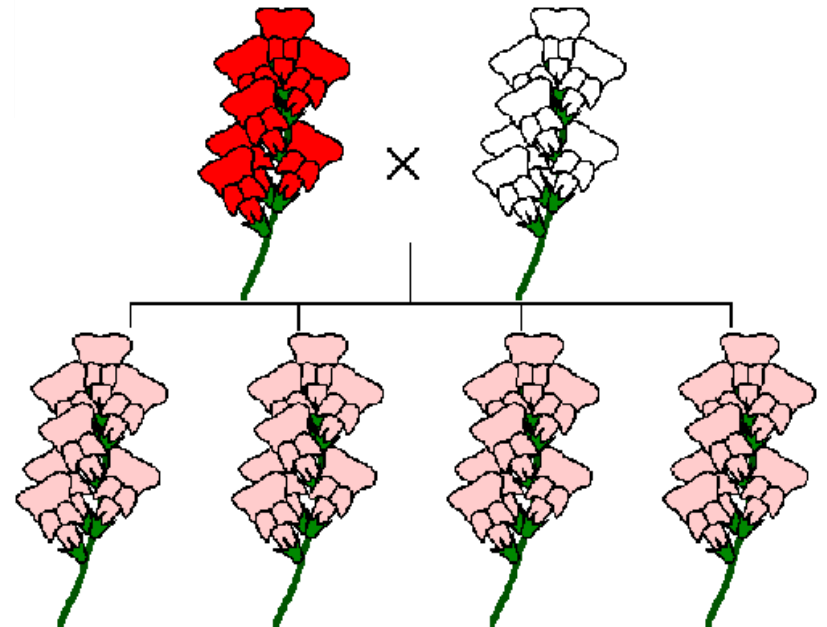
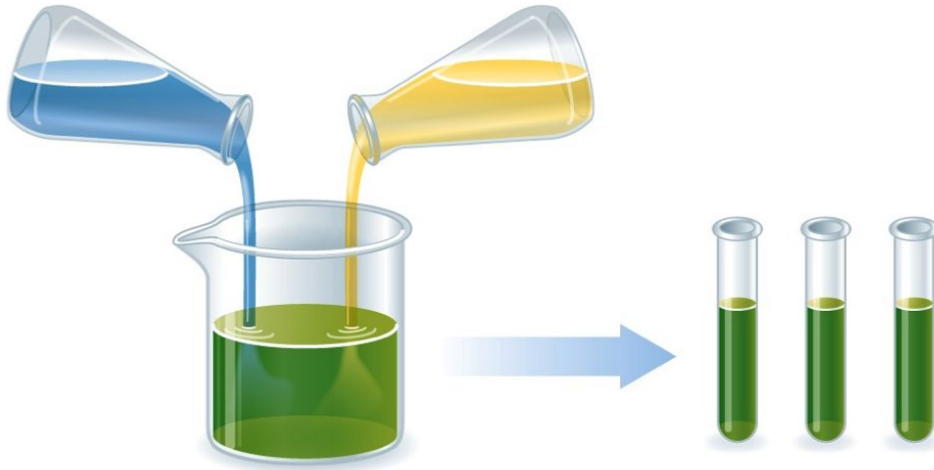


Inheritance

Inheritance



Blending inheritance



PŘEDSTAVTE SI BĚLOCHA, KTERÝ ZTROSKOTAL NA OSTROVĚ
OBYDLENÉM ČERNOCHY...

Tak mládenci, nechte
mě si vybrat některé
z vašich manželek.
Koneckonců
jsem Brit...

Chci také nějaké
dobrovolníky,
aby odnesli moje
zavazadla.

Bláznivý
běloch

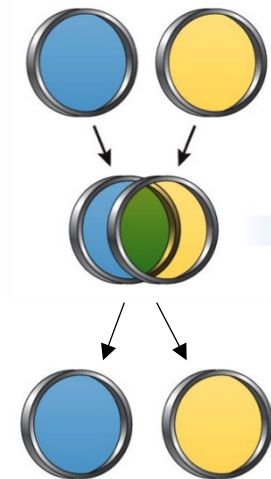
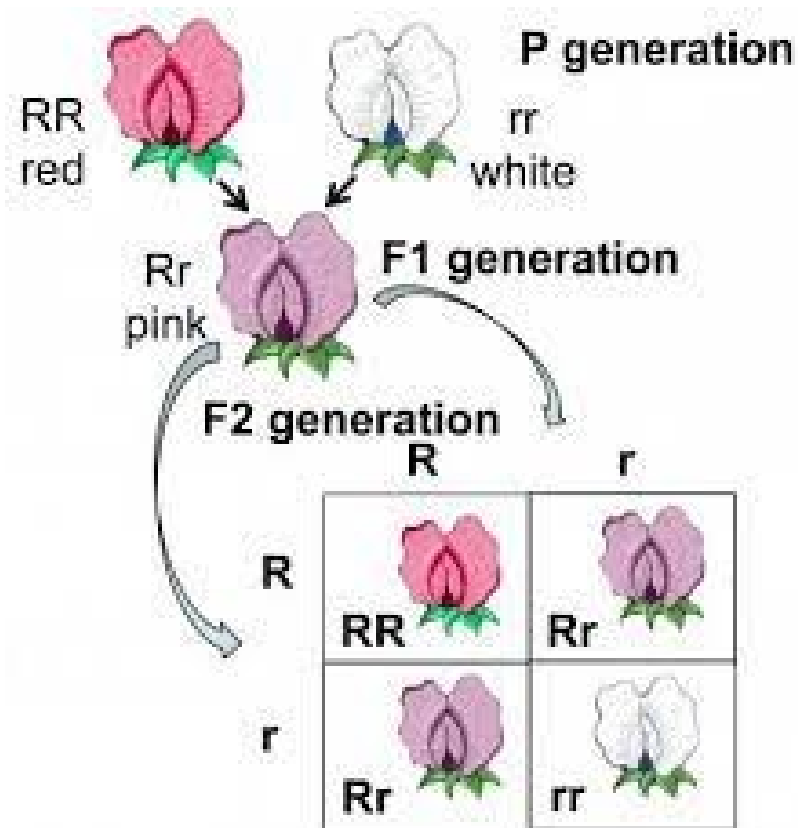
MĚL BY SPOUSTU MANŽELEK A NADPRŮMĚRNÝ POČET DĚTÍ...

Fleeming Jenkin
criticizes Darwin's
theory of evolution.

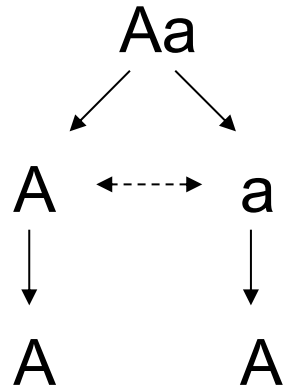
...ALE UVĚŘIL BY NĚKDO TOMU, ŽE NA
CELÉM OSTROVĚ POSTUPNĚ VZNIKNE
BÍLÁ NEBO I JEN ŽLUTÁ POPULACE?

Mendel's theory of inheritance (1866)

