

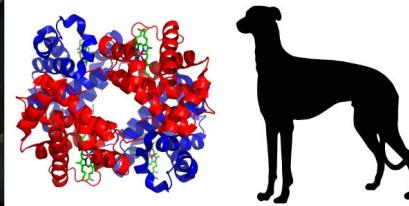
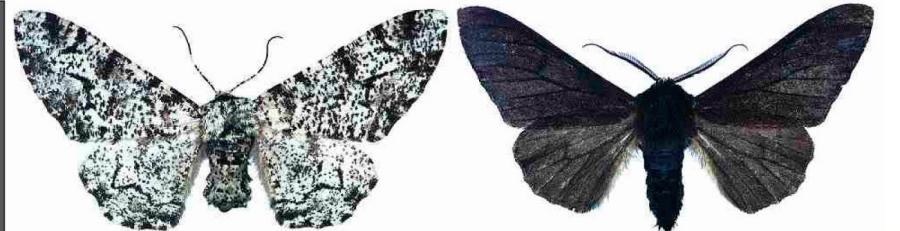
Meta-analysis of the genes underlying phenotypic changes

**Virginie Courtier-Orgogozo
Institut Jacques Monod, Paris**

www.gephebase.org

GepheBase

The Database of Evolutionary Genotype-Phenotype Relationships



www.gephebase.org

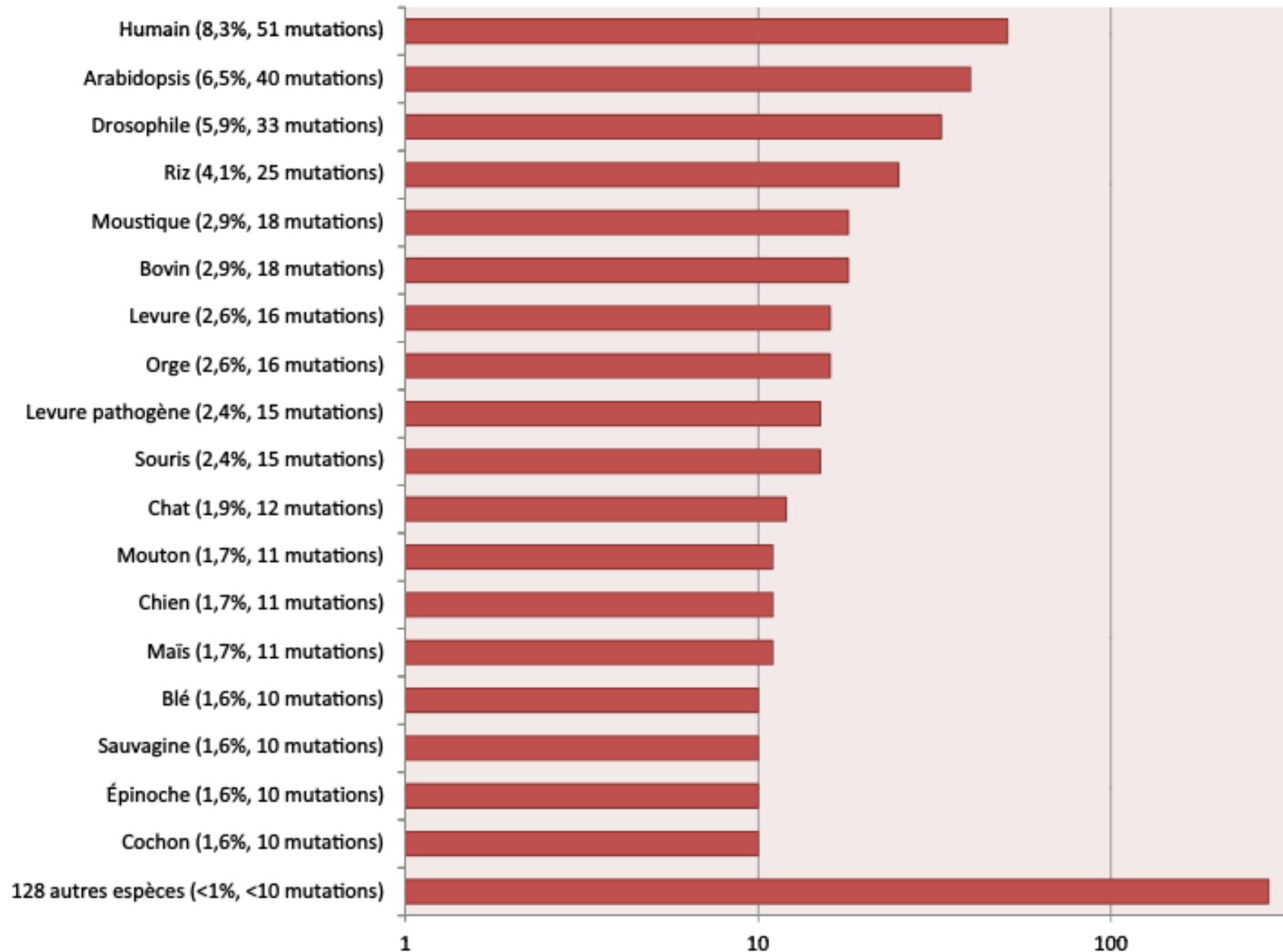
**Includes Natural, Domesticated and Experimental Variation
but *NO LAB MUTANTS* and *NO CLINICAL TRAITS***

>1600 genes and mutations
associated with
natural phenotypic changes
in animals and plants

Virginie Courtier-Orgogozo
Arnaud Martin (Washington DC)
Laurent Arnoult (former postdoc)
Stéphane Prigent (former postdoc)

Biases in current data

Bias toward certain species



Coding: global change

OVATE coding region
(Liu 2002)



myostatin coding region
(Grobet 1997)



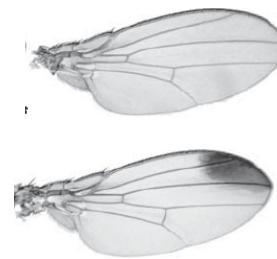
Mc1r coding region
(Eizirik 2003)



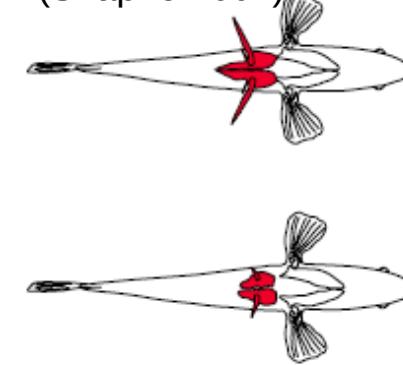
Cis-regulatory: local change

spatially

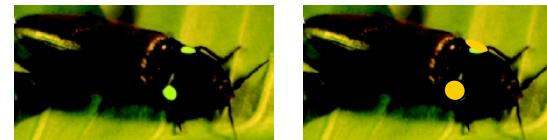
yellow cis-reg. region
(Prud'homme 2006)



Pitx1 cis-reg. region
(Shapiro 2004)



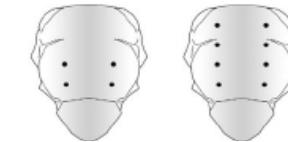
luciferase coding region
(Stoltz 2003)



anthocyanin-2 coding region
(Quattrocchio 1999)



scute cis-reg. region
(Marcelini 2006)

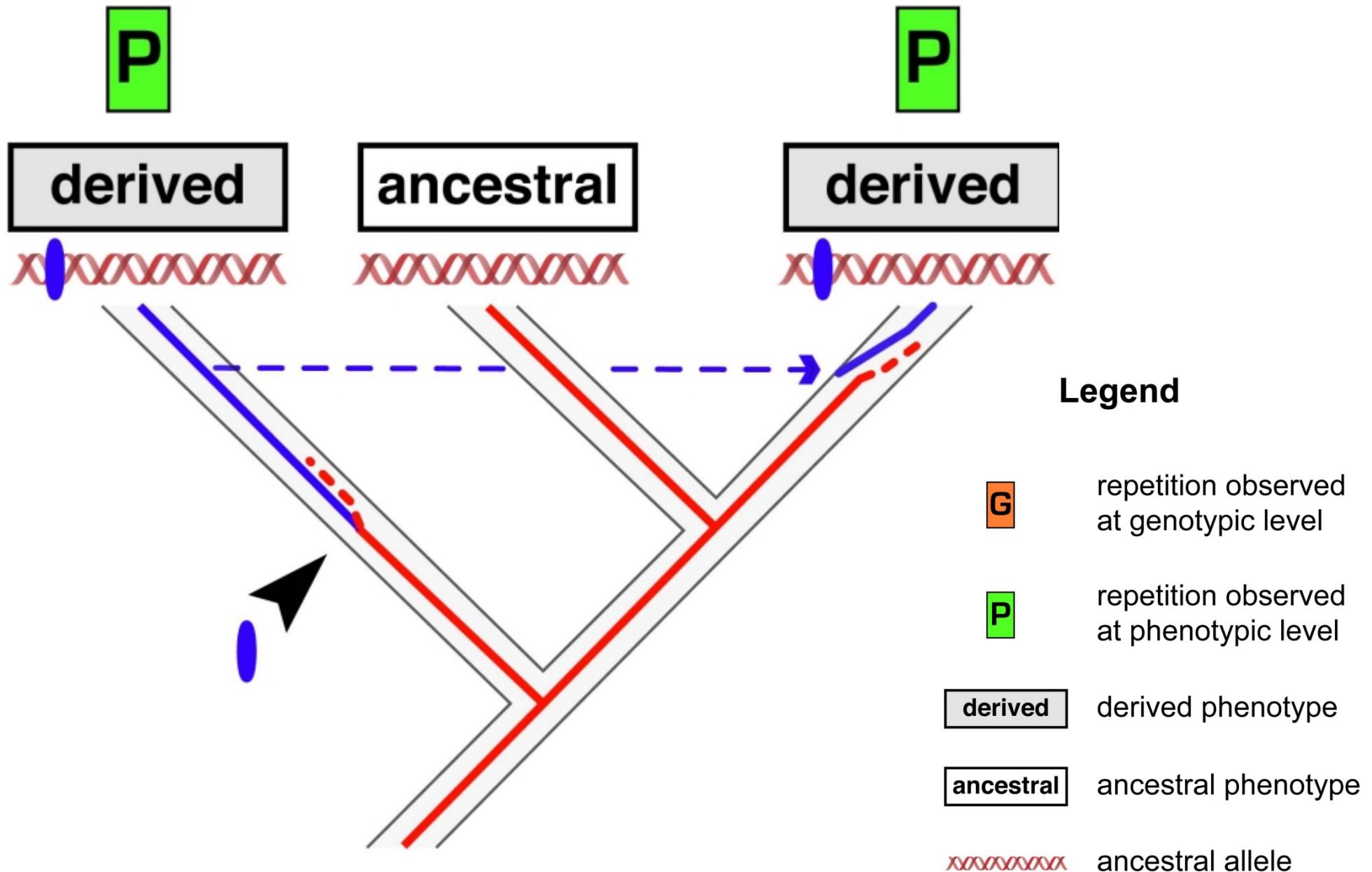


agouti cis-reg. region *temporarily*
(Jones 2018)



The tree of life is not a tree

Lateral transfer





L.S. Mills research photo by Jaco and Lindsey Barnard

Genetic plagiarism of body color in aphids

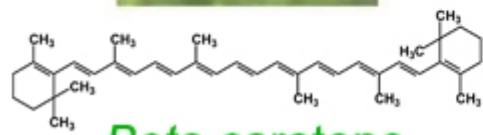
Green morph



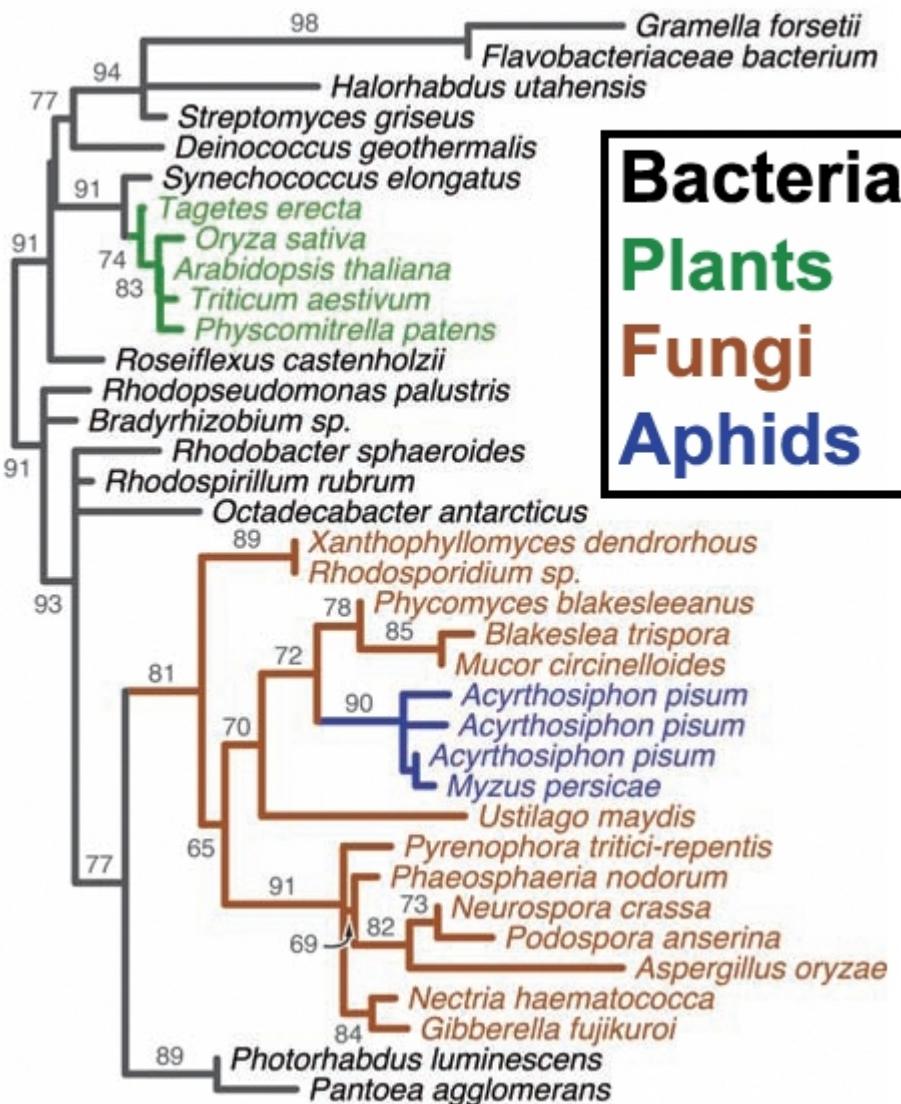
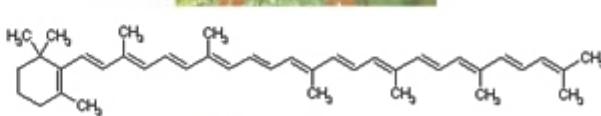
Red morph



