

Relations génotype-phénotype et Evolution

Virginie COURTIER-ORGOGOZO



ROBUST

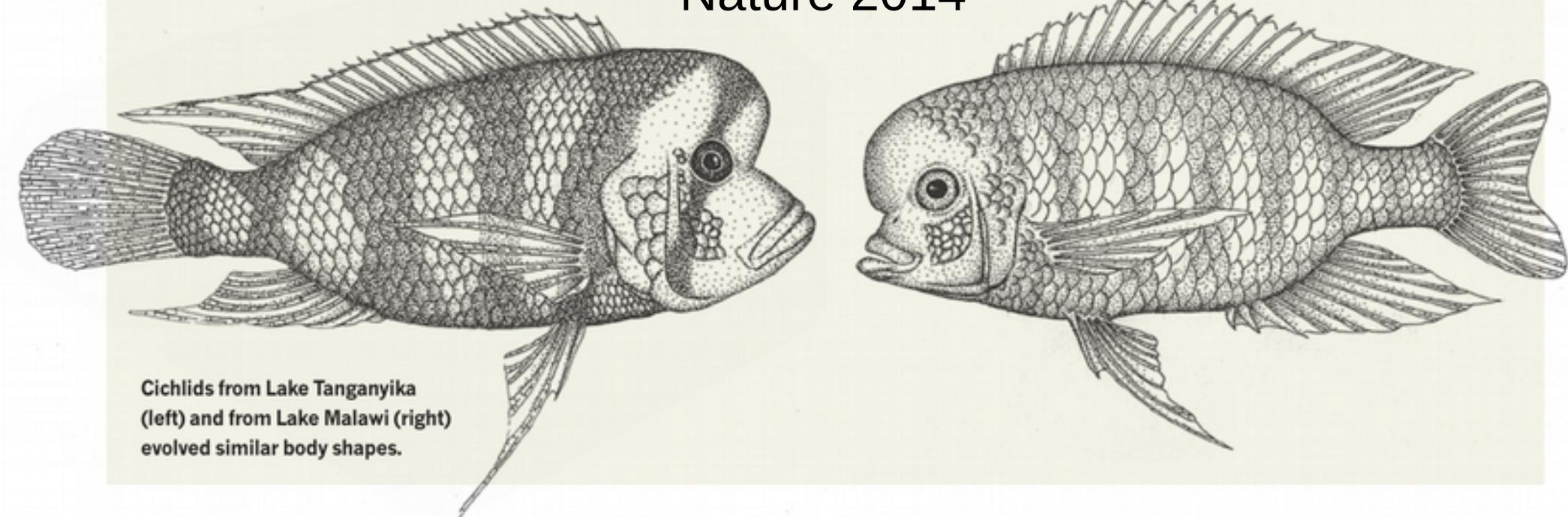


JOHN TEMPLETON
FOUNDATION



université
PARIS
DIDEROT
PARIS 7

Nature 2014



Cichlids from Lake Tanganyika
(left) and from Lake Malawi (right)
evolved similar body shapes.

Does evolutionary theory need a rethink?

Researchers are divided over what processes should be considered fundamental.

POINT

Yes, urgently

Without an extended evolutionary framework, the theory neglects key processes, say Kevin Laland and colleagues.

COUNTERPOINT

No, all is well

Theory accommodates evidence through relentless synthesis, say Gregory A. Wray, Hopi E. Hoekstra and colleagues.