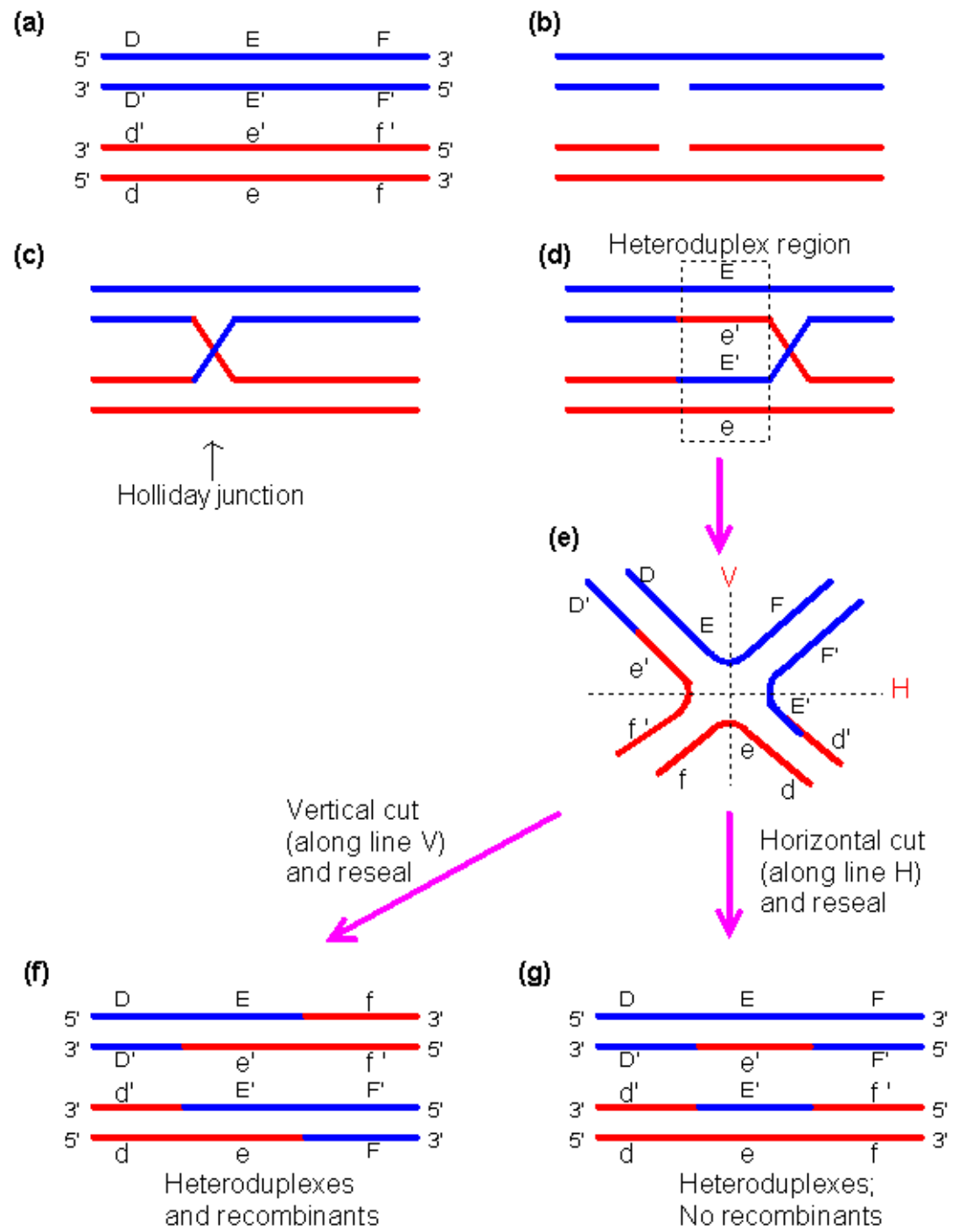
Law of segregation



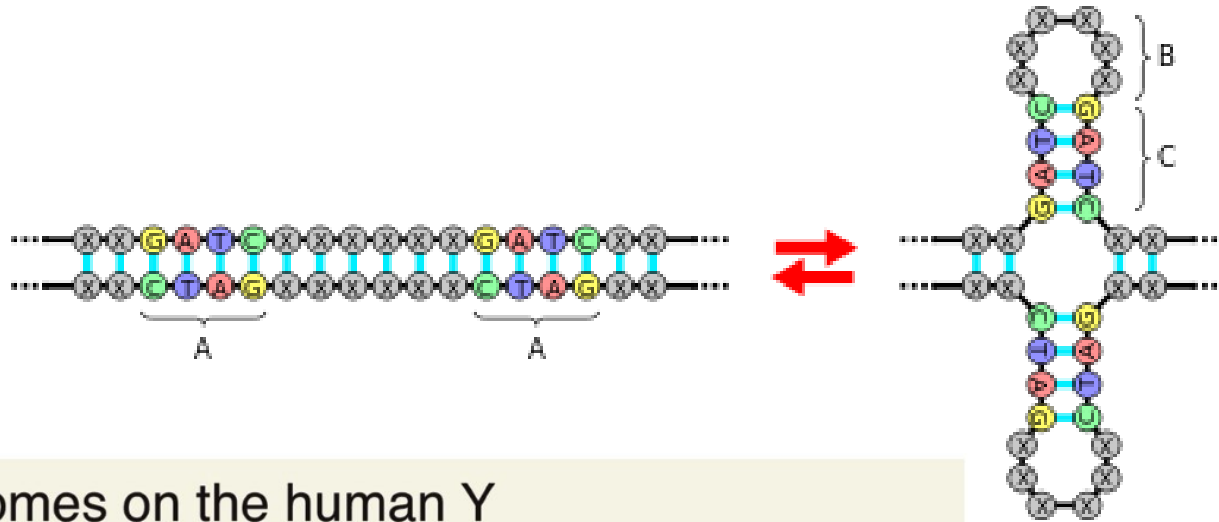
Gene conversion



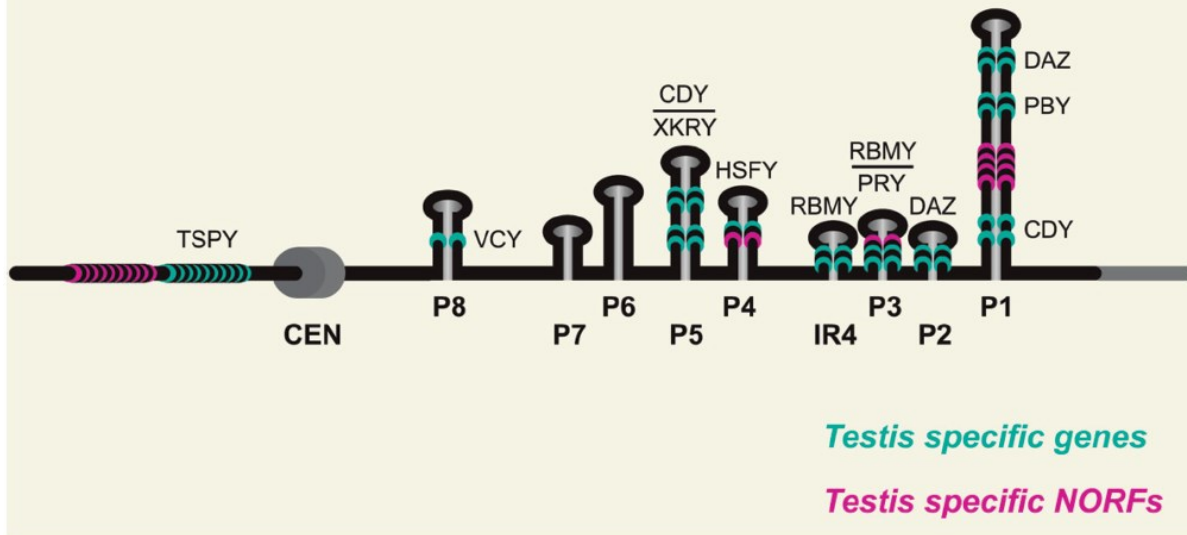
- Gene conversion during meiotic recombination.



Gene conversion between paralogous sequences

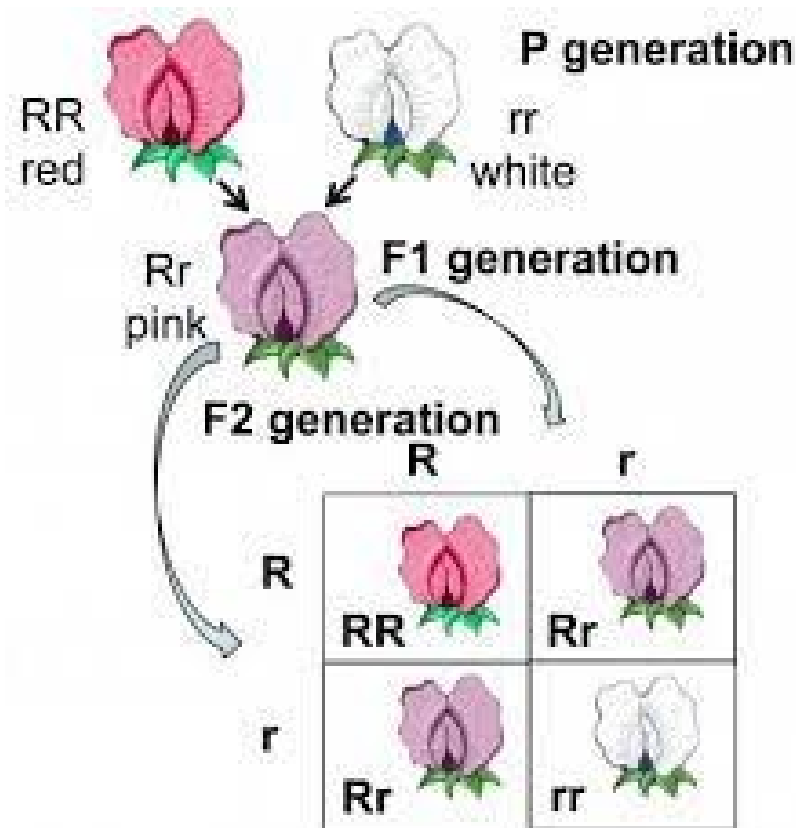


Palindromes on the human Y



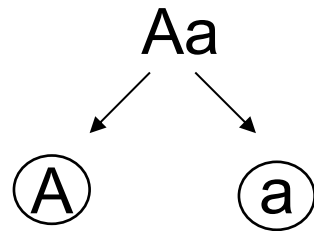
Mendel's theory of inheritance (1866)