Beta-carotene



Torulene



- Moran and Jarvik 2010
Science

Genetic plagiarism of body color in aphids

Green morph

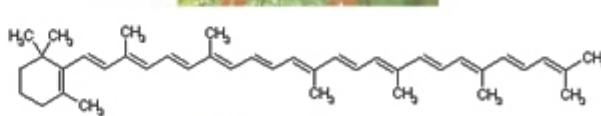


Red morph

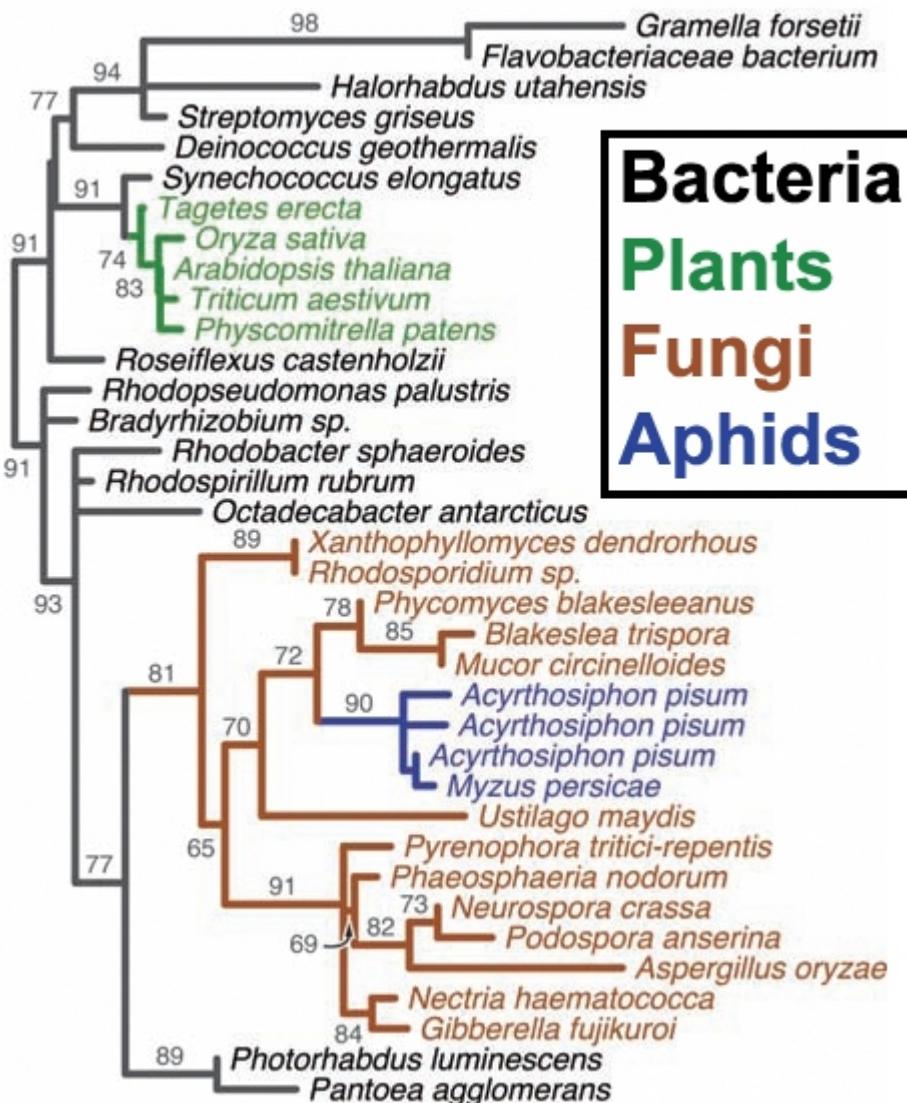


Beta-carotene

Torulene



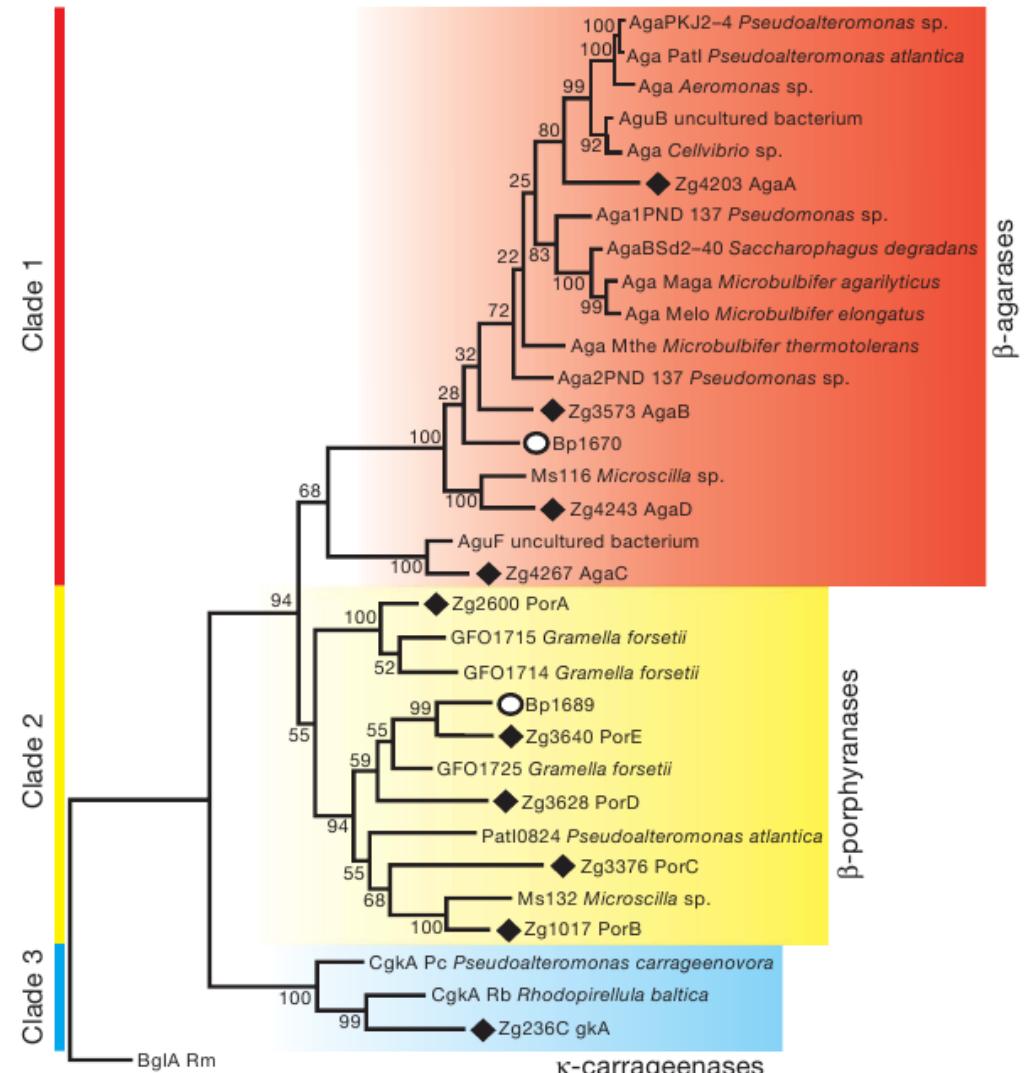
Carotenoid biosynthesis enzymes transferred
also from fungi to the two-spotted spider mite
(Altincicek 2012)



- Moran and Jarvik 2010
Science

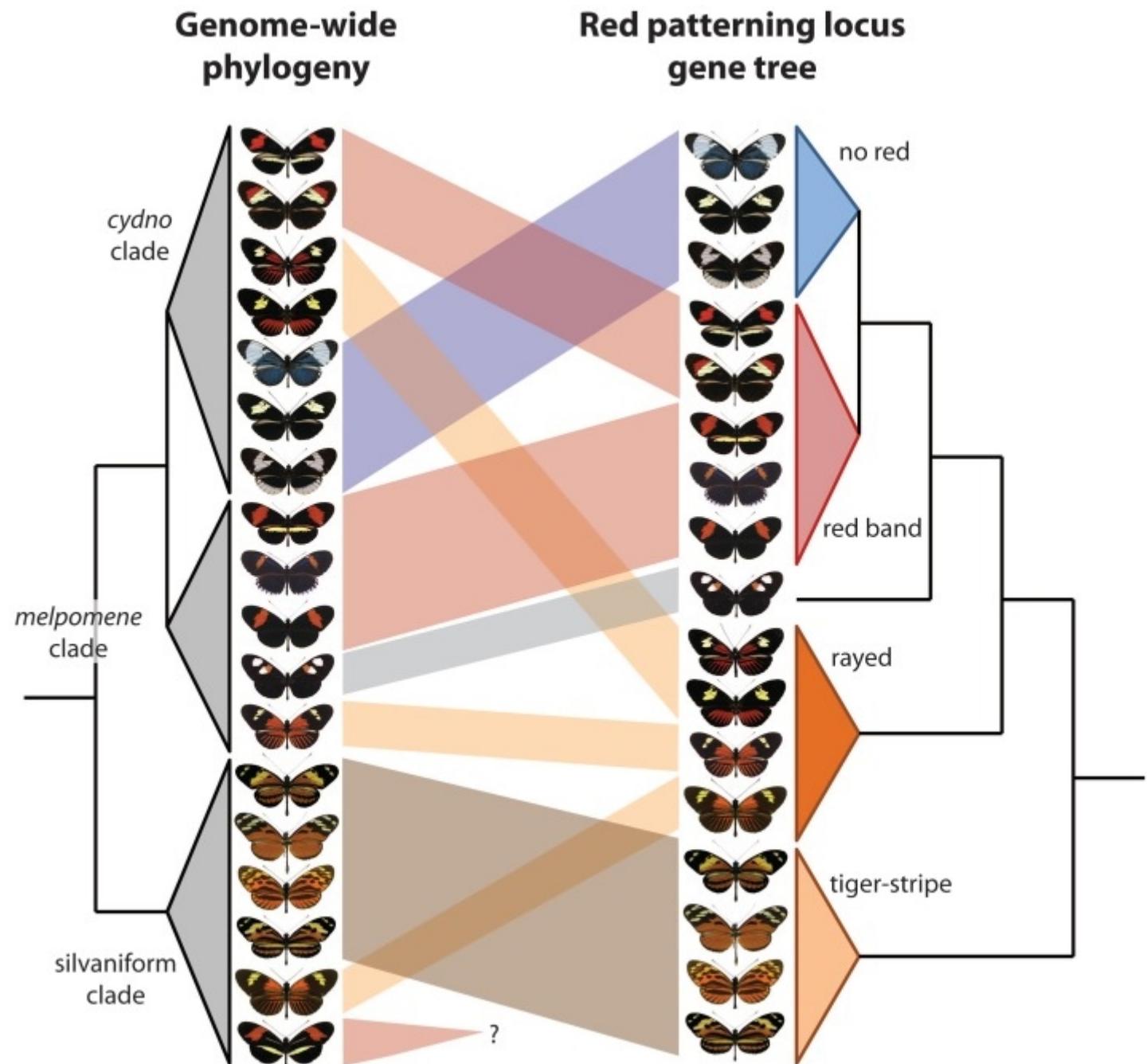
We are what we eat

Seaweed digestion in Japanese people

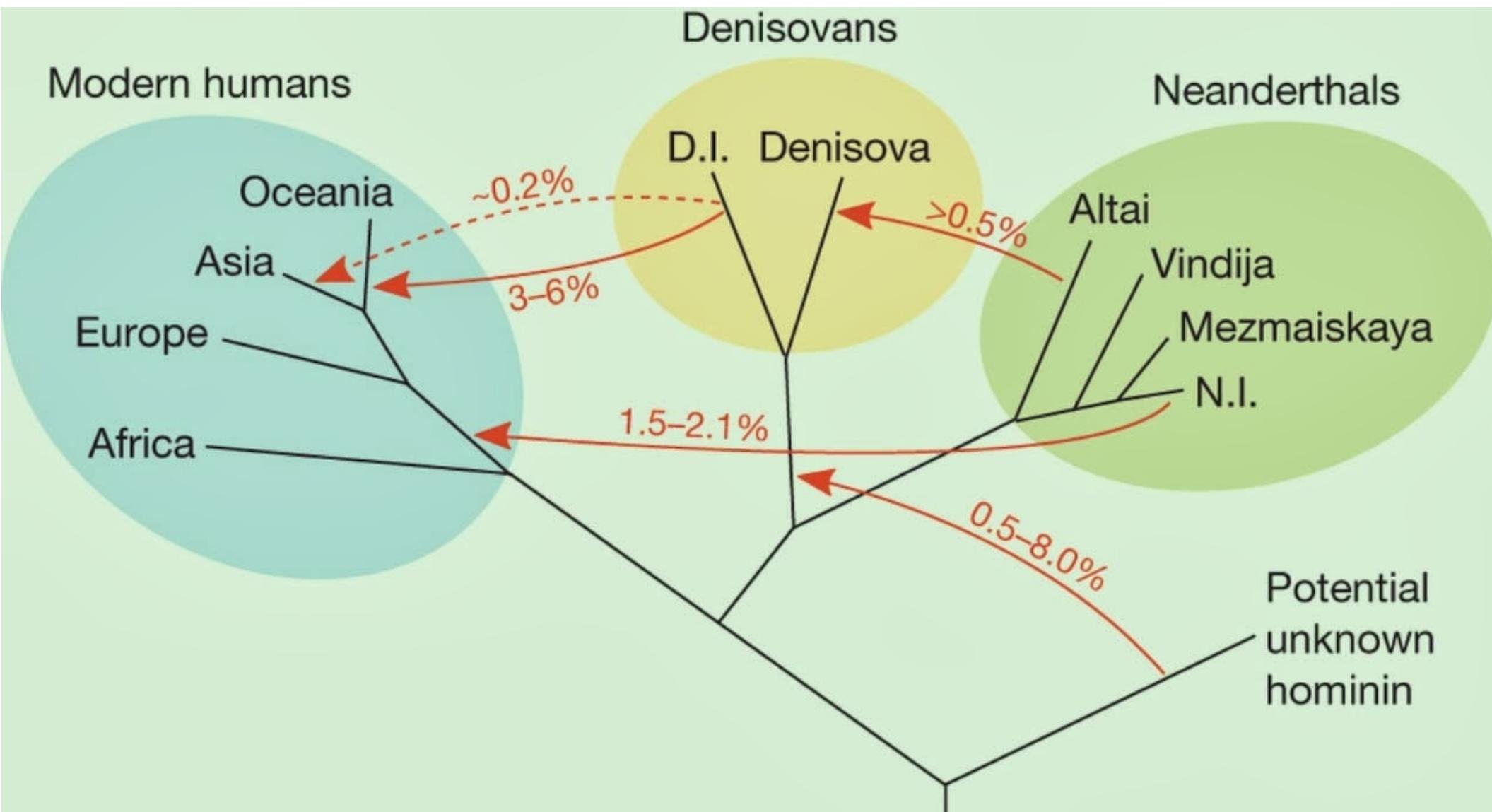


○ *Bacteroides plebeius*
from Japanese microbiome only!