Didier Raoult dénigre la théorie de l'évolution

Pr Didier Raoult

Dépasser Darwin

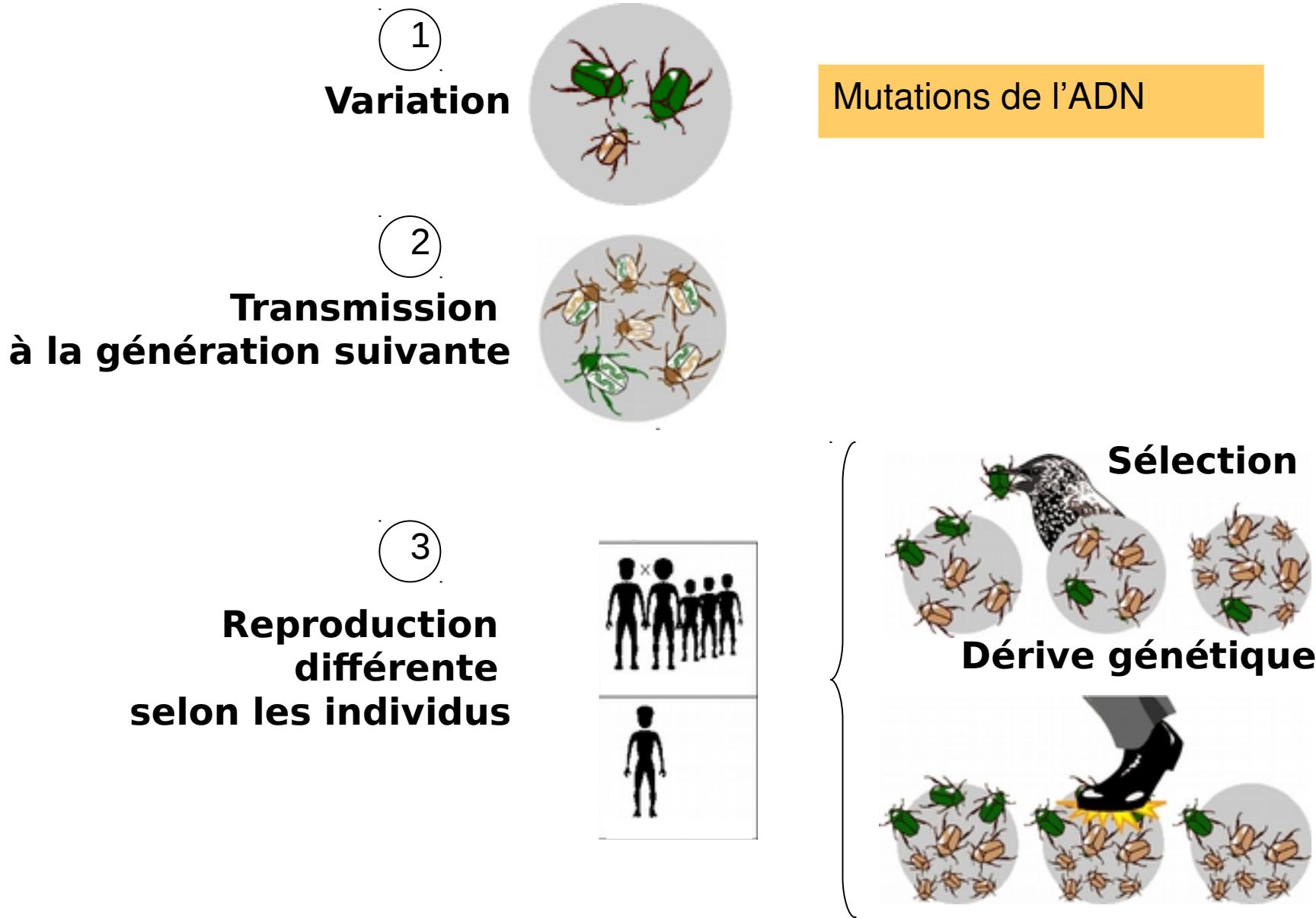
L'évolution comme vous ne l'aviez jamais imaginée

PLON

Actuellement, la théorie la plus castratrice en science est celle de Charles Darwin, car elle empêche une réanalyse des données contemporaines. C'est juste une théorie de WASP (Anglo-Saxons protestants blancs), ceux qui dominaient le Royaume-Uni

Enfin, Darwin définit la sélection naturelle par analogie à la sélection humaine. Il pense que la nature agit pour trier les espèces comme les hommes l'ont fait pour le bétail, les poules, les chevaux, les chiens et les chats, choisissant les «étalons» les plus capables et ségrégant progressivement les caractères. Mais la nature ne fonctionne pas ainsi, la plupart des évolutions résultent de sélections multiples dues au hasard, les plus capables à un instant ne seront pas les plus fertiles ni les plus performants à l'avenir. Le tri se fait souvent par catastrophe. La nature se développe aussi par imitation et par adaptation à l'environnement, et cette acquisition de compétences est transmissible. C'est ce