Law of segregation



1:2:1

Meiotic drive

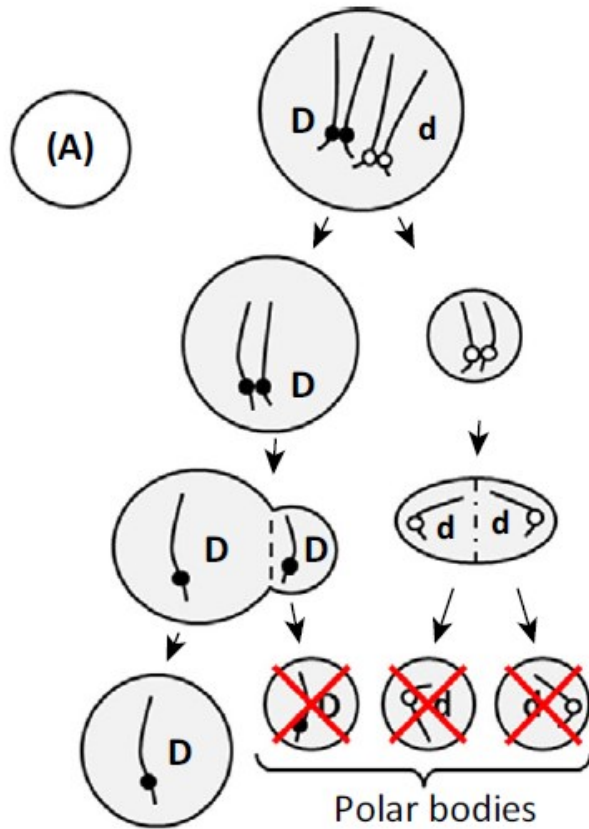


~~1 : 1~~

10 : 1
1 : 10

Female meiotic drive

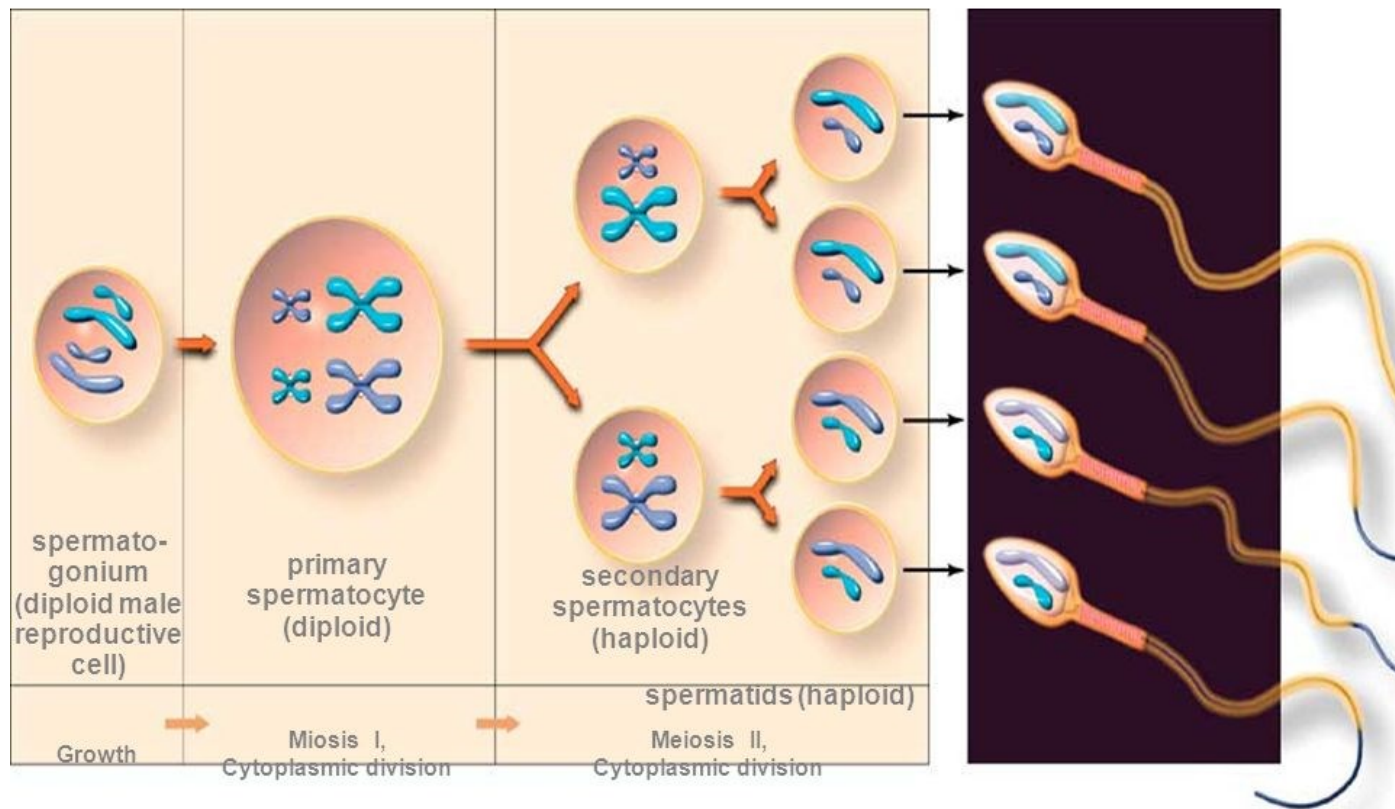
Oogenesis



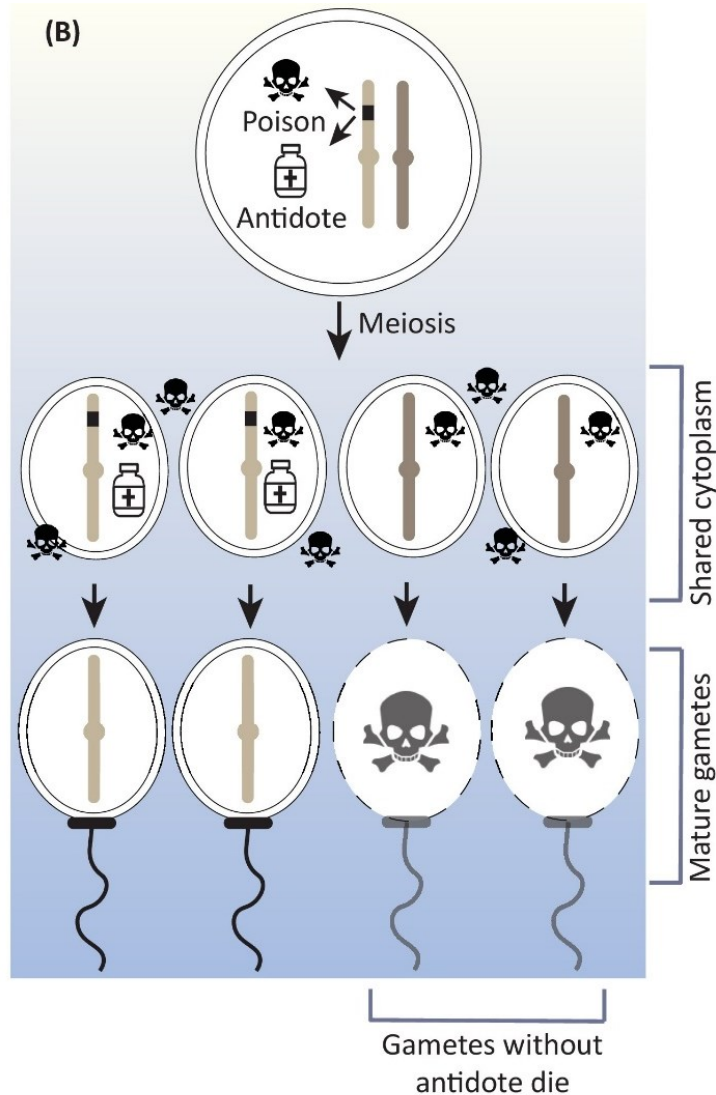
Monkey flower (*Mimulus guttatus*)

Male meiotic (gametic) drive

Spermatogenesis



Mechanisms of male meiotic drive

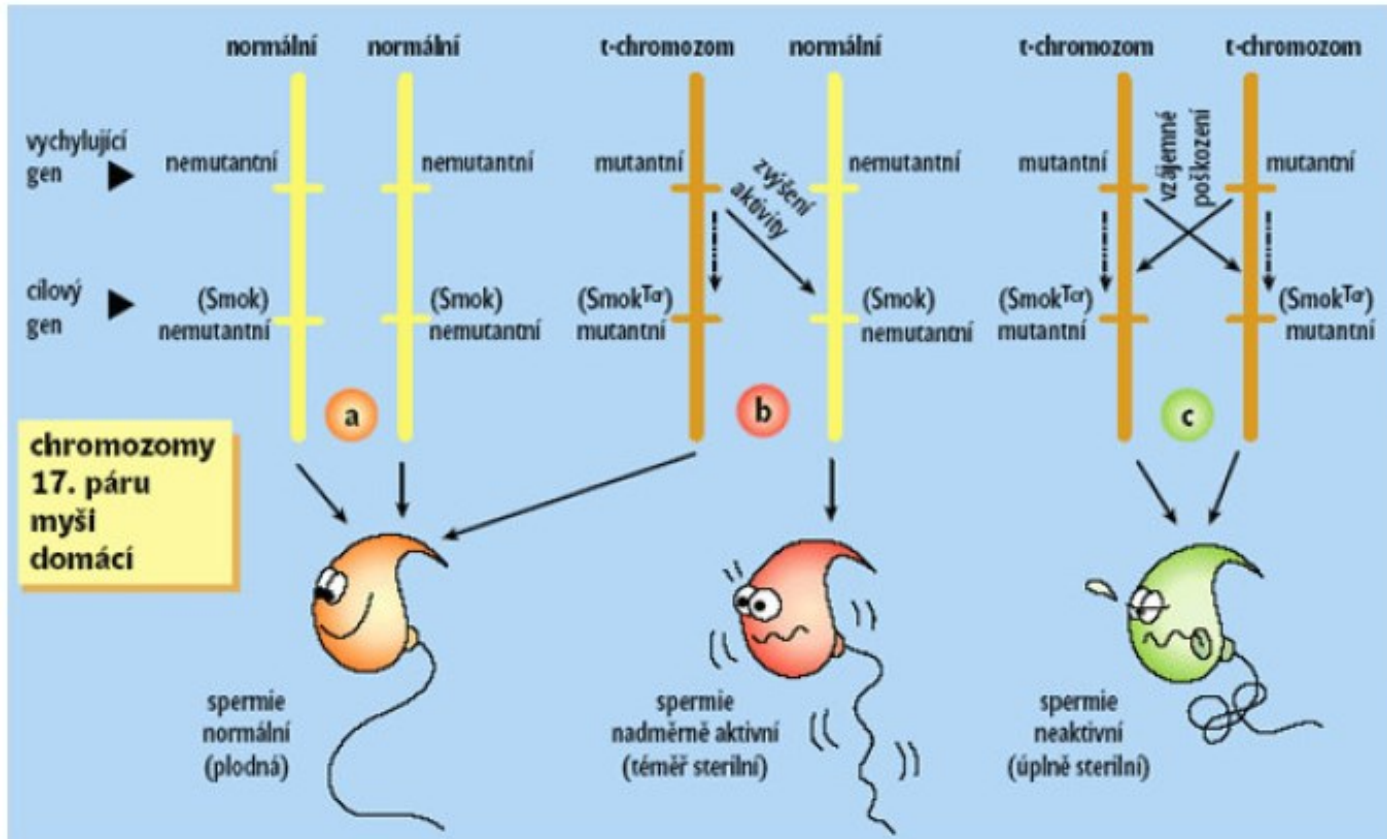


Drive mostly occur in non-recombining regions

- Inversions
- Sex chromosomes

t-haplotype

- Inversion on chromosome 17

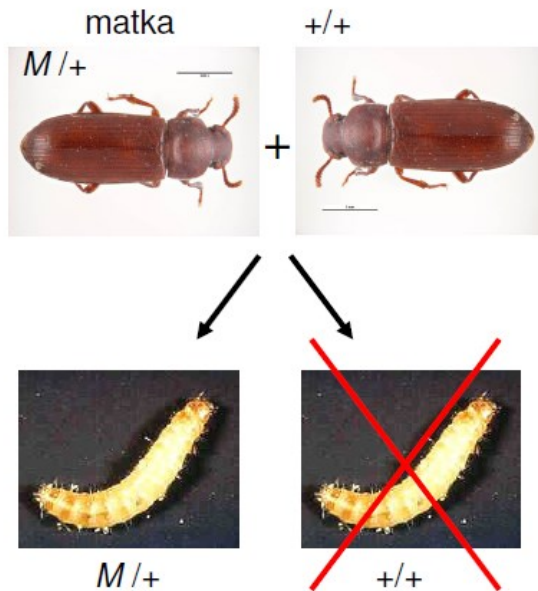


Zygotic drive

Medea (Maternal-Effect Dominant Embryonic Arrest)

- Allele causes death of progeny that do not inherit it
- Maternally expressed poison and zygotically expressed antidote.