Genetic plagiarism of the red wing color pattern

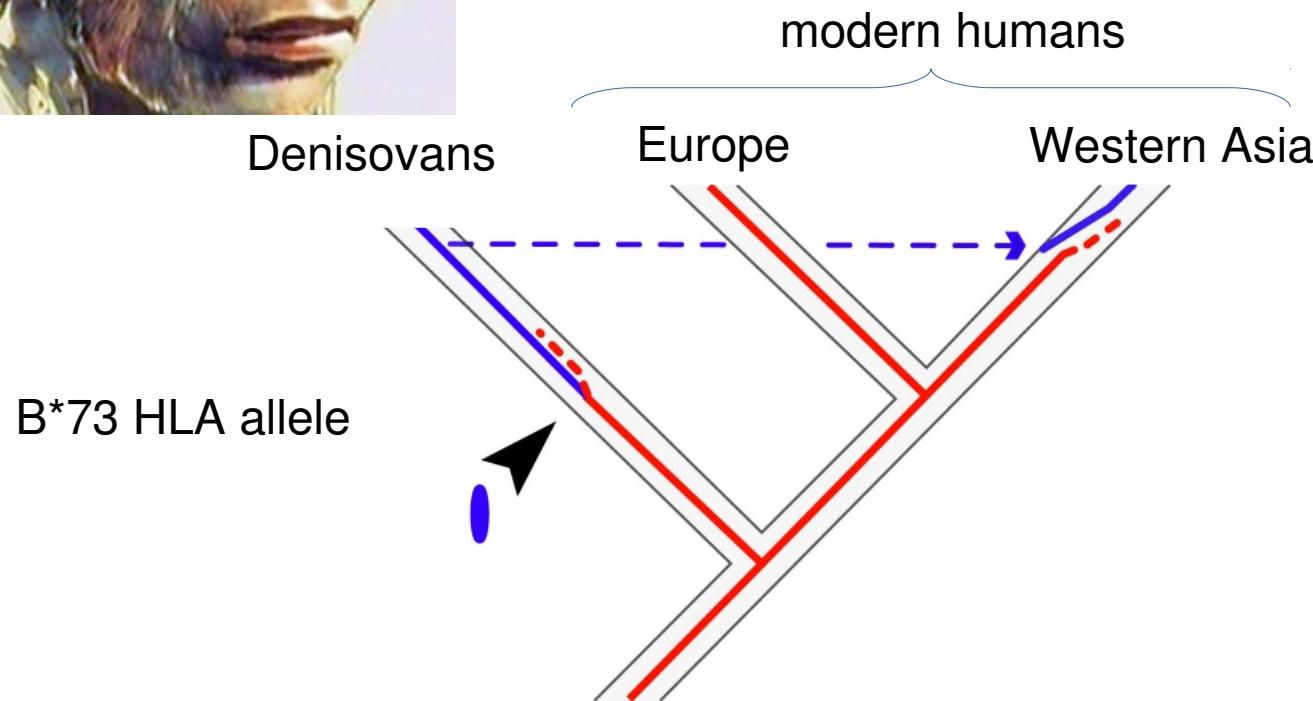


(Reed 2011, Pardo-Diaz 2012,
Heliconius Consortium 2012)



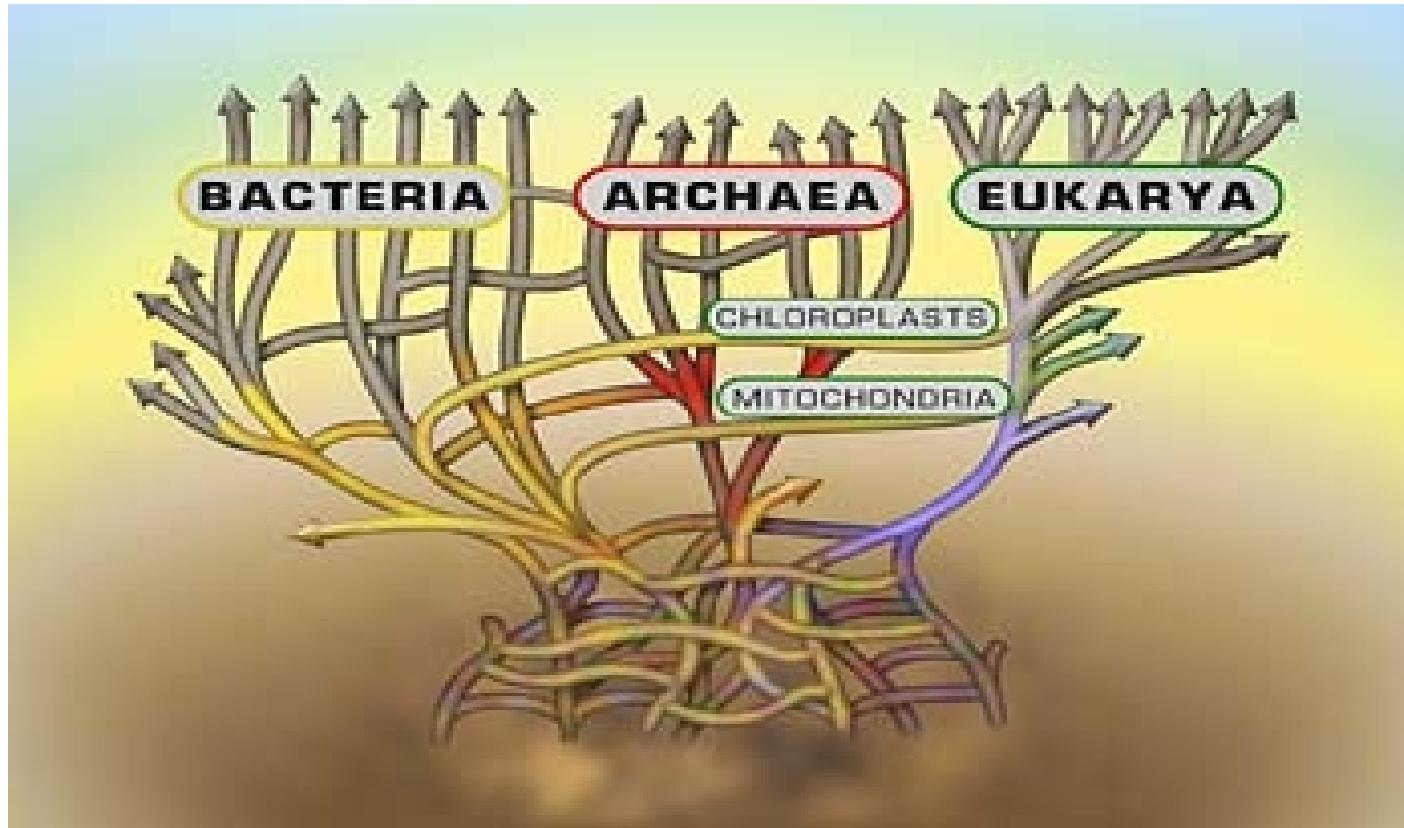
Genetic plagiarism of immune resistance genes

Siberia to Southeast Asia



Abi-Rached et al 2011 Nature

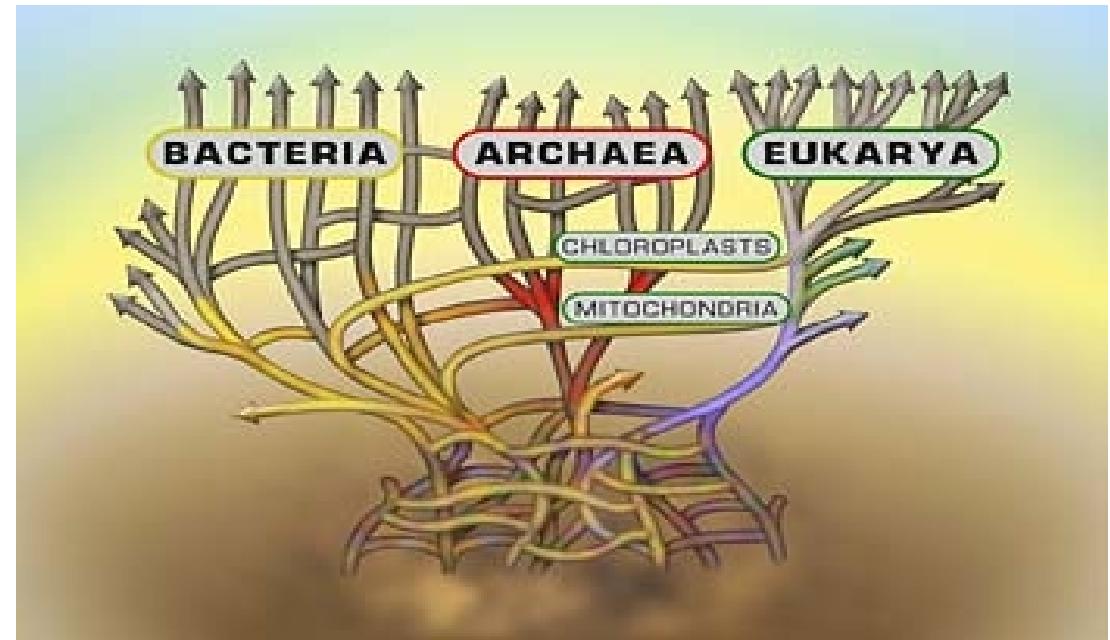
The tree of life is not a tree



Even more interconnections!

Exchange of metabolites throughout the tree of life

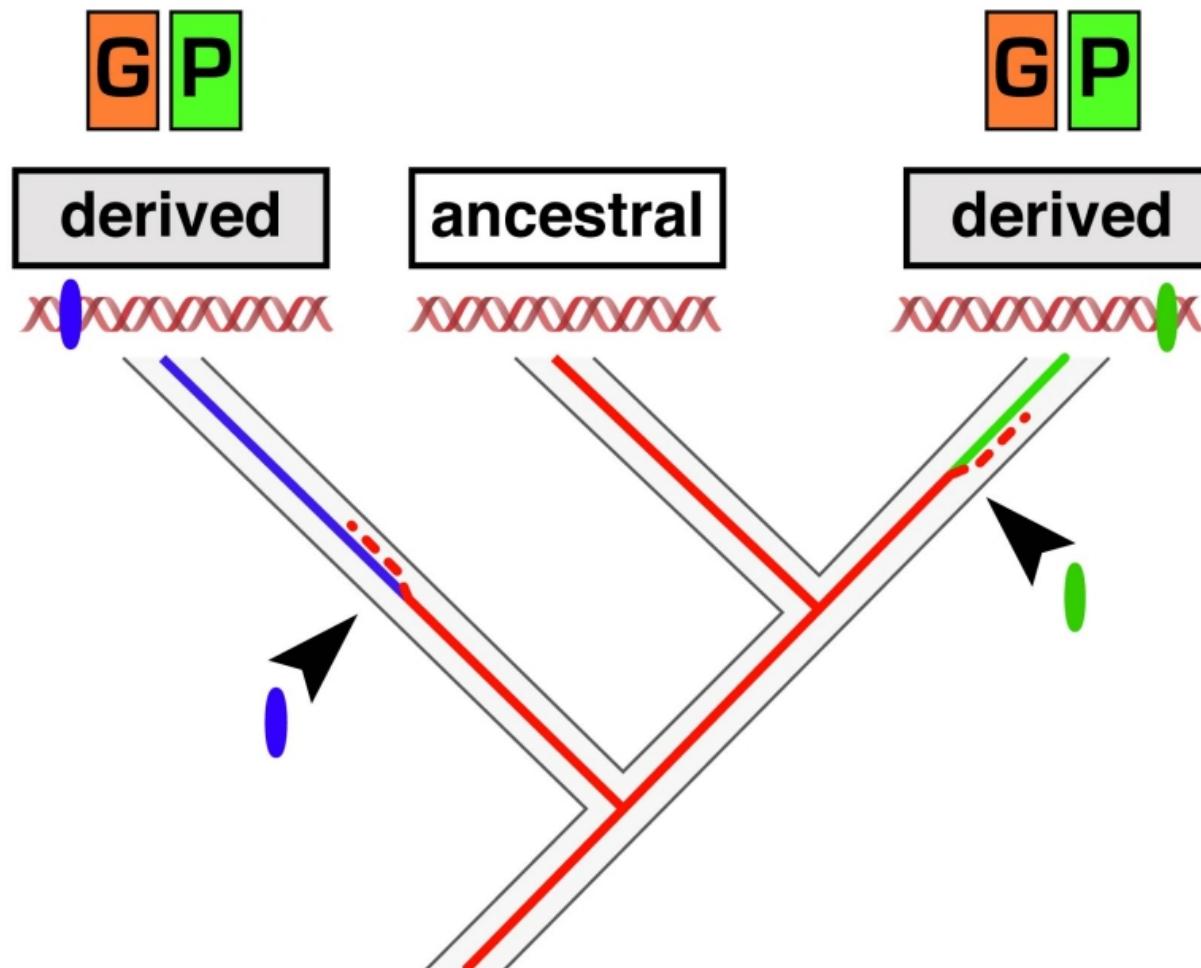
Arginine
Histidine
Isoleucine
Leucine
Lysine
Methionine
Phenylalanine
Threonine
Valine
Ascorbic acid (Vitamin C)
Biotin (Vitamin H)
Folic acid (Vitamin M)
Riboflavin (Vitamin B2)
Thiamine (Vitamine B1)
Cobalamin (Vitamin B12)



Even more interconnections!

Few genetic paths to evolution

Repeated evolution sensu stricto



Legend

- G** repetition observed at genotypic level
- P** repetition observed at phenotypic level
- derived** derived phenotype
- ancestral** ancestral phenotype
- xxxxxx** ancestral allele

Repeated evolution

clam



garter snake

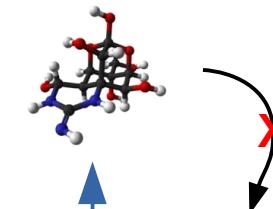


Saxitoxin



toxic plancton

Tetrodotoxin



pufferfish (fugu)

toxic newt



Bricelj 2005
Geffeney 2005
Venkatesh 2005

Repeated evolution via the same amino acid change

clam



↑ Nav1.4
X E945D

Saxitoxin



toxic plancton

garter snake



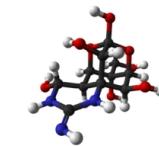
↑ Nav1.4 sodium channel
X E945D

Tetrodotoxin



toxic newt

↑ Nav1.4
E945D



pufferfish (fugu)

Repeated evolution via the same amino acid change

Locus	Mutation	Homoplastic lineages	Variation	Xenobiotic
<i>Ace-1</i>	Gly119Ser	5 (insects)	Intraspecific	Insecticides (organophosphorus)
<i>ERG3</i>	Trp205Stop	4 (yeast lines)	Experimental	Fungicide (nystatin)
<i>ERG6</i>	Gln44Stop	3 (yeast lines)		
	Gly127Arg	4 (yeast lines)	Experimental	Fungicide (nystatin)
	Tyr223Stop	4 (yeast lines)		
<i>Esterase E3</i>	Gly137Asp	3 (flies)	Intraspecific	Insecticides (diazinon)
	Trp251Leu/Ser	2 (blowflies)		Insecticides (malathion)
<i>Na,K-ATPase α</i>	Asn122His	5 (insects)		
	Glu111Val	3 (insects)		
	Glu111Leu	3 (insects)	Interspecific	Host plant toxins (cardenolides)
	Iso315Val	2 (insects)		
	Thr797Ala	2 (insects)		
<i>Nav1.4 channel</i>	Glu945Asp	1 (pufferfish)	Interspecific	Endogenous toxin (tetrodotoxin)
	Glu945Asp	1 (snake)	Interspecific	Salamander toxin (tetrodotoxin)
	Glu945Asp	1 (bivalve mollusk)	Intraspecific	Plankton toxin (saxitoxin)
<i>para (kdr)</i>	Leu1014His	2 (insects)		
	Leu1014Phe	11 (insects)		
	Leu1014Ser	2 (mosquitoes)	Intraspecific	Insecticides (pyrethroids)
	Met918Thr	5 (insects)		
	Thr929Ile	3 (2 moths, 1 louse)		
<i>Rdl</i>	Ala302Gly	3 (insects)	Intraspecific	Insecticides (cyclodienes)
	Ala302Ser	11 (insects)		
<i>Vkorc1</i>	Leu128Ser/Gln	3 (rodents)	Intraspecific	Pesticide (warfarin)
	Tyr139Cys	2 (rodents)		

Resistance to xenobiotics

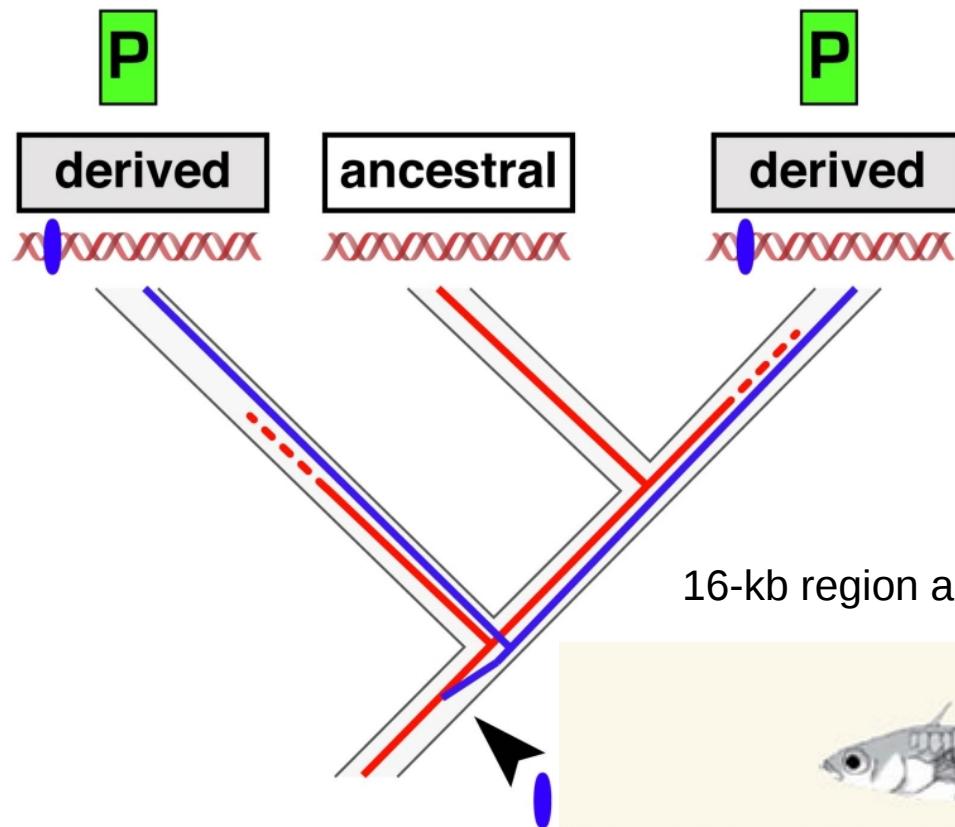
Repeated evolution



Also in:
Humans
Horses
Quails
Chickens
Mice
Pigeons

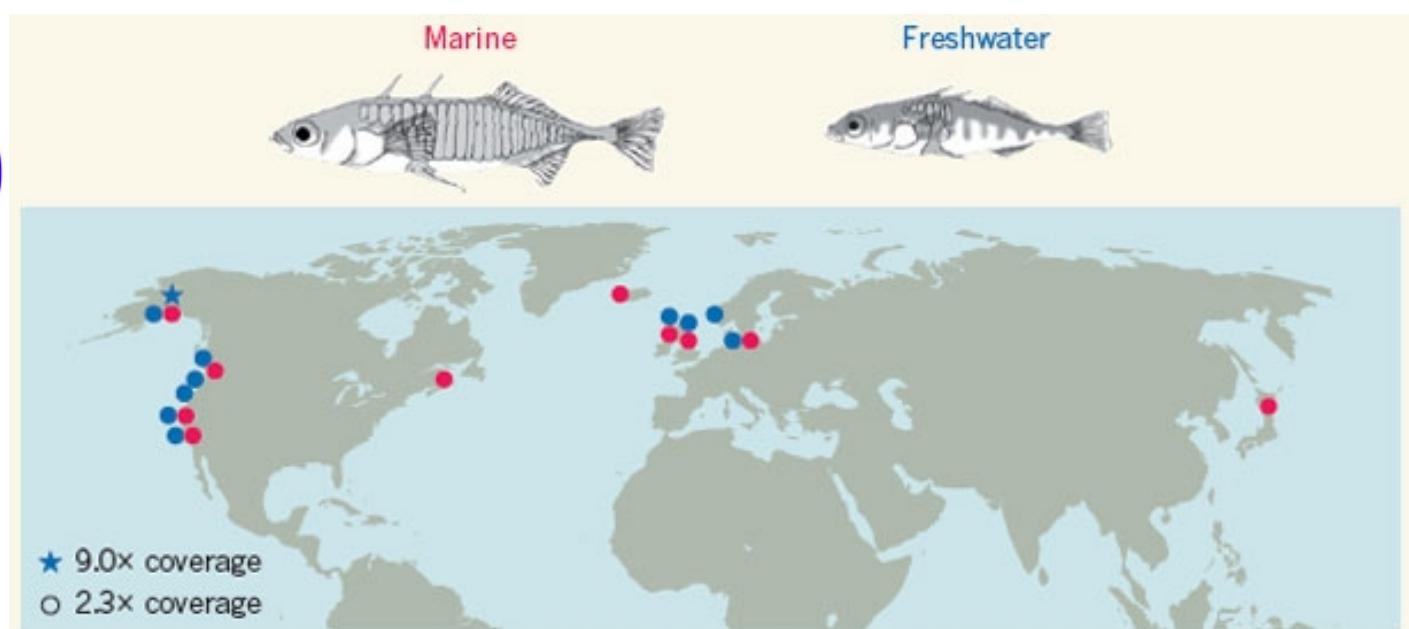


Repeated evolution via ancestral polymorphisms

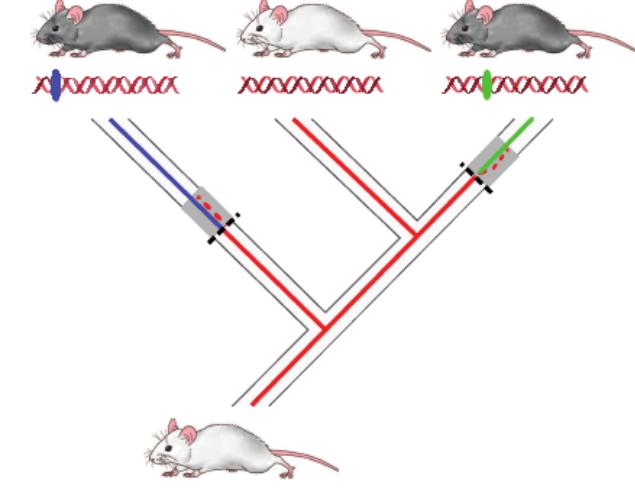


Incomplete lineage sorting
Standing genetic variation

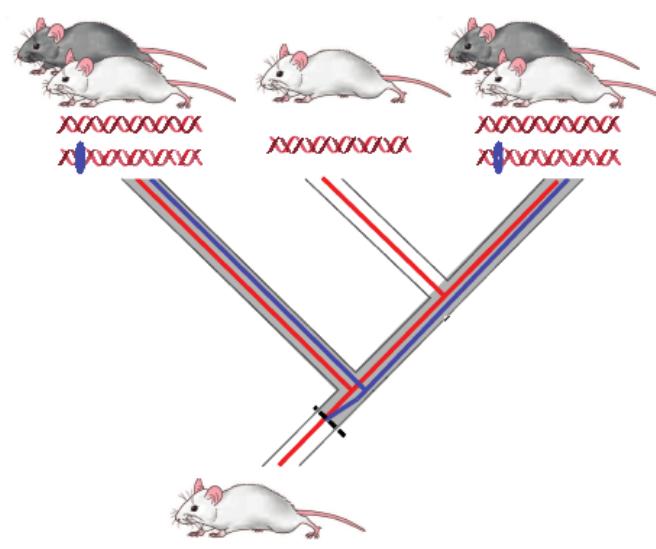
16-kb region around *Eda* shared by all freshwater fishes



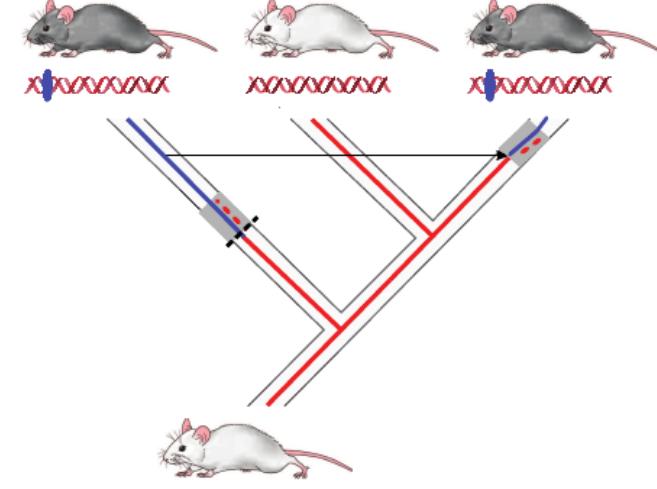
Armor plate
(Colosimo 2005
Jones 2012)