Tribolium castaneum

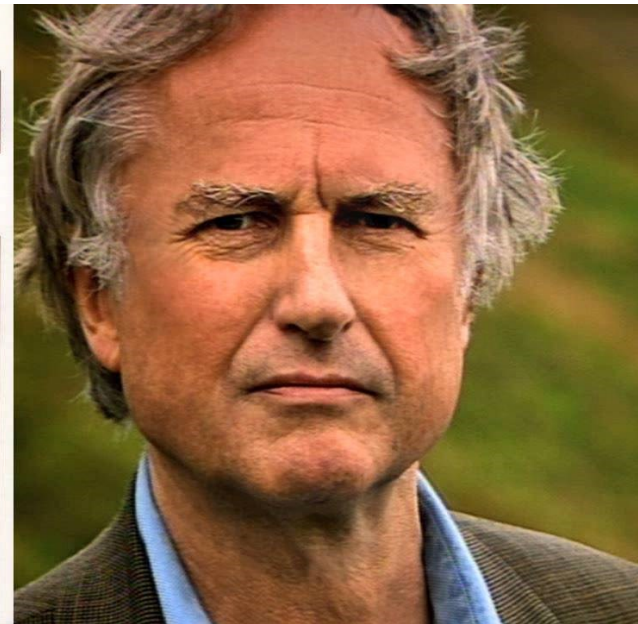
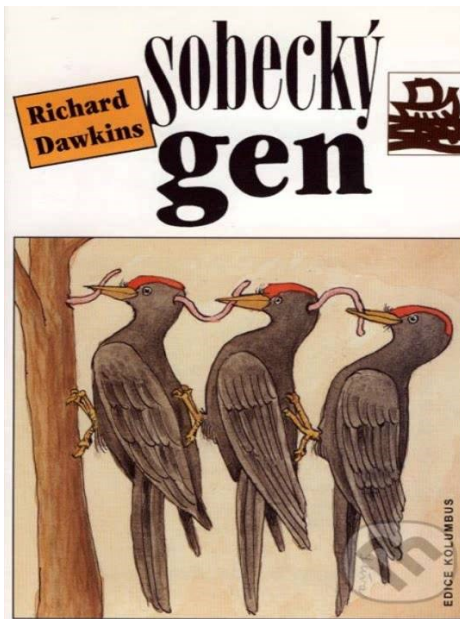
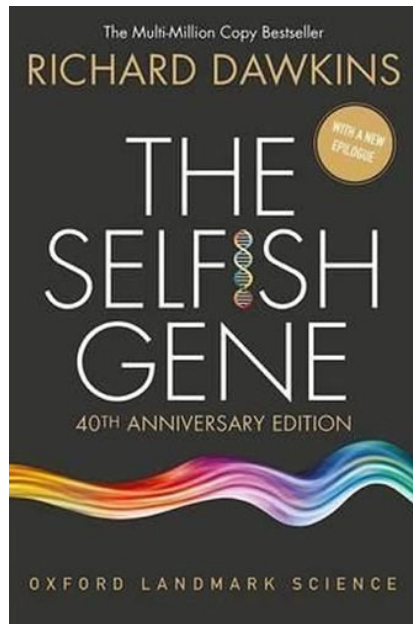


Syntetic drive

- Could be used to introduce quickly some allele to the population.
- Possible practical applications: regulation of diseseases etc.



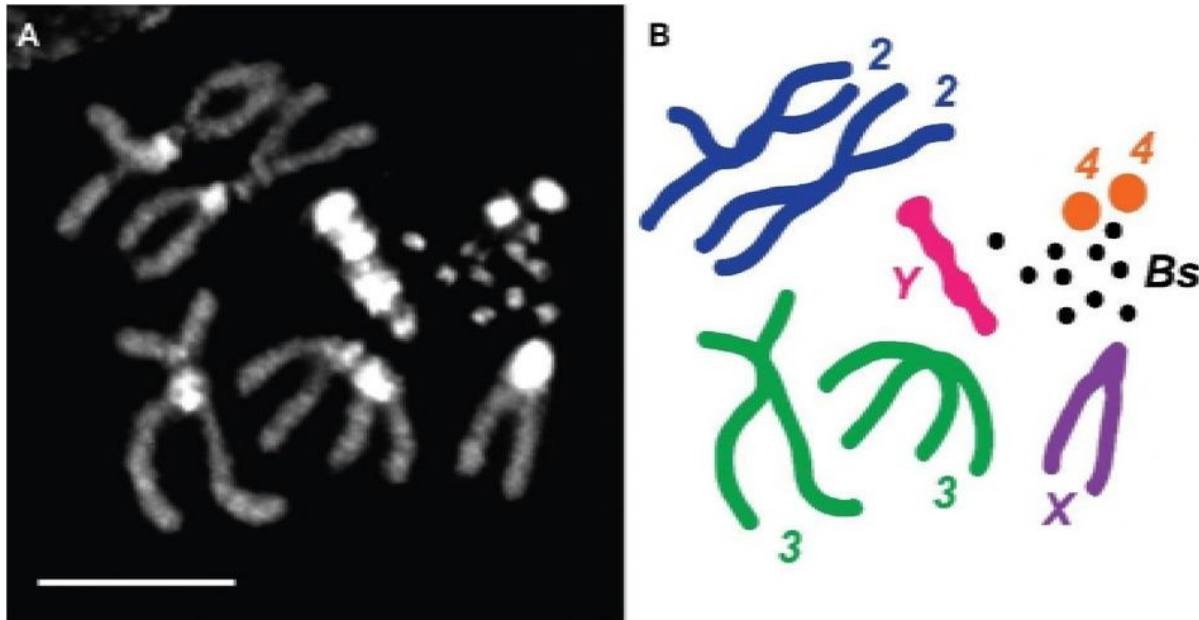
Selfish genes



- Drive genes can spread in the population even if they do not provide any advantage to the individuals carrying them.

Selfish B chromosomes

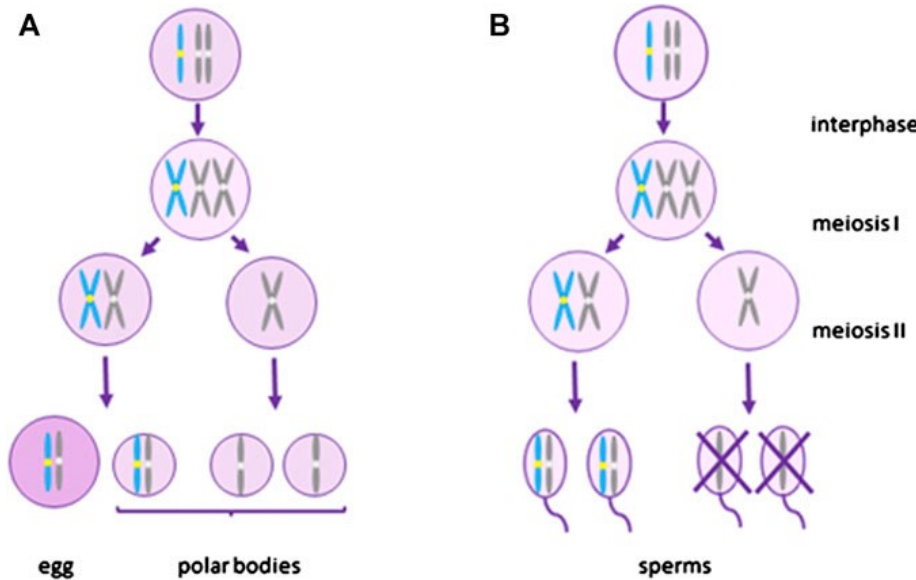
- Additional parasitic chromosomes
- Only in some individuals in population, in one or more copies
- Often spread via meiotic or mitotic drive mechanisms



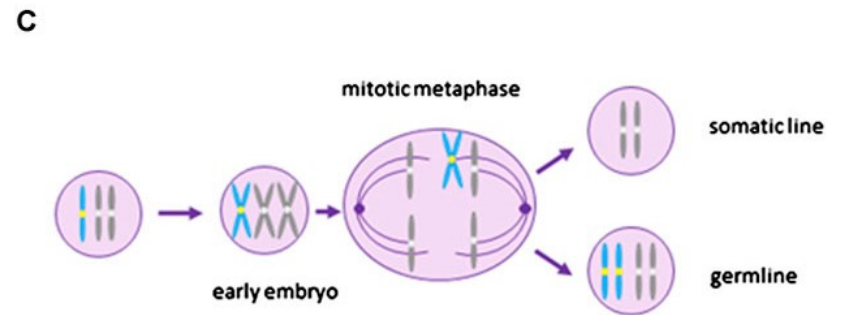
D. melanogaster B chromosomes

Selfish B chromosomes

Meiotic drive



Mitotic drive associated with gonotaxis



Lilium callosum



Pseudococcus affinis



Camnula pellucida

Selfish B chromosomes

Postmeiotic mitotic drive in angiosperms



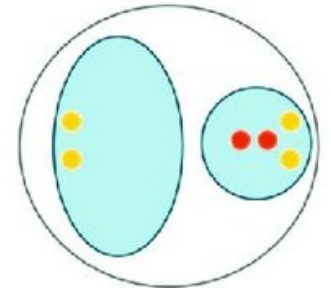
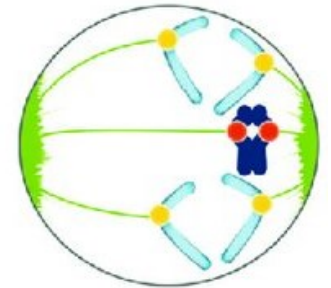
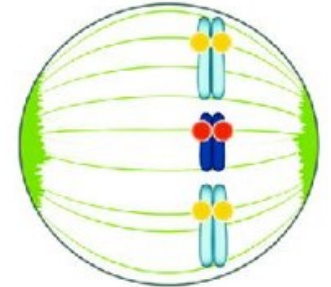
Rye (*Secale cereale*)