Genetic convergence

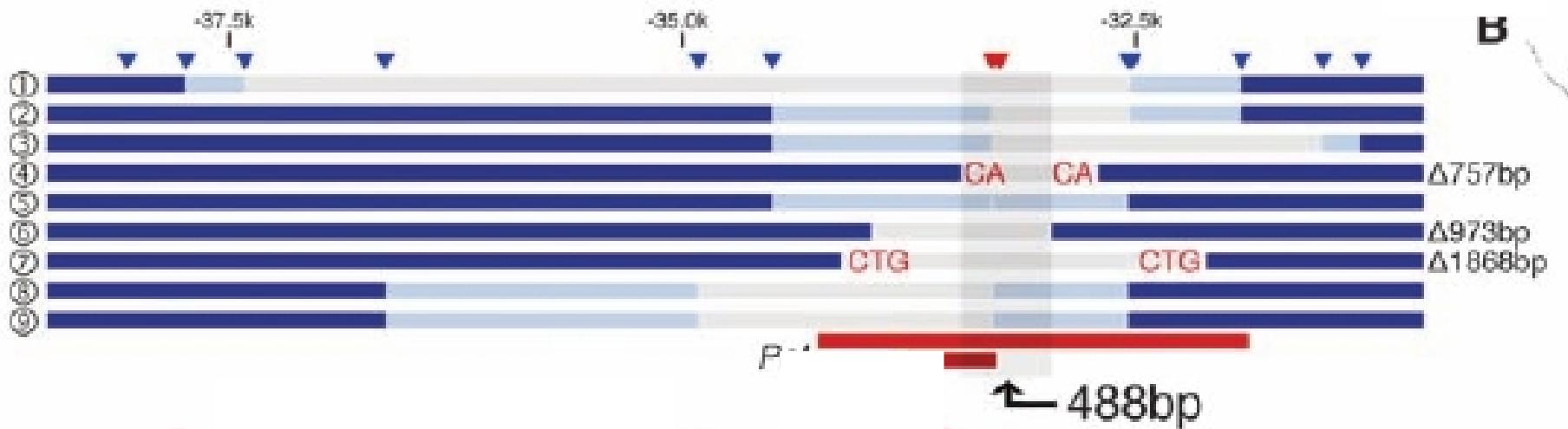
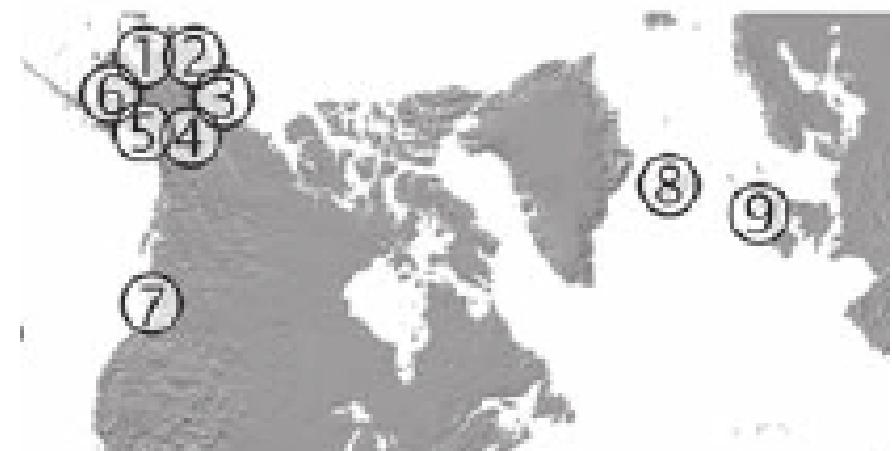
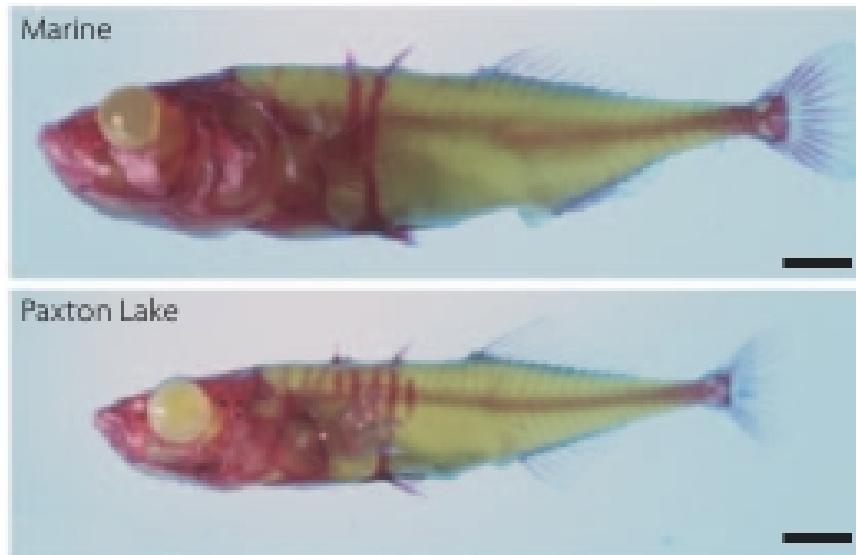


Ancestral polymorphism

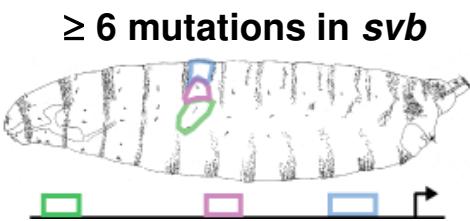


Lateral transfer

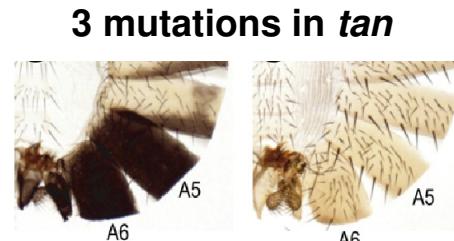
9 independent deletions in the cis-regulatory region of *Pitx1*



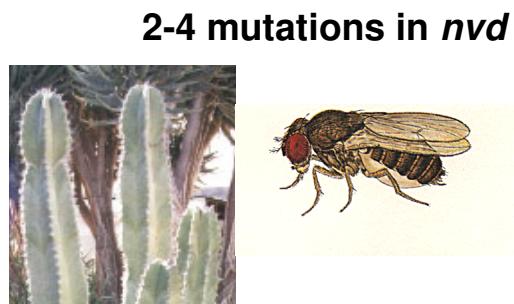
Accumulation of evolutionary-relevant mutations at the same locus



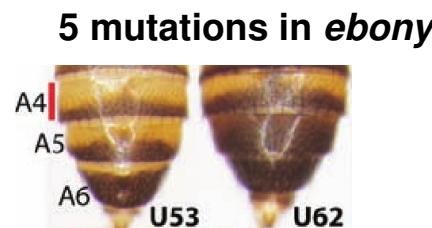
(McGregor, Orgogozo et al. 2007 Nature)



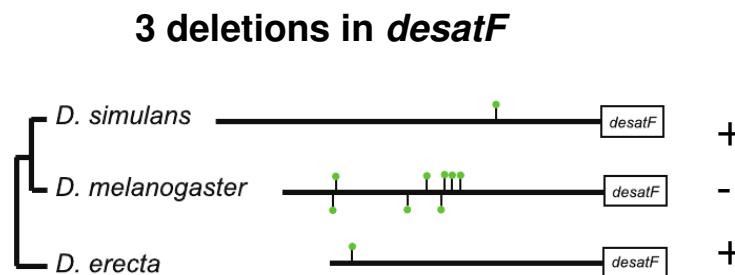
(Jeong et al., 2008 Cell)



(Lang et al. 2012 Science)

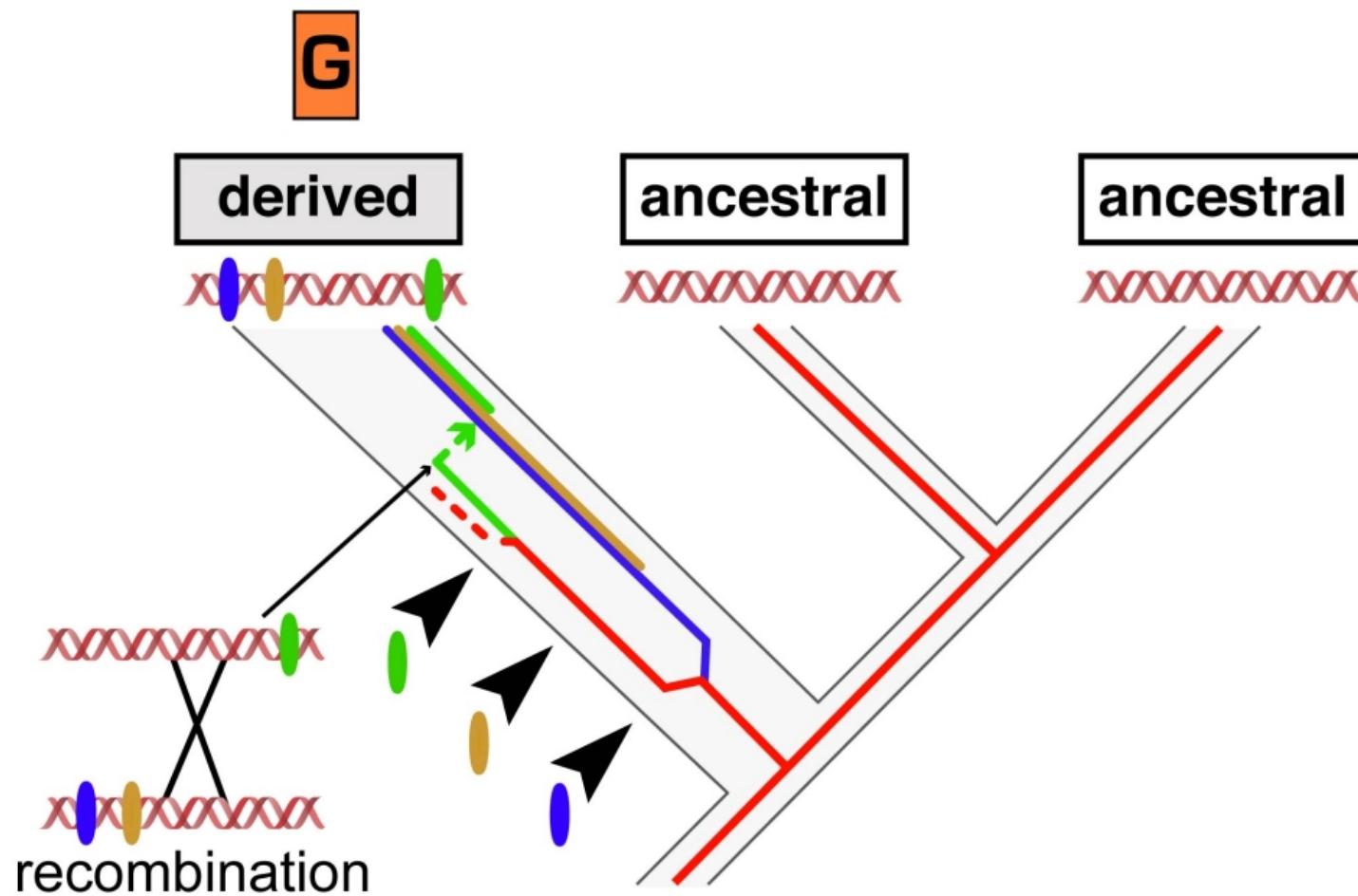


(Rebeiz et al., 2009 Cell)



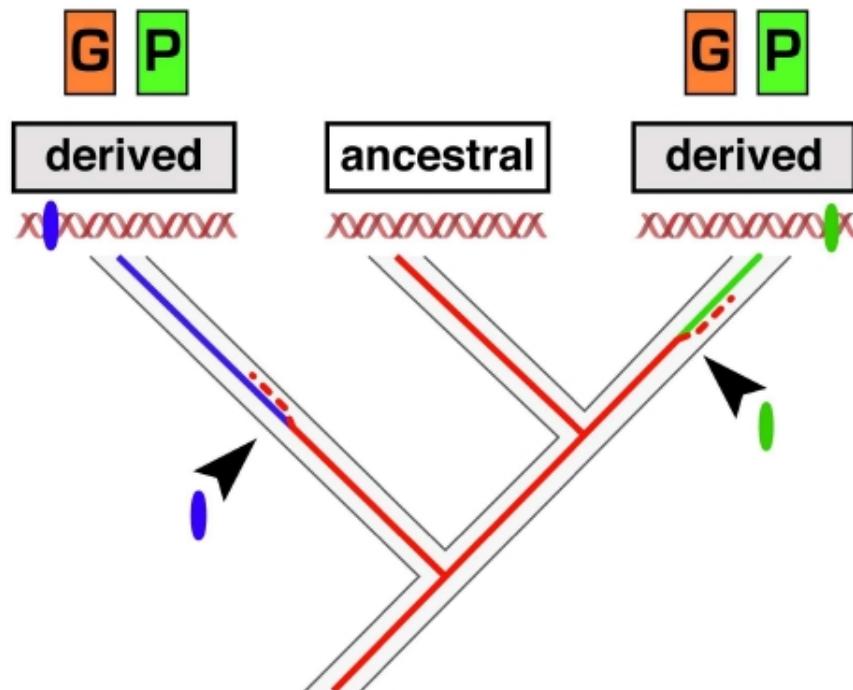
(Shirangi et al., 2009 PloS Biol)

Intralineage hotspot

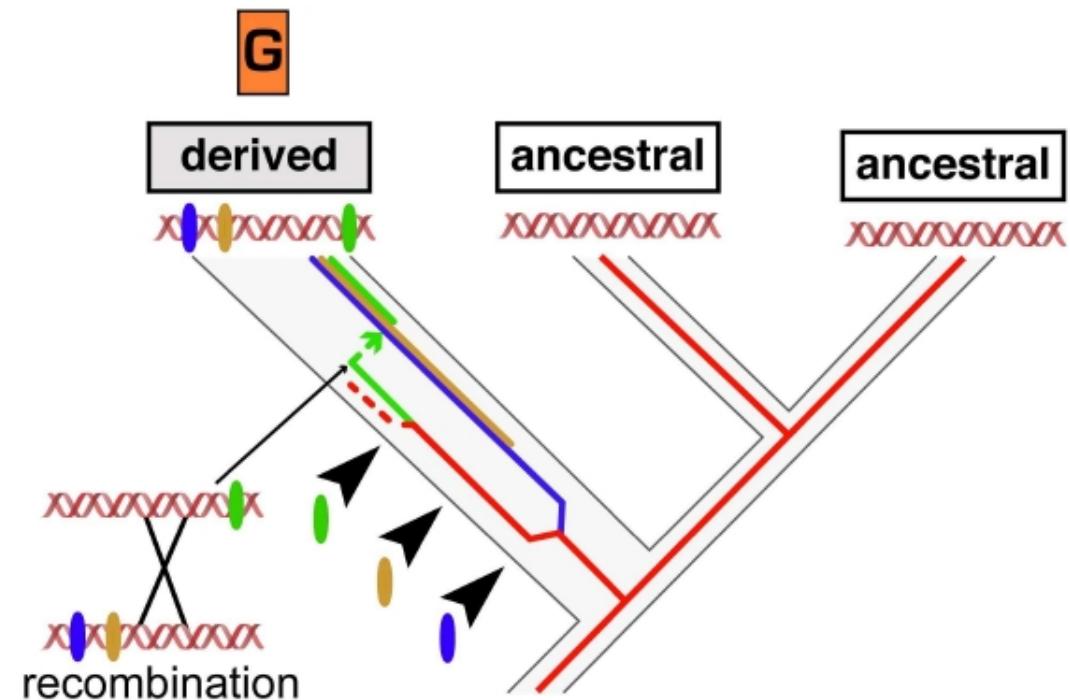


Hotspot genes: preferred targets of evolution

Interlineage hotspot



Intralineage hotspot



Why is the set of genetic paths limited?

There are specialized genes in a genome

Steroid hormone biosynthesis

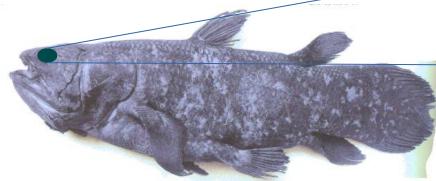


a specialized tissue
specialized enzymes

2-4 mutations in *nvd*



Color vision



a specialized tissue
specialized molecules

mutations in *opsin*
genes

Hypoxia resistance



a specialized tissue
specialized molecules

mutations in
haemoglobin genes



McCracken
2009

Specialized genes are usually genes that interact
with external parameters

Why is the set of genetic paths limited?

genes with specialized functions

But what about phenotypes involving multifunctional genes?