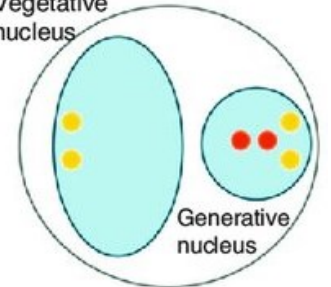
Vegetative cell

Generative cell

Drive of Bs
Asymmetric cell division



Vegetative nucleus



Generative nucleus

Tissue specific elimination of B chromosomes in plants



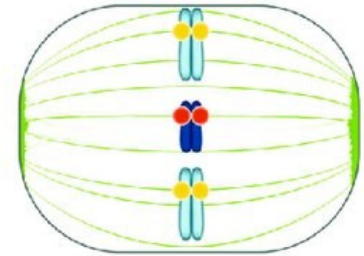
goatgrass



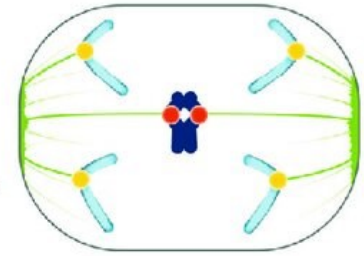
sorghum

Elimination of Bs
Symmetric cell division

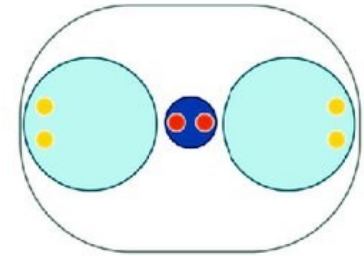
Metaphase



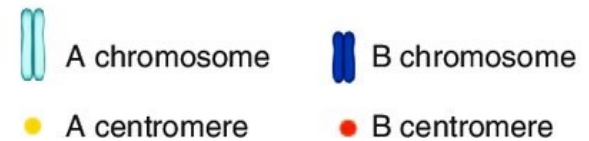
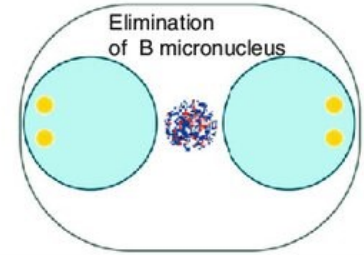
Anaphase
Nondisjunction of Bs



Telophase

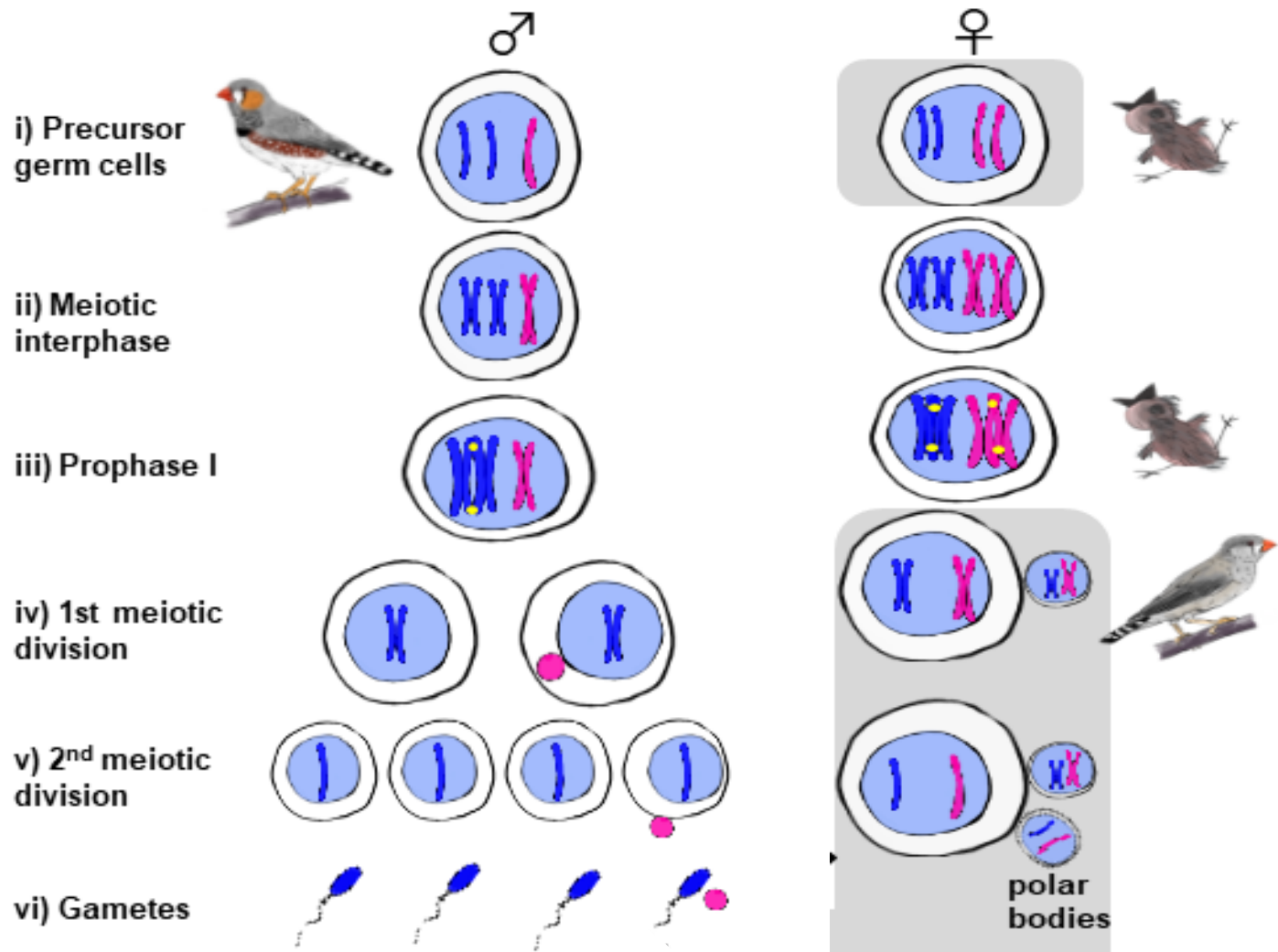


Late
Telophase



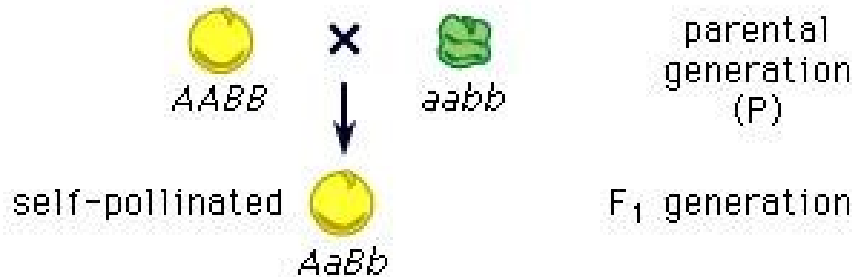
Germline-restricted chromosome

Possibly „domesticated“ B chromosome?



Mendel's theory of inheritance

Law of independent assortment



♀ \ ♂		pollen			
		AB	Ab	aB	ab
ovules	AB	$AABB$	$AABb$	$AaBB$	$AaBb$
	Ab	$AABb$	$Aabb$	$AaBb$	$Aabb$
	aB	$AaBB$	$AaBb$	$aaBB$	$aaBb$
	ab	$AaBb$	$Aabb$	$aaBb$	$aabb$

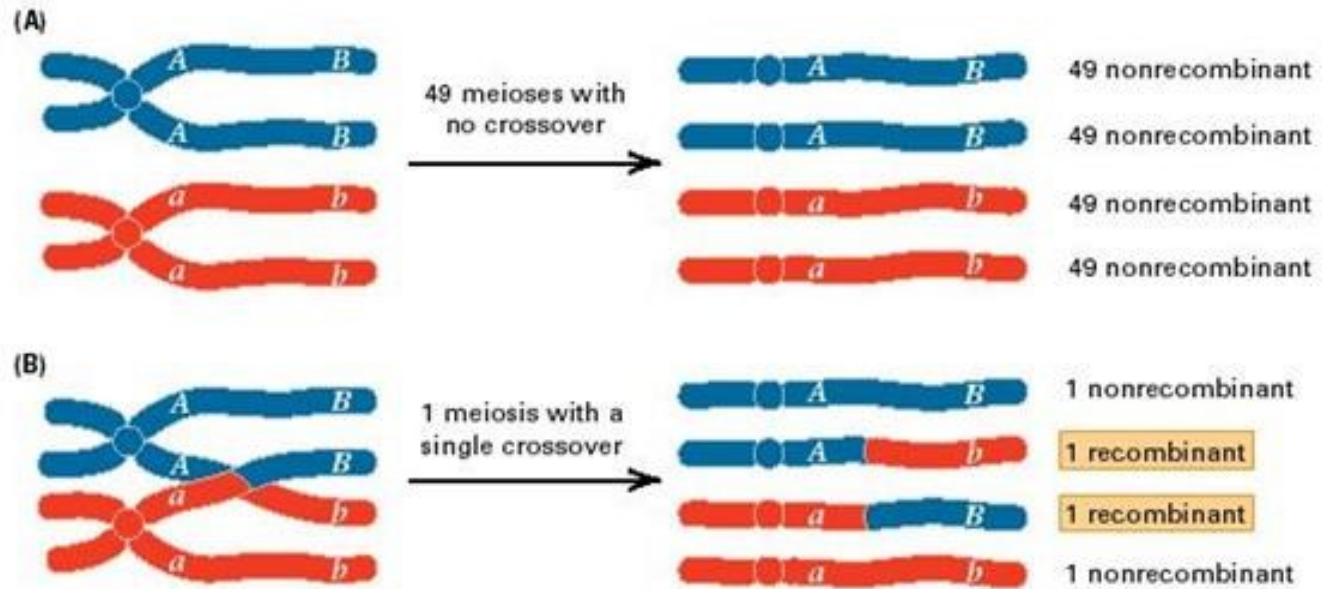
F₂ generation

Chromosomal theory of inheritance and gene linkage



Thomas Morgan

- Alleles of different genes that are localized on the same chromosome tend to be inherited together.
- 1 cM ~ 1% recombinant genotypes