Evolution appears to use a restricted set of all possible paths

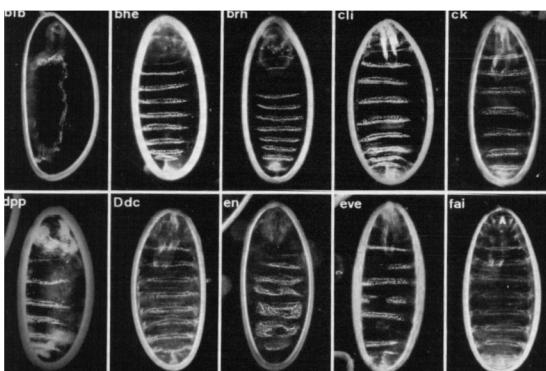
Changes in trichome pattern

EVOLUTION

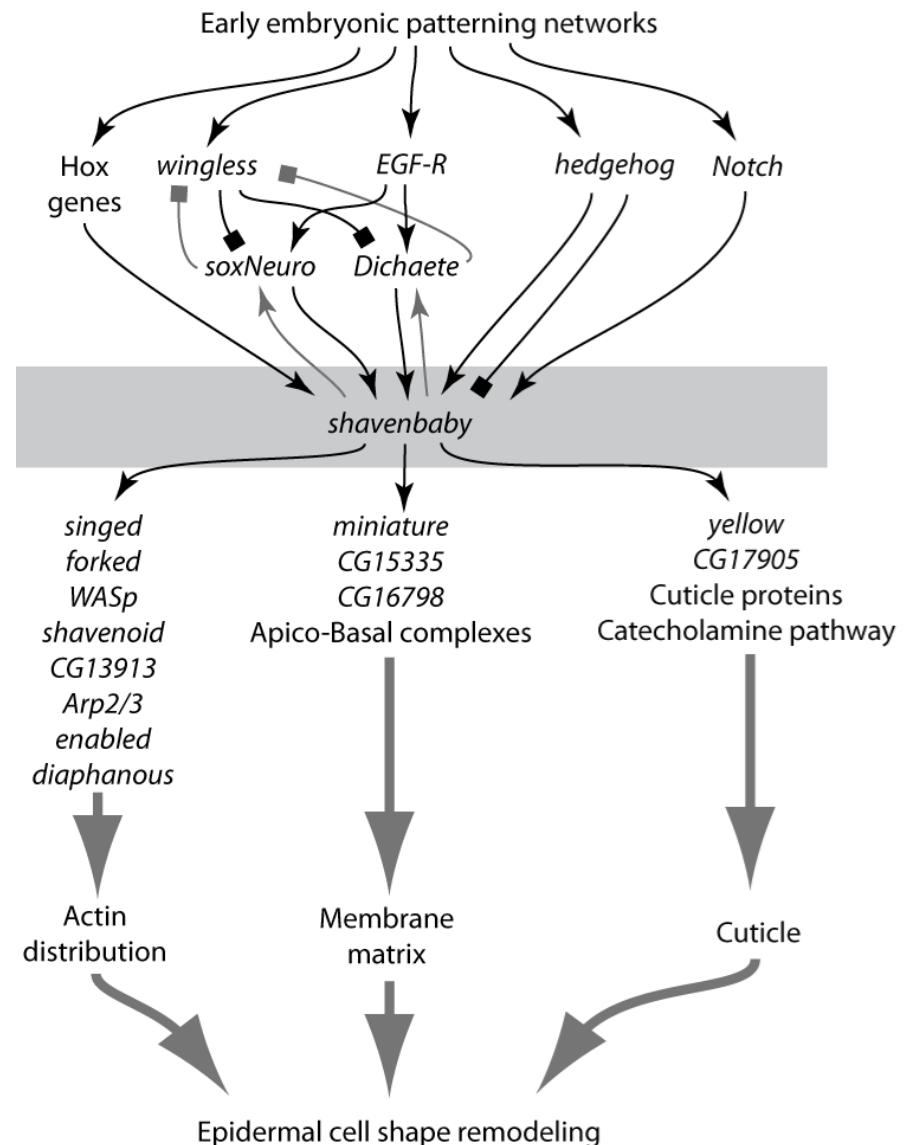


A single gene

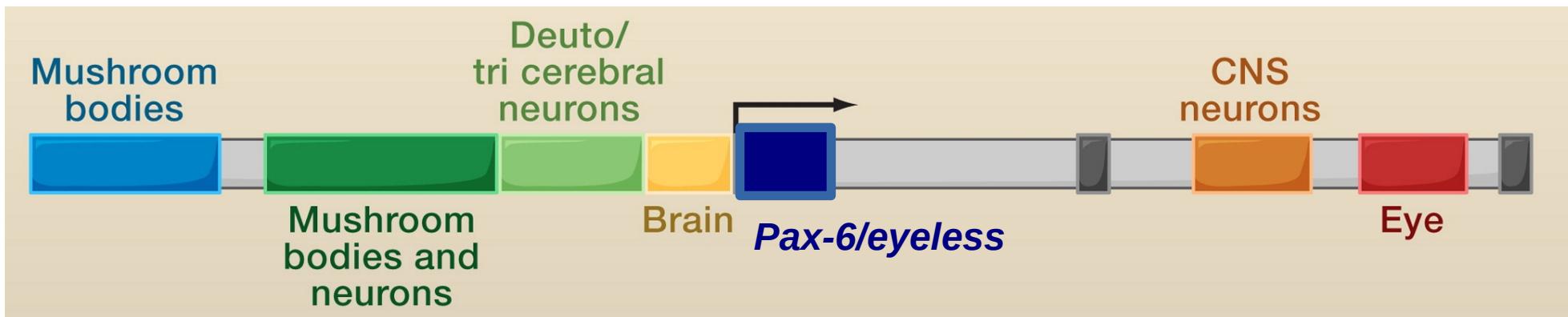
MUTAGENESIS



~100 genes

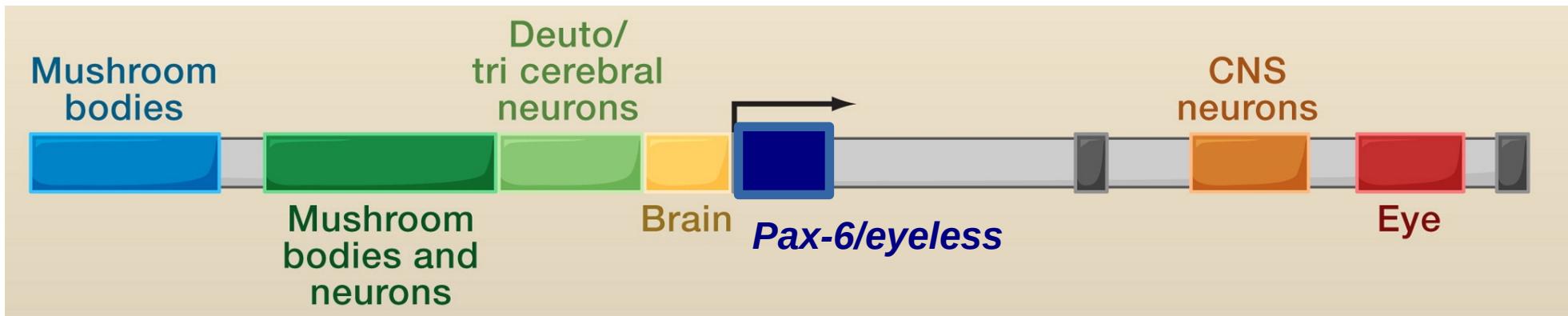


Specialized loci within multifunctional genes



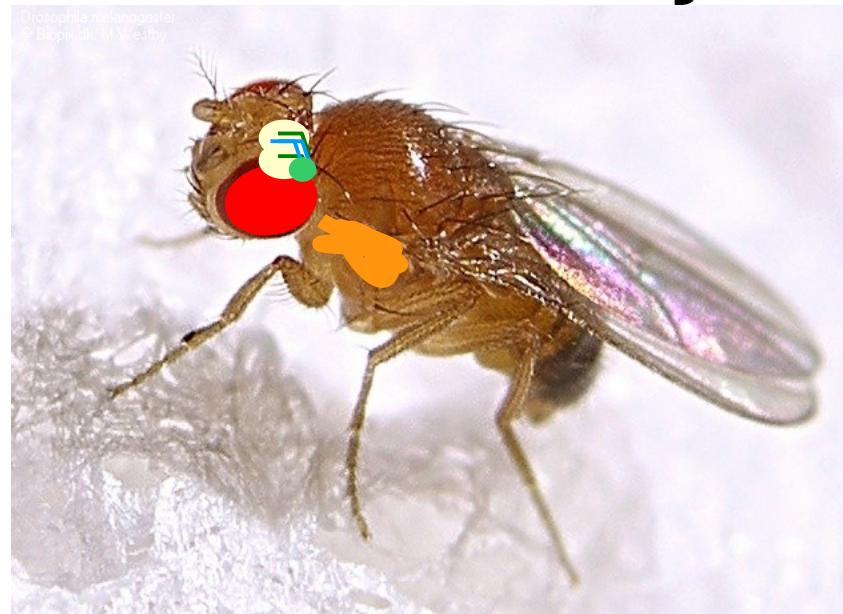
Carroll 2008

Specialized loci within multifunctional genes



Carroll 2008

Modularity of cis-regulatory elements
is reflected in modularity of body parts



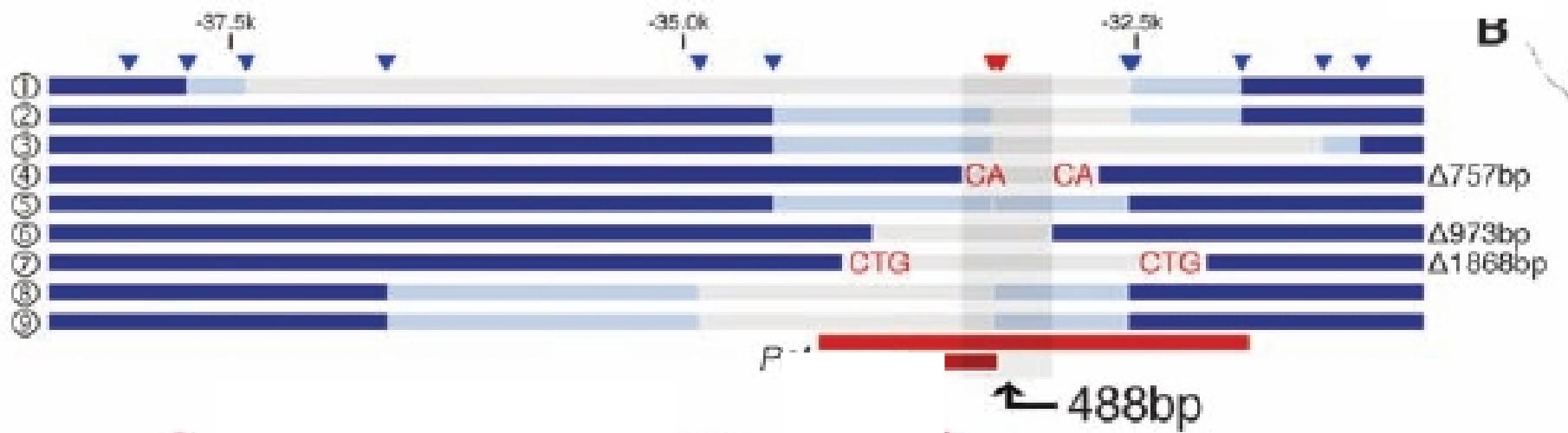
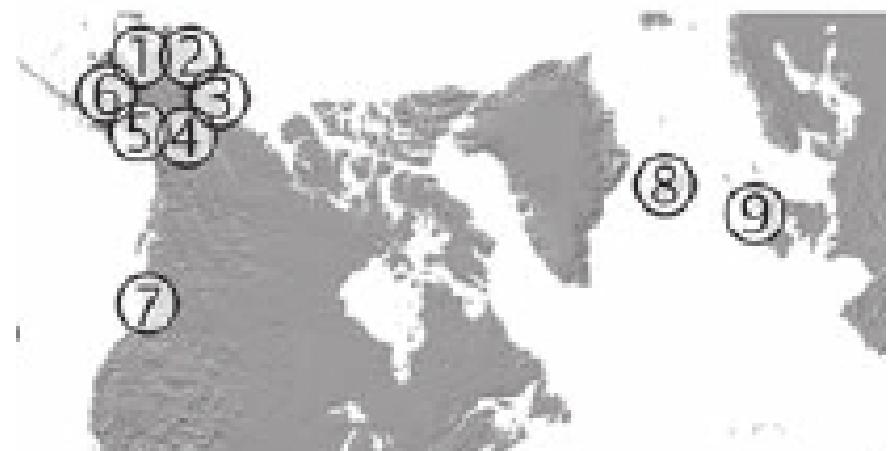
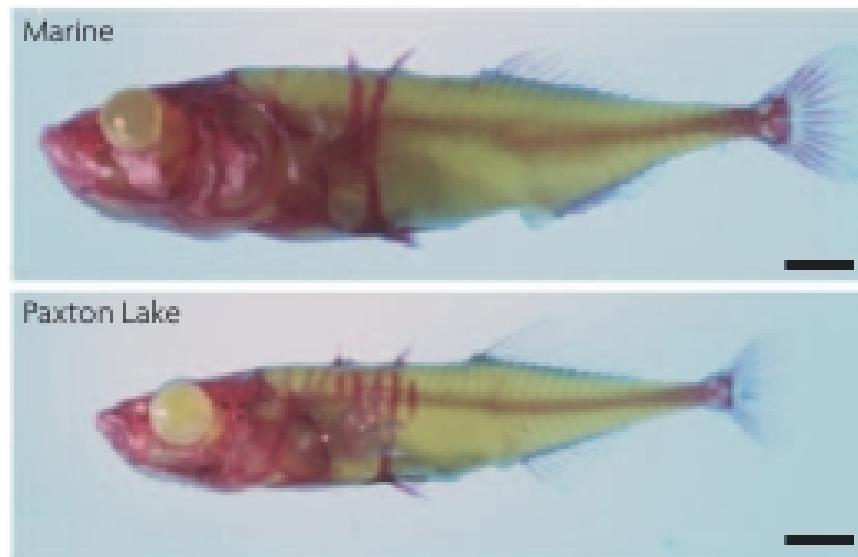
Why is the set of genetic paths limited?