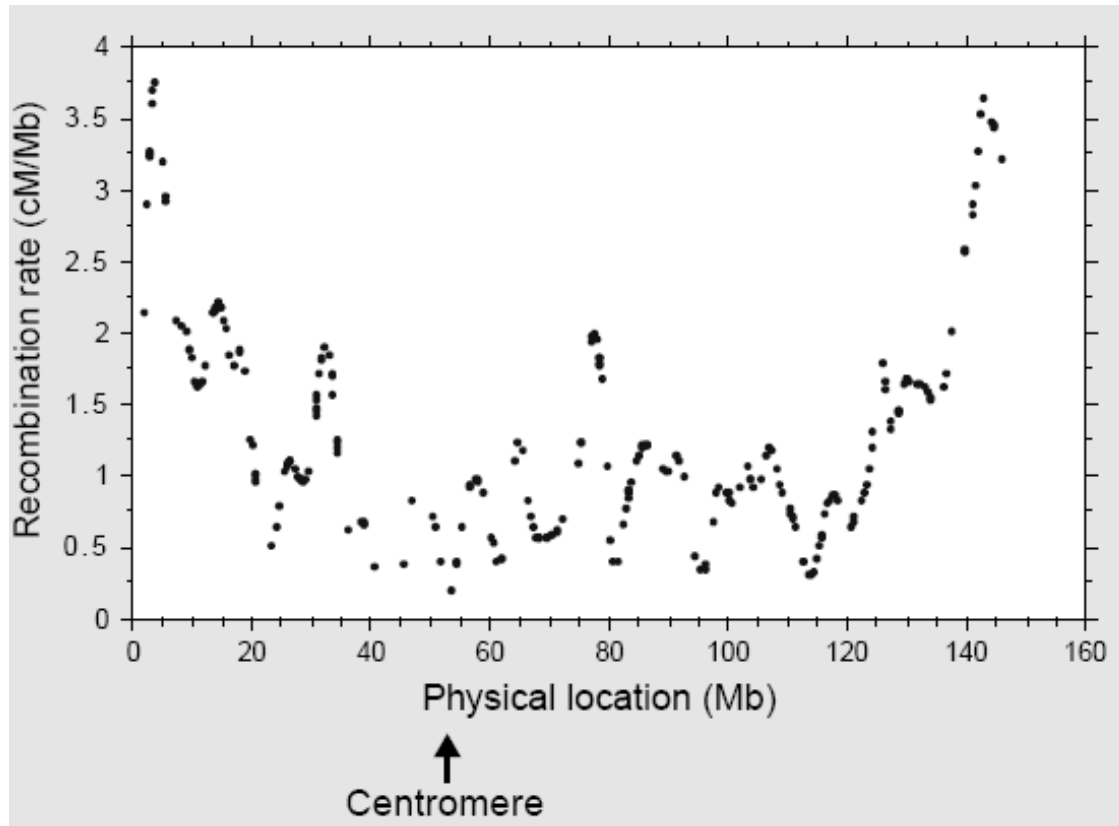
(C) Frequency of recombination:

$$r = \frac{1 + 1}{49 + 49 + 49 + 49 + 1 + 1 + 1 + 1} = \frac{2}{200}$$

= 1 percent = 1 map unit = 1 cM

Recombination rate (r)

- Higher at the ends of chromosomes (near telomeres), lower around centromeres.
- Recombination hotspots in some organisms (e.g. mammals).
- Crossing-over interference. Usually 1 crossing-over per chromosome (max. 3).
Leads to higher recombination rate in smaller chromosomes.



Recombination rate (r)

- Males usually have higher recombination rates than females.
humans: 1,7 x ; mouse 1,3 x
- **Haldane-Huxley rule.** If one sex do not recombine, it is the heterogametic sex (např. *Drosophila*, *Bombyx*).
- Absence of recombination on non-pairing sex chromosomes (Y,W).

XY males



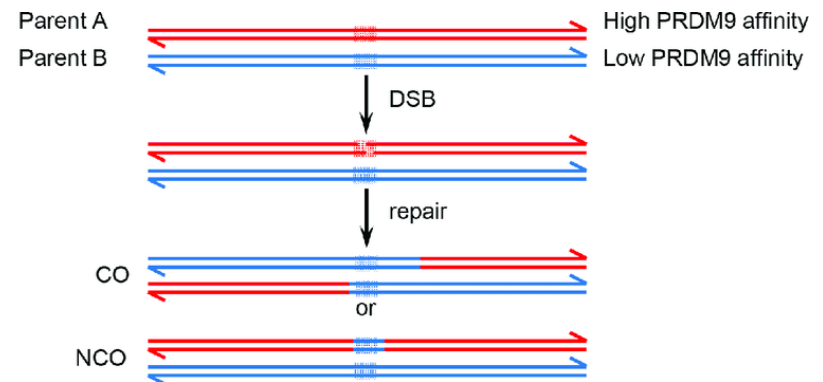
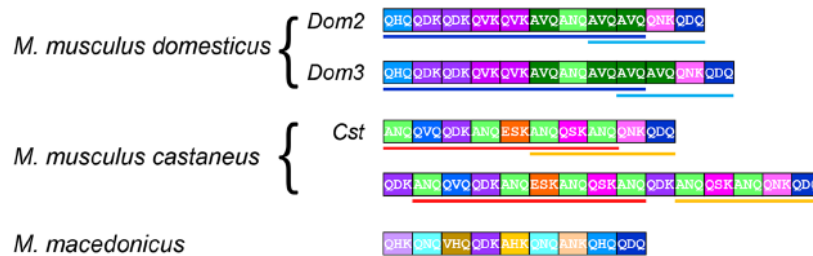
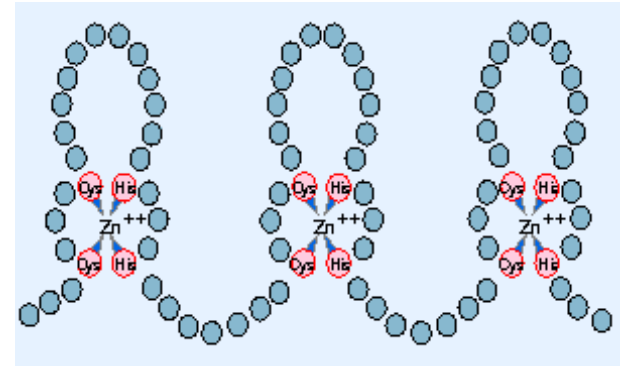
ZW females



Hotspots of recombination

Prdm9

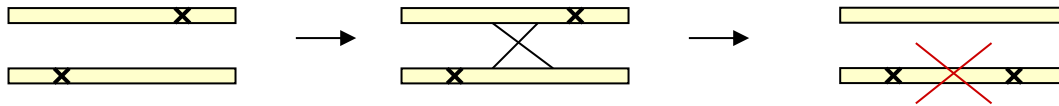
- DNA binding protein (zinc-finger domain). Determines the position of double strand breaks in meiosis (hotspots of recombination).
- Fast molecular evolution due to gene conversions.



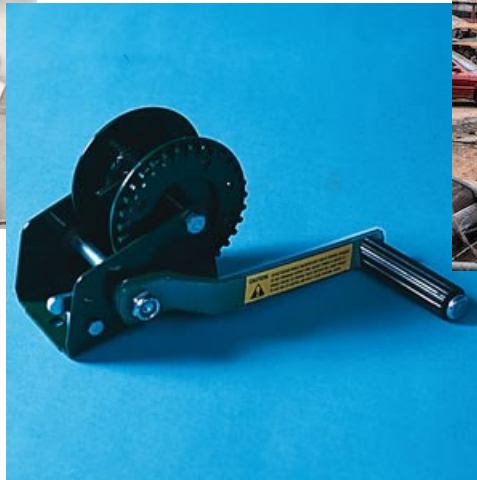
Asymmetrical gene conversion.

Evolutionary importance of recombination

- Slows down accumulation of deleterious mutations (Muller's ratchet).

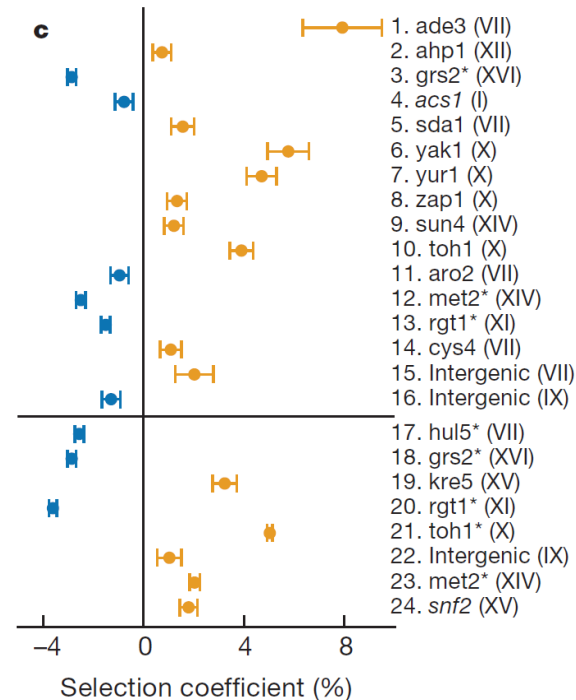
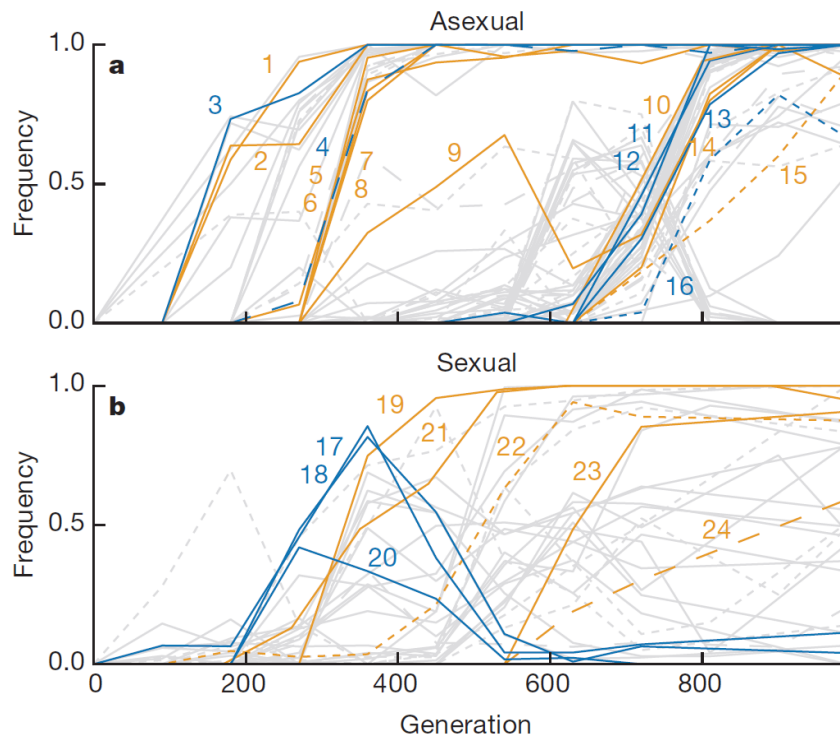
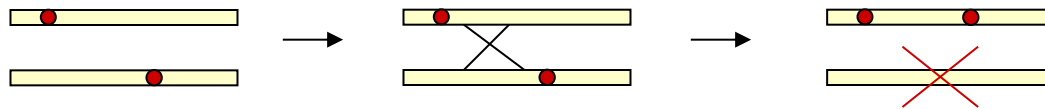


J. H. Muller



Evolutionary importance of recombination

- Allows to combine multiple advantageous mutations.
Speed up adaptive evolution.



Sometimes recombination between specific genes is disadvantageous

Supergenes



Heterostyly in *Primula vulgaris*

Received: 31 January 2024 | Revised: 24 May 2024 | Accepted: 17 June 2024
DOI: 10.1111/1755-0998.13988

RESOURCE ARTICLE

MOLECULAR ECOLOGY
RESOURCES WILEY

The *Primula edelbergii* S-locus is an example of a jumping supergene

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Funding information
Schweizerischer Nationalfonds zur Förderung der Wissenschaftlichen Forschung, Grant/Award Number: 175556

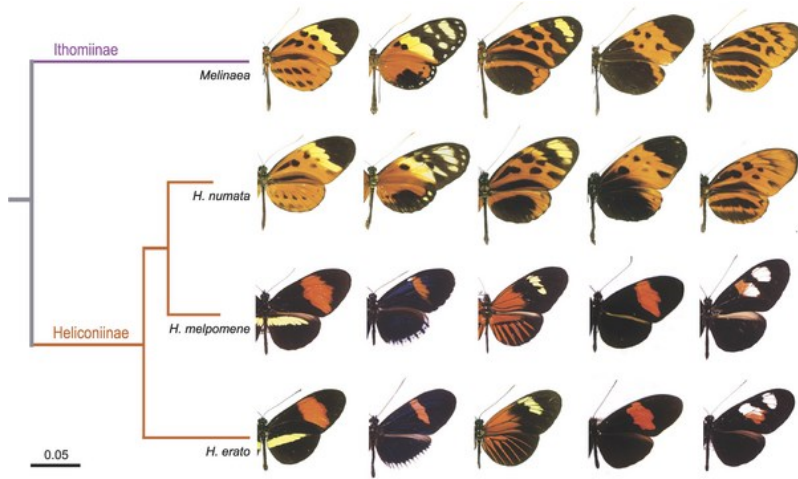
Abstract

Research on supergenes, non-recombining genomic regions housing tightly linked genes that control complex phenotypes, has recently gained prominence in genomics. Heterostyly, a floral heteromorphism promoting outcrossing in several angiosperm families, is controlled by the S-locus supergene. The S-locus has been studied primarily in close relatives of *Primula* and other groups that independently evolved heterostyly; however, it remains unknown whether genetic architecture and composition of the S-locus are maintained among species that share a common origin of heterostyly and subsequently diverged across larger time scales. To address this research gap, we present a chromosome-scale genome assembly of *Primula edelbergii*, a species that shares the same origin of heterostyly with

- Co-adapted gene complexes underlying complex traits. Inherited as a single gene.

Supergenes

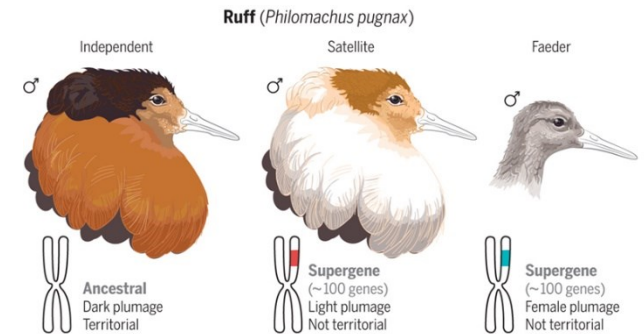
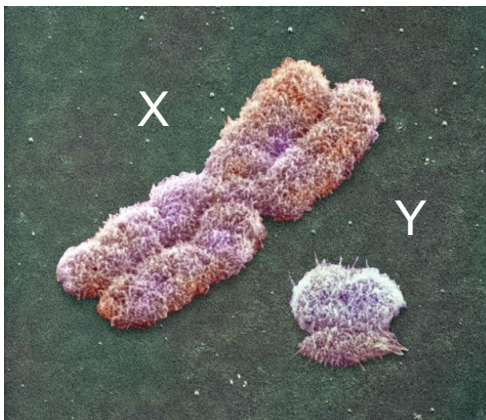
Mimetic polymorphism in *Heliconius numata*



Male morphs in ruffs

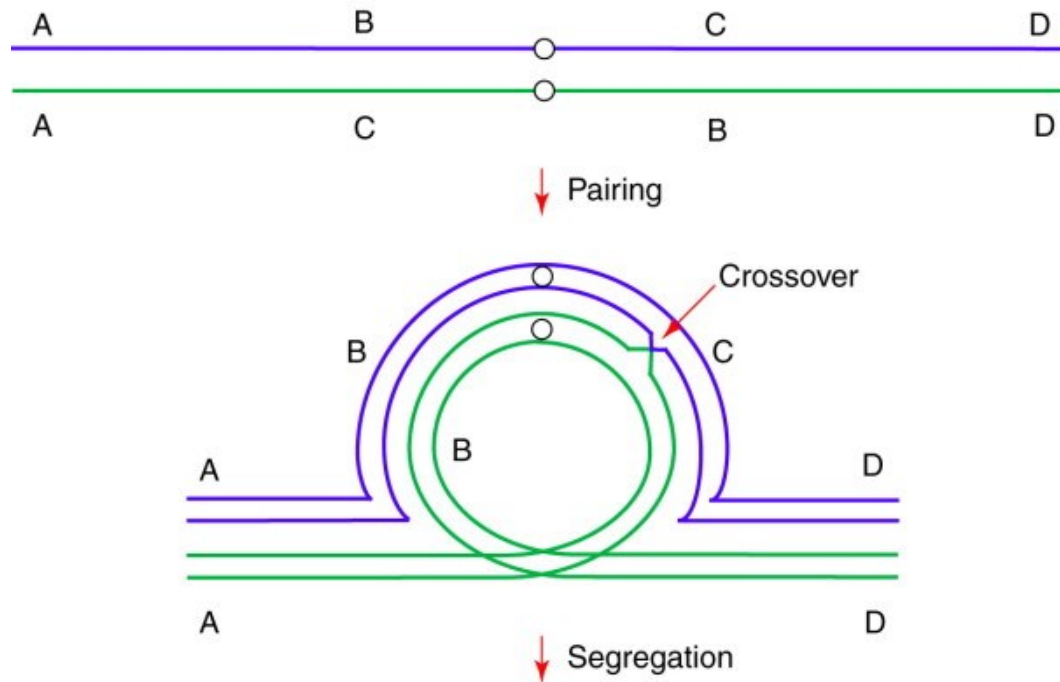


Sex chromosomes

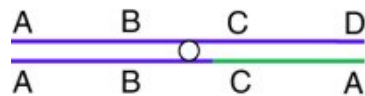


Kupper et al. 2016 Nat. Genet.

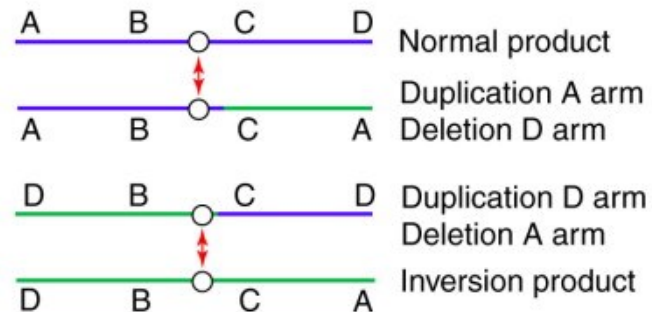
Inverze účinně potlačuje rekombinaci



End of meiosis I

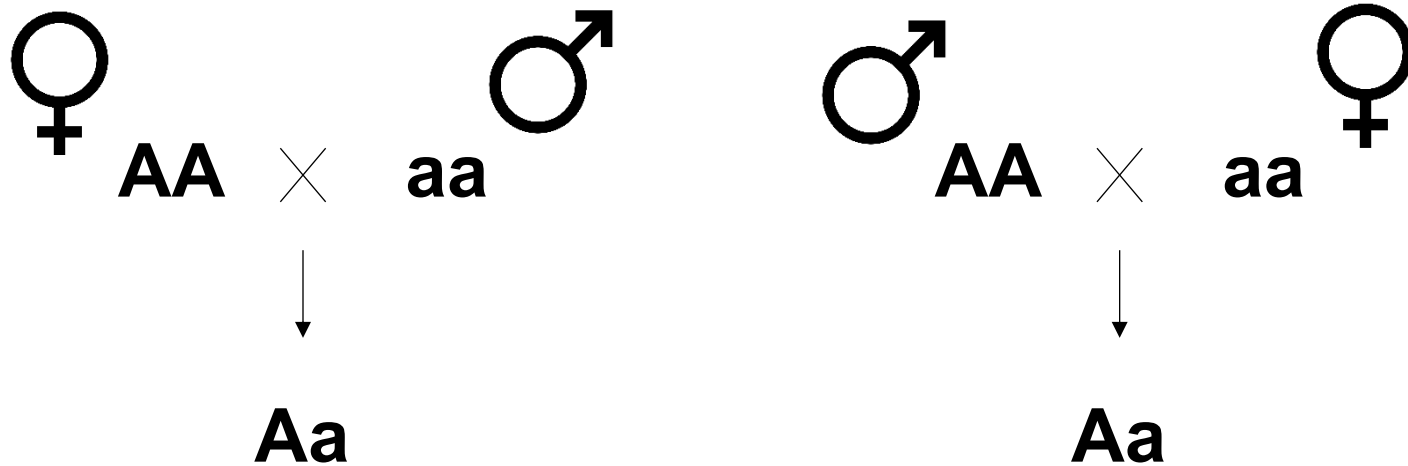


End of meiosis II



Mendel's theory of inheritance

Law of uniformity and identity of reciprocal F1 hybrids



Uniparental inheritance

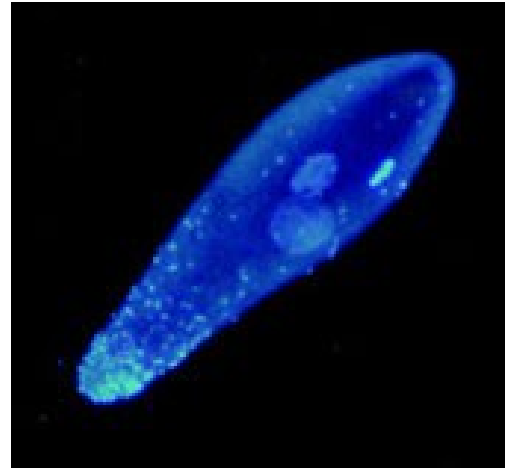
Mitochondrial and plastid DNA

- Mostly maternal inheritance
- „Mothers curse“
- In rare cases paternal inheritance (molluscs).
- Most genes from mitochondrial and plastid DNA moved to nukleus.