genes with specialized functions

multifunctional genes with specialized regions

mutational bias

9 deletions in the cis-regulatory region of *Pitx1* due to region sensitive to chromosome breaks



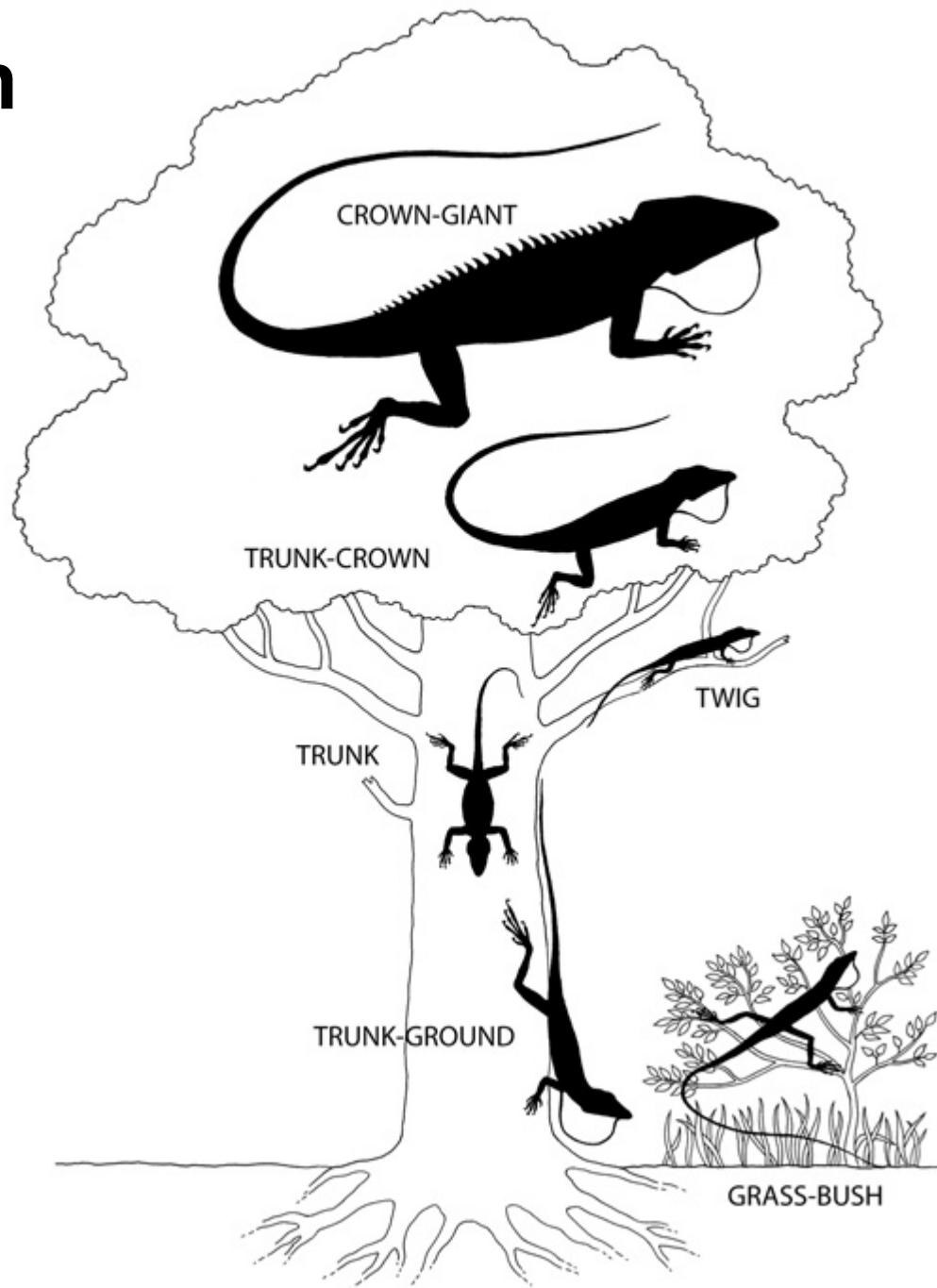
Why is the set of genetic paths limited?

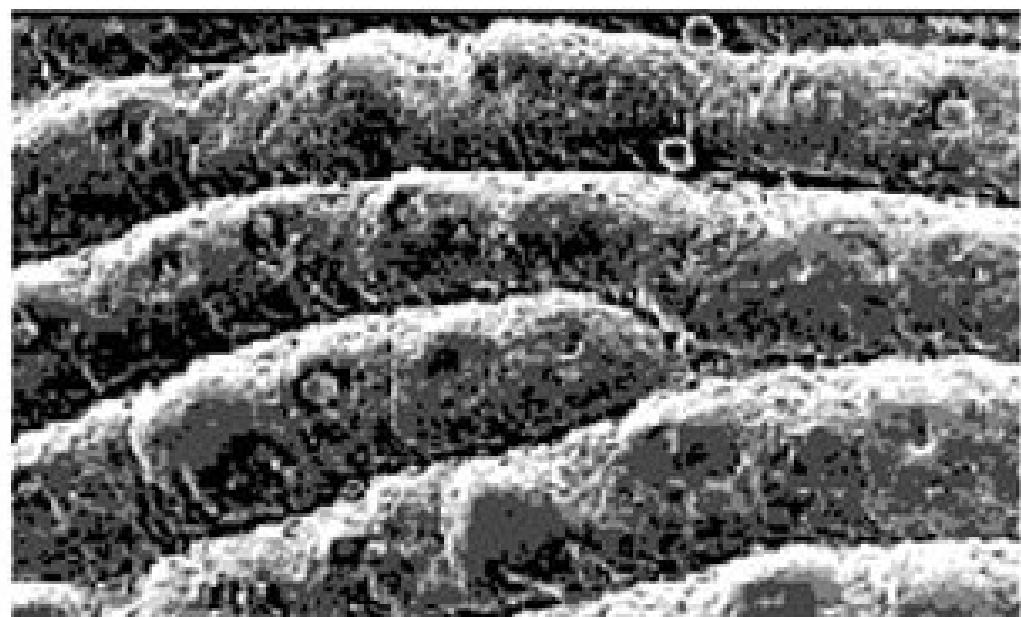
genes with specialized functions

multifunctional genes with specialized regions

mutational bias

Repeated evolution in diverse Caribbean islands





Evolution repeats itself in..



.. the genes responsible for natural evolution

Ex : *hemoglobin* in dogs and humans in Tibet
(Wang et al 2014 GBE)



.. the genes responsible for experimental evolution

Ex : *sulfate transporter SUL1* in yeasts in low sulfate
(Gresham et al 2008 PloS Genetics)