Intracellular parasite of Arthropods

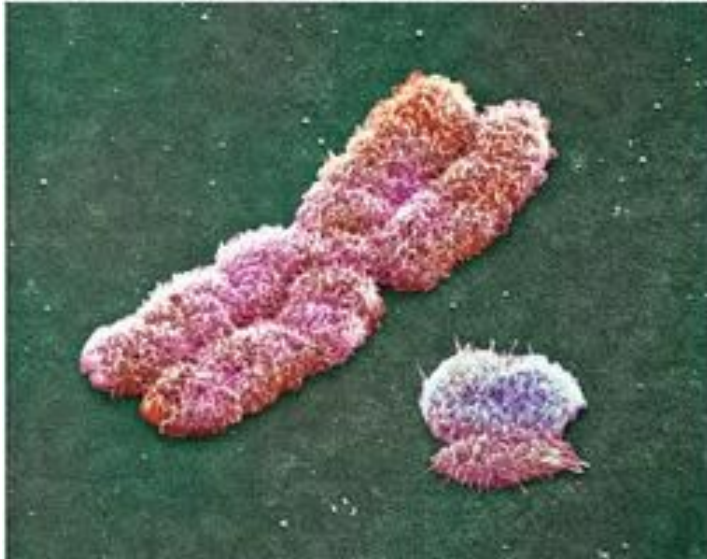
Wolbachia



- cytoplasmic incompatibility (infected males cannot reproduce with uninfected females)
- parthenogenesis (infected females can reproduce without males)
- feminization of males
- killing males

Sex chromosomes

- Chromosom Y paternal inheritance.
- Chromosom W maternal inheritance.



XY males
XX females

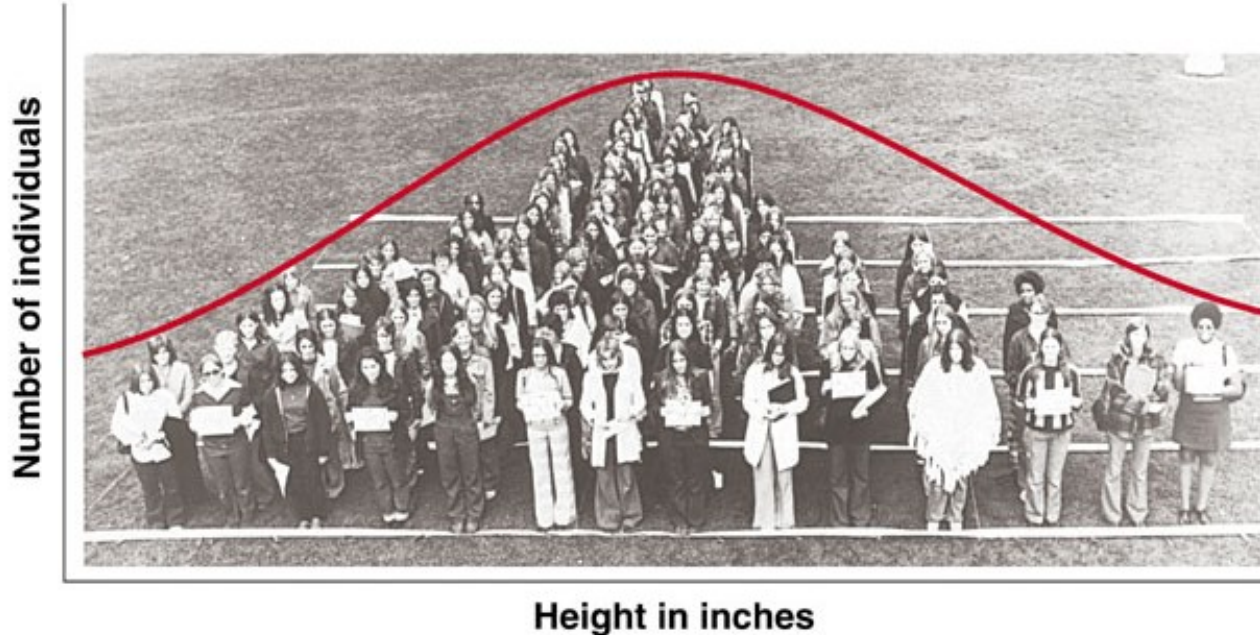


ZW females
ZZ males



Inheritance of quantitative traits

- continuous variation of traits
- traits underlined by many genes (interactions among genes)
- traits are often affected by environment
- inheritance can be less predictable



Heritability (H^2 , h^2)

- Proportion of variance in a phenotype caused by genetic factors.

$$H^2 = V_G/V_P$$

$$V_P = V_G + V_E$$

V_P - phenotype

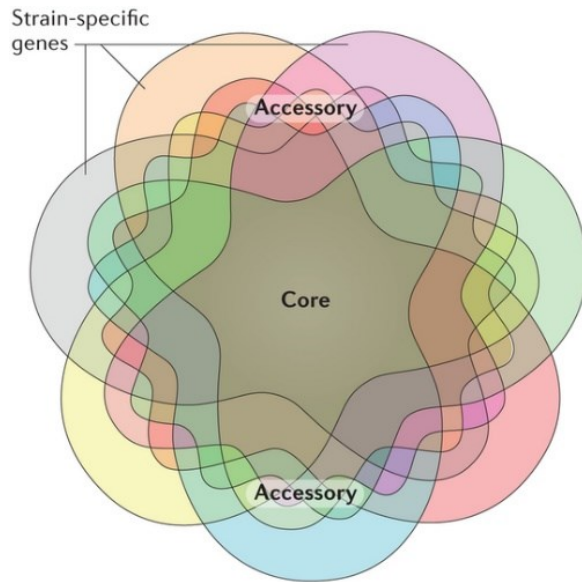
V_G - genotype

V_E - environment

- Range between 0 and 1.
- Traits with higher heritability better respond to selection .

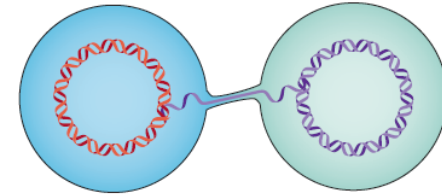
Horizontal gene transfer

- Widespread in prokaryotes
- pangenom = set of all genes of the given taxonomical group

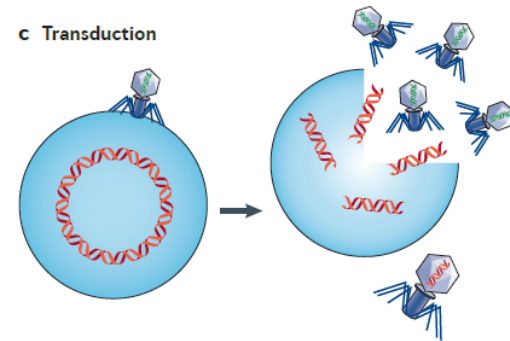


Nature Reviews | Genetics

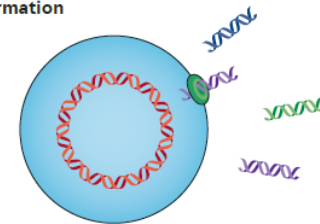
a Conjugation



c Transduction



e Transformation






PRESENTATION

PNAS

RESEARCH ARTICLE | EVOLUTION

 OPEN ACCESS

Contingency, repeatability, and predictability in the evolution of a prokaryotic pangenome

Alan J. S. Beavan ^a, Maria Rosa Domingo-Sananes ^{a,b}, and James O. McInerney ^{a,1}

Edited by W. Doolittle, Dalhousie University, Halifax, NS, Canada; received March 27, 2023; accepted November 5, 2023

December 26, 2023 | 121 (1) e2304934120 | <https://doi.org/10.1073/pnas.2304934120>

Significance

Different strains of the same prokaryotic species often show significant variation in gene content. Whether this variation is due to genetic drift or selection is not well understood. If the latter, we expect sets of genes to be consistently and repeatedly gained or lost together, or sequentially. We used machine learning to predict the presence of variable genes in a large set of *Escherichia coli* strains, using other variable genes as predictors. We find a large proportion of genes are predictable, suggesting selection plays a role in their acquisition, loss, and maintenance. We show that some genes are consistently associated with the presence or absence of others. These results have implications for understanding evolutionary dynamics in prokaryotic genomes.

Horizontal gene transfer

- Less frequent in multicellular organisms.
- Movement of genes from mtDNA to nucleus.
- Transfer of genes from endosymbionts to the host.



Wolbachia → hmyz
(např. *Drosophila*),
hlístice



Elysia chlorotica



*Acyrtosiphon
pisum* (aj.) – syntéza
karotenoidů (původ:
houby)

Sex determination in *Armadillidium*

- *Wolbachia*, can cause feminization of males.
- Horizontal gene transfer from *Wolbachia* to *Armadillidium*.
The transferred *Wolbachia* genes determine female sex in *Armadillidium*..



Wolbachia

Svinka obecná
Armadillidium vulgare