.. the phenotypes evolving in certain environments

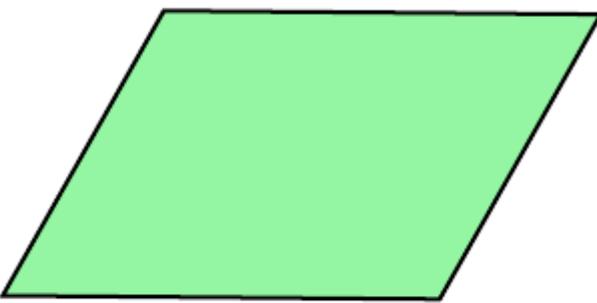


Ex : flying marsupial phalanger and placental flying squirrel

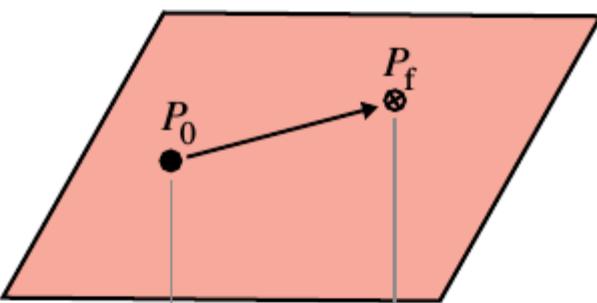
Orgogozo 2015 Interface Focus

Synchronous predictability

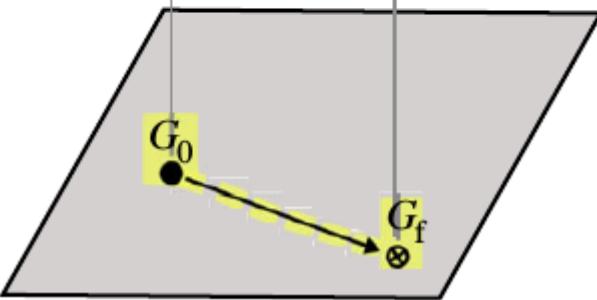
Natural Evolution



environment
space

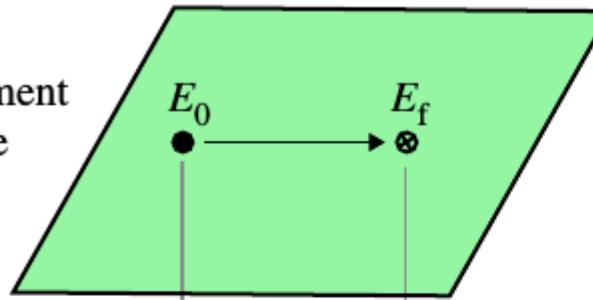


phenotype
space

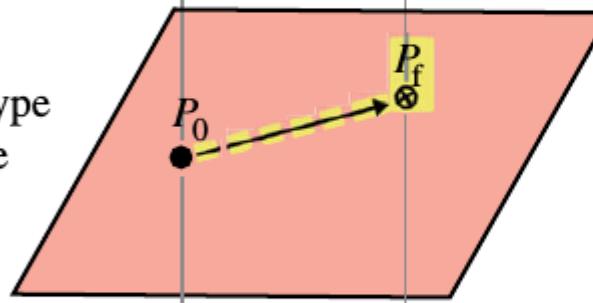


genotype
space

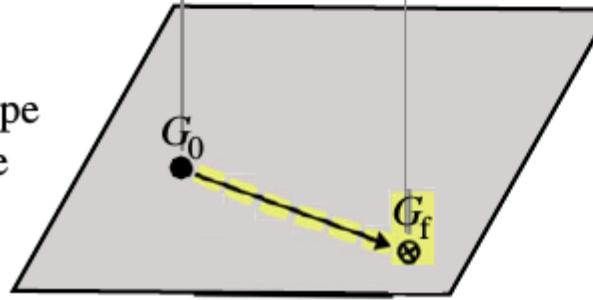
Experimental Evolution



E_0 → \otimes

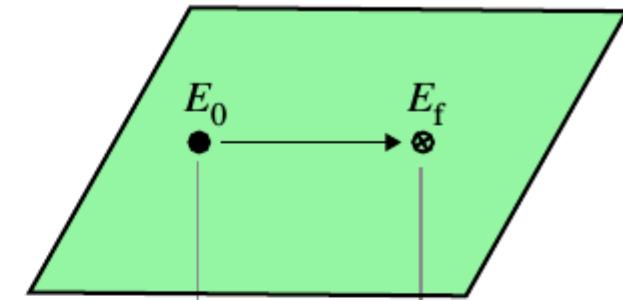


P_0 → \otimes

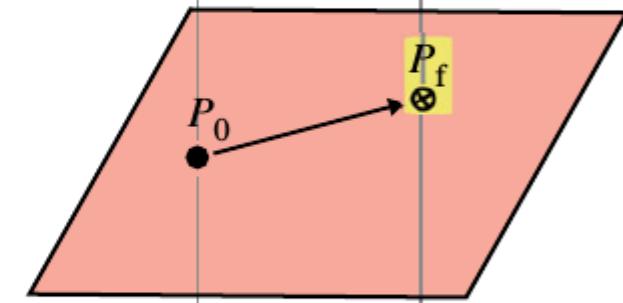


G_0 → \otimes

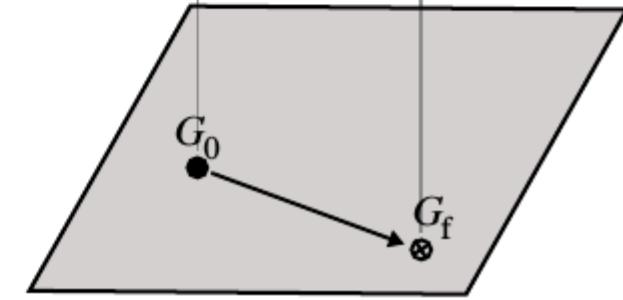
Convergent Evolution



E_0 → \otimes

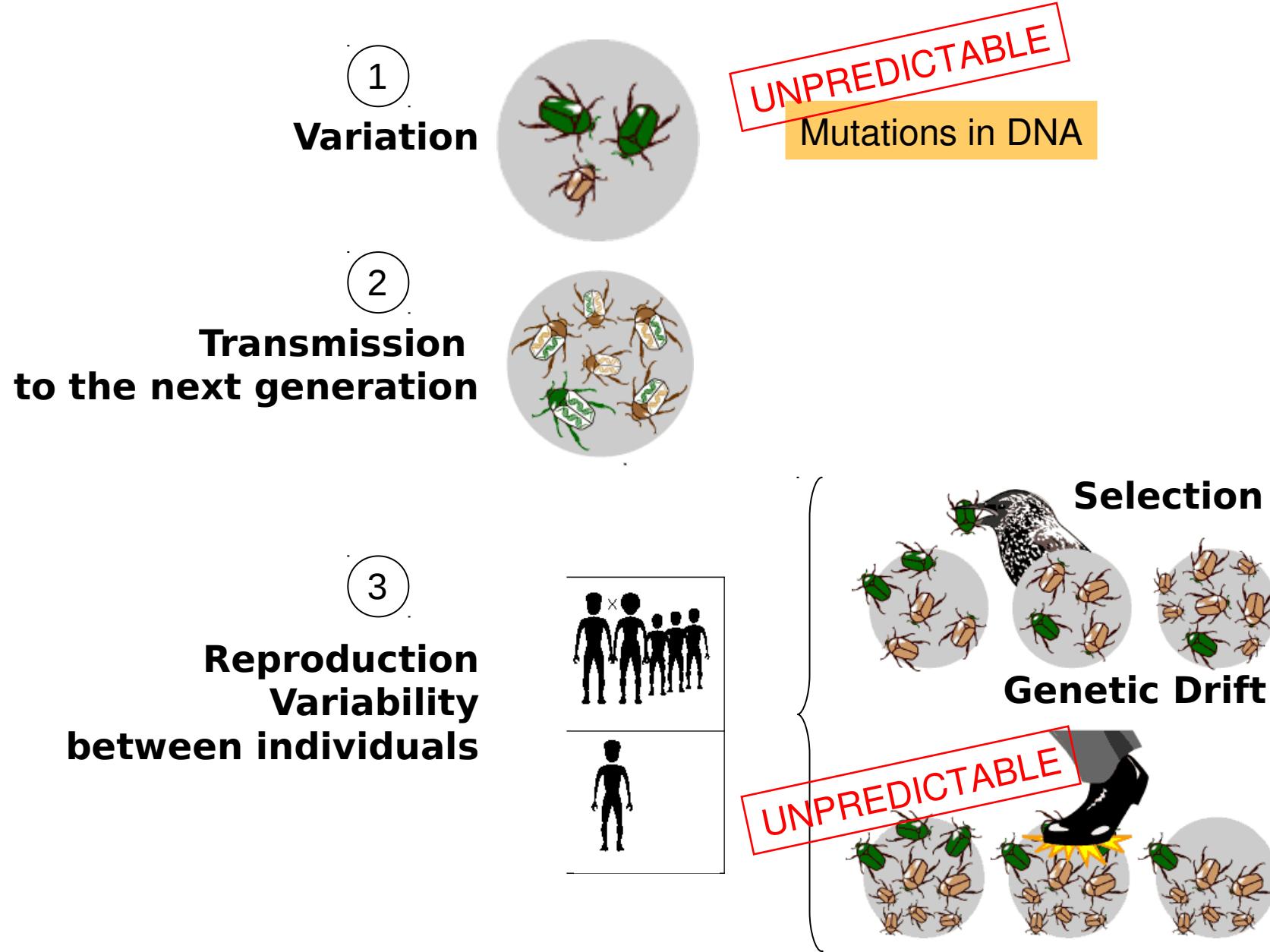


P_0 → \otimes



G_0 → \otimes

Classical Darwinian Evolution



**Many unpredictable processes
at a low level**



**Predictable Evolution
at the genetic level**

Mutations in DNA
Chromosome segregation during meiosis
Assortative mating
Gamete competition during fecondation
Life history traits
Genetic linkage
Environmental changes (meteorite, etc.)
...



From random processes can emerge predictability

Many unpredictable processes
at a low level

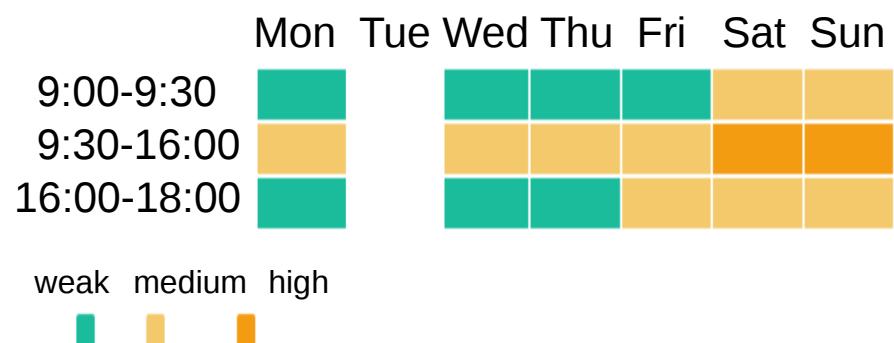


Predictable
at a higher level

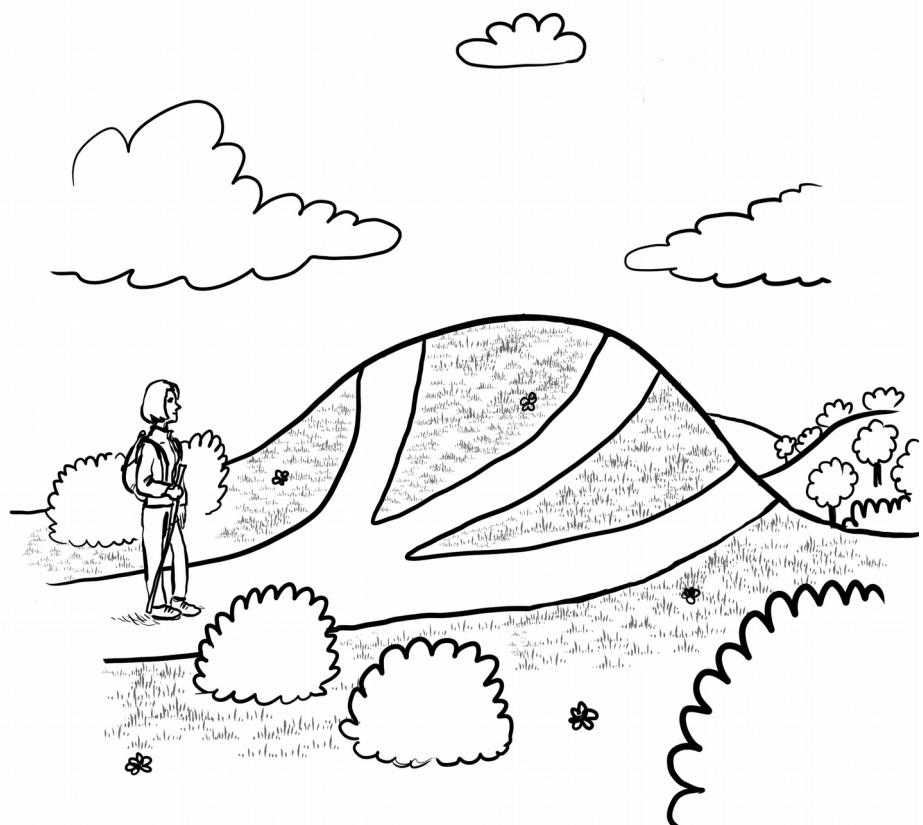
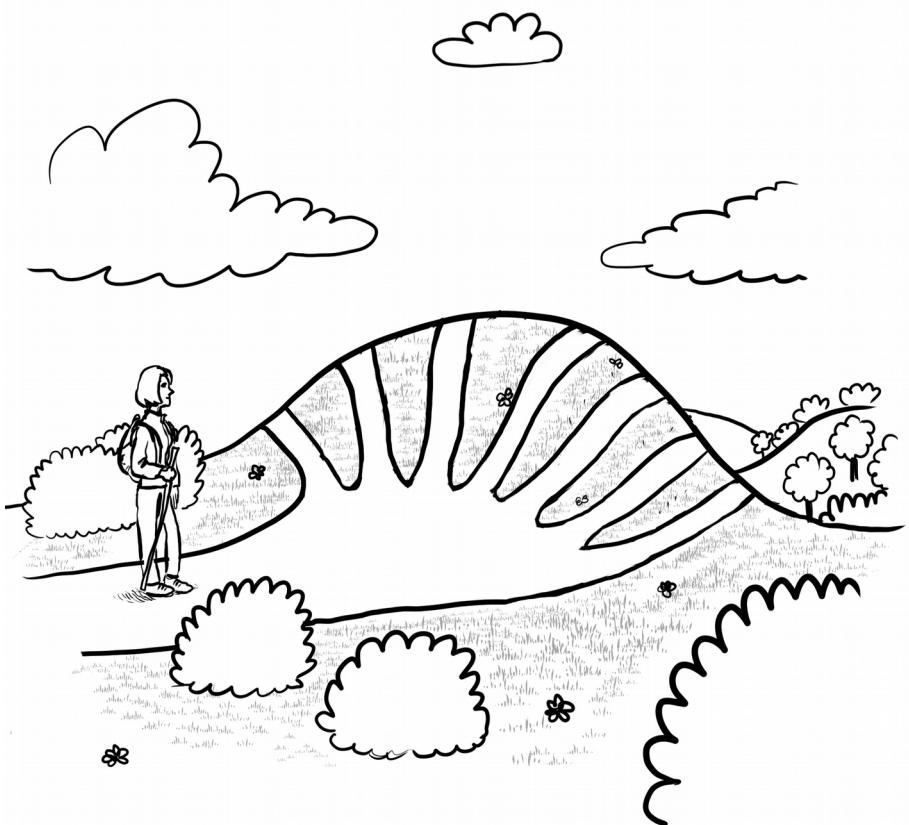
Trajectory of single individuals



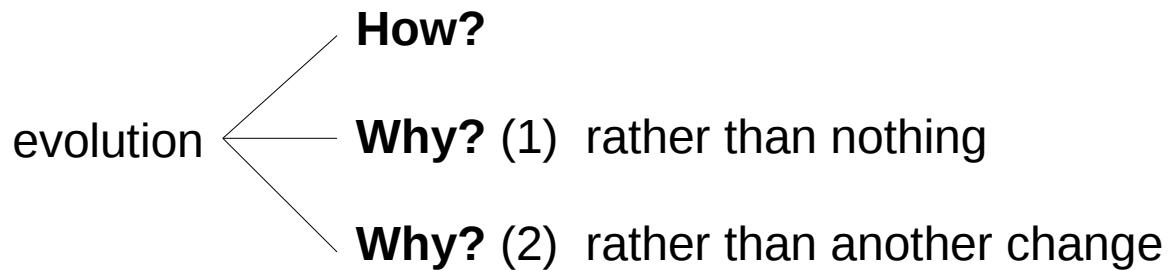
Estimated waiting time for Louvre museum



The number of possible paths for evolution is smaller than previously thought



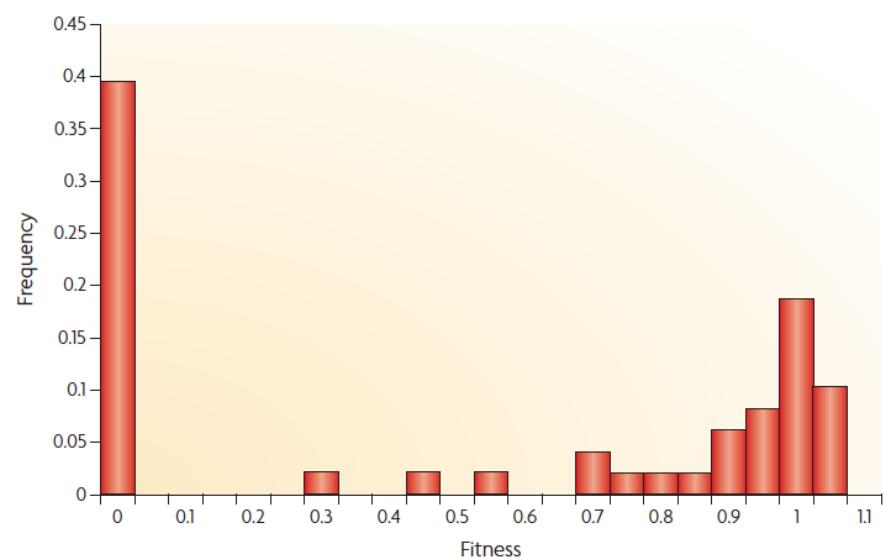
What are all the possible evolutionary paths?



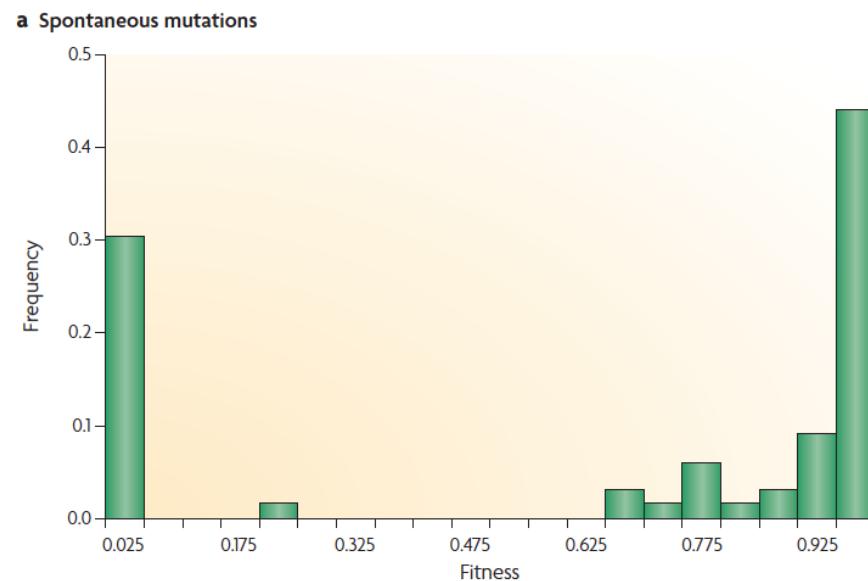
Test all possible paths:

Examine all possible mutations in a protein sequence,
in a cis-regulatory sequence, etc.

Mostly deleterious and neutral mutations



Vesicular Stomatitis Virus



Saccharomyces cerevisiae
diploids

Eyre-Walker and Keightley NRG 2007

Evolution: unconstrained and unpredictable?

[past and present organisms are] a subset of workable, but basically fortuitous, survivals among a much larger set that could have functioned just as well, but either never arose, or lost their opportunities, by historical happenstance.

Stephen Jay Gould, 2002

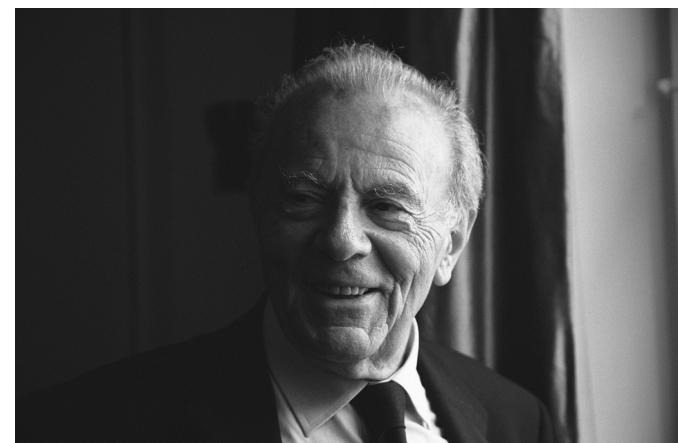


It is hard to realize that the living world as we know it is just one among many possibilities; that its actual structure results from the history of the earth.

1977

Evolution and Tinkering

François Jacob



Would life evolve again, would it produce similar living beings?



evolution

How?

Why? (1) rather than nothing

Why? (2) rather than another change