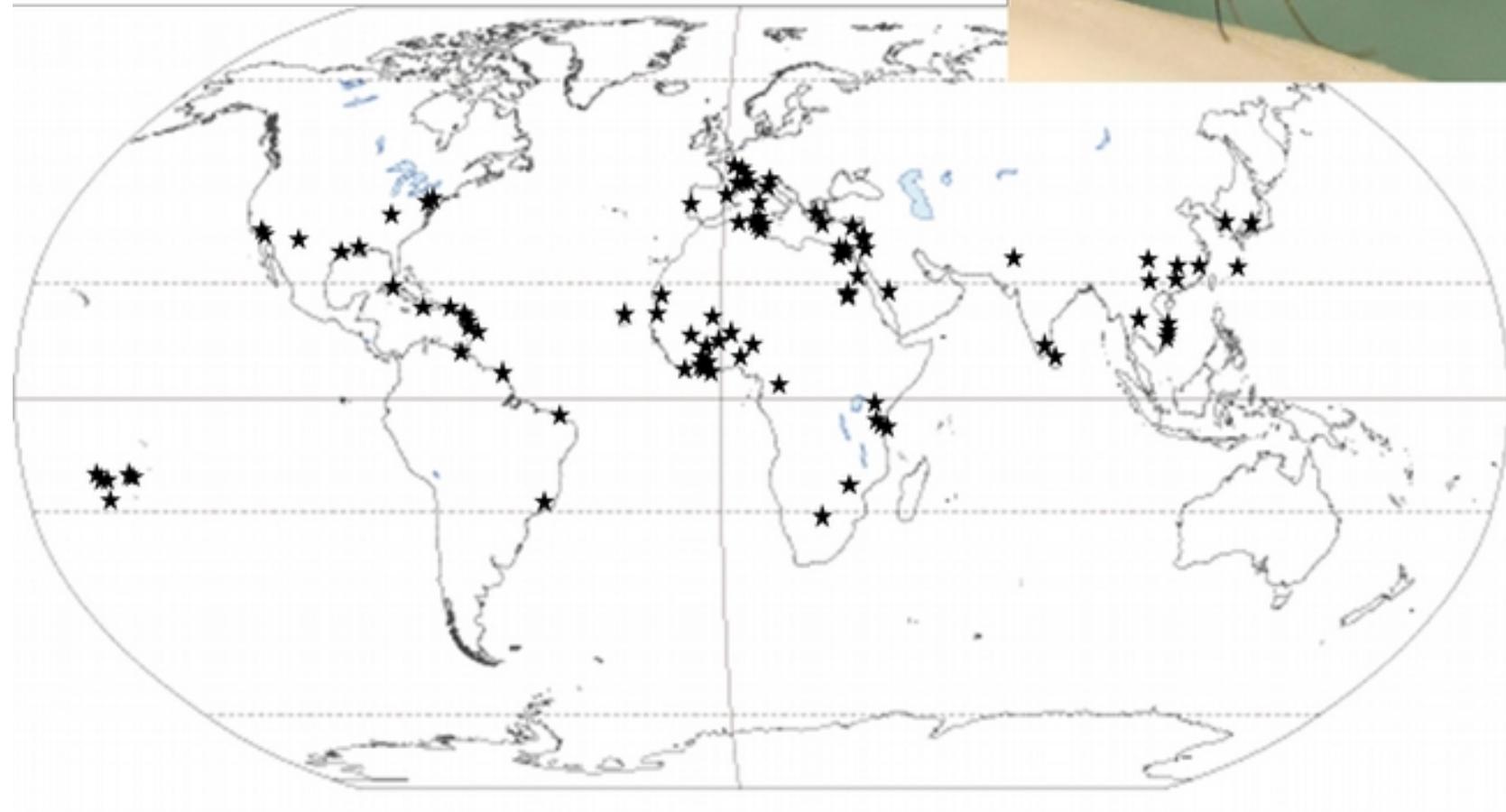
Classical Darwinian Evolution

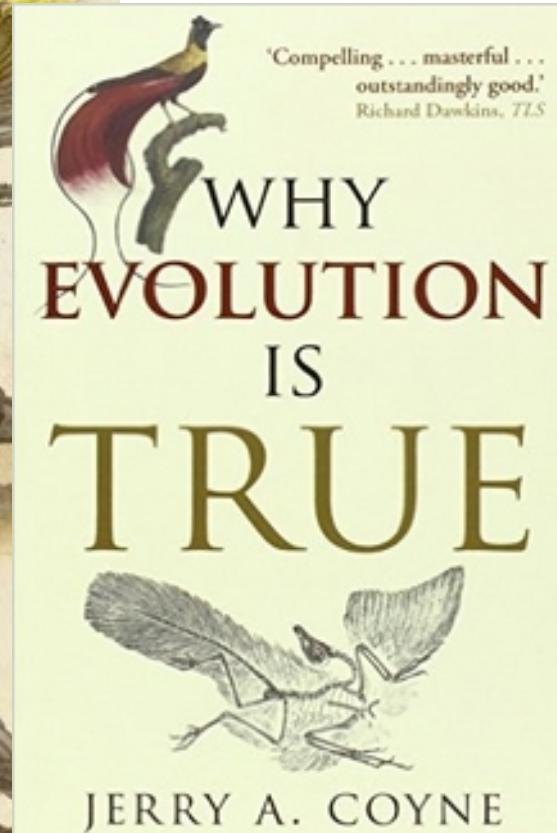
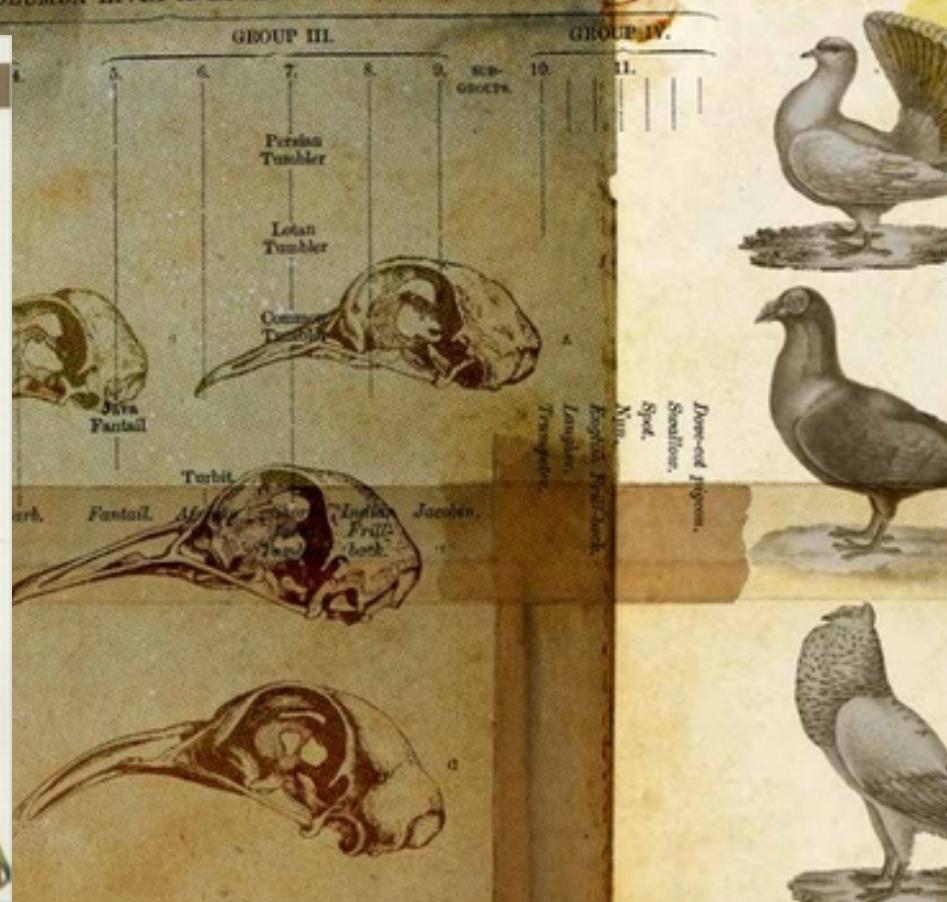
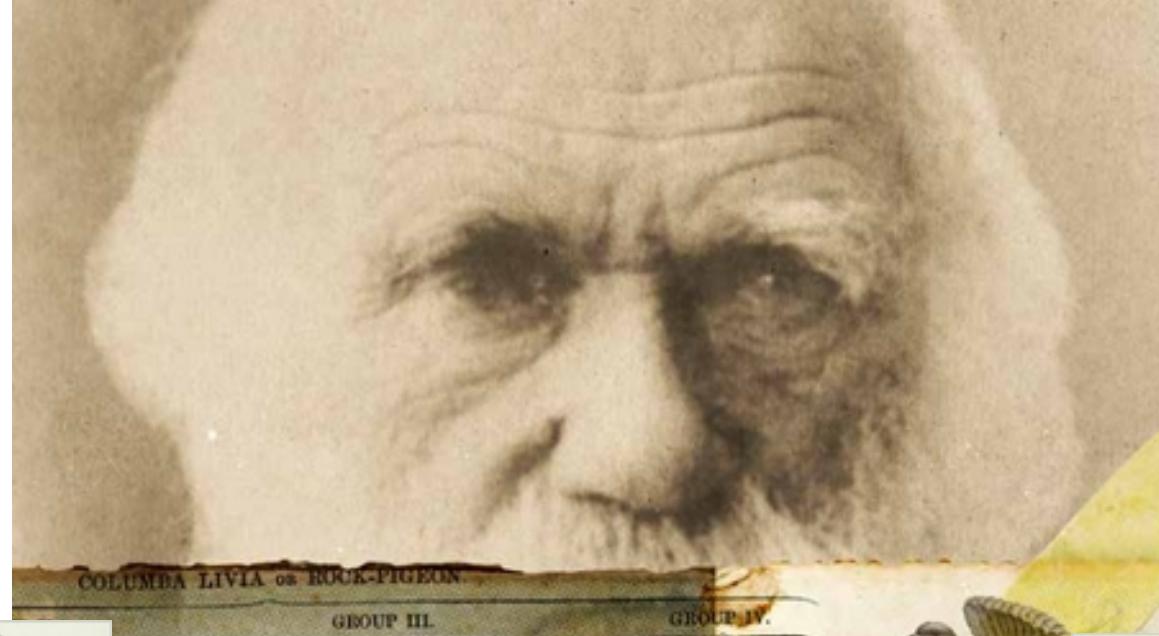
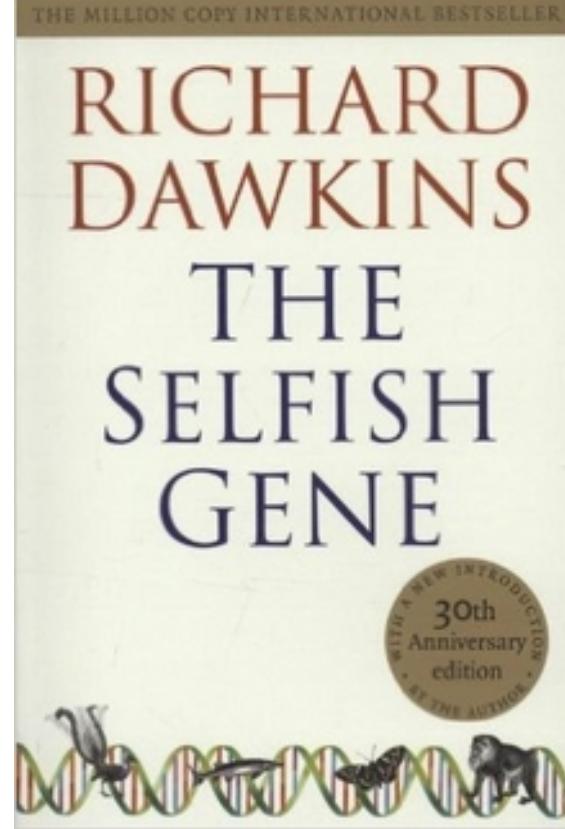


Insecticide resistance in *Culex pipiens*

Ester²

First detected in Liberia and Nigeria in 1977
and in France in 1986





Current critics

New view = EES = Extended Evolutionary Synthesis

Evolution is not random:

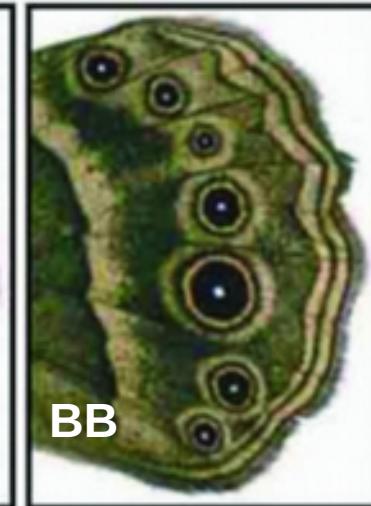
- physical development influences the generation of variation (developmental bias)
- the environment directly shapes organisms' traits (plasticity)
- organisms modify environments (niche construction)
- organisms transmit more than genes across generations (extra-genetic inheritance)

Developmental bias

Selection for bigger/smaller spots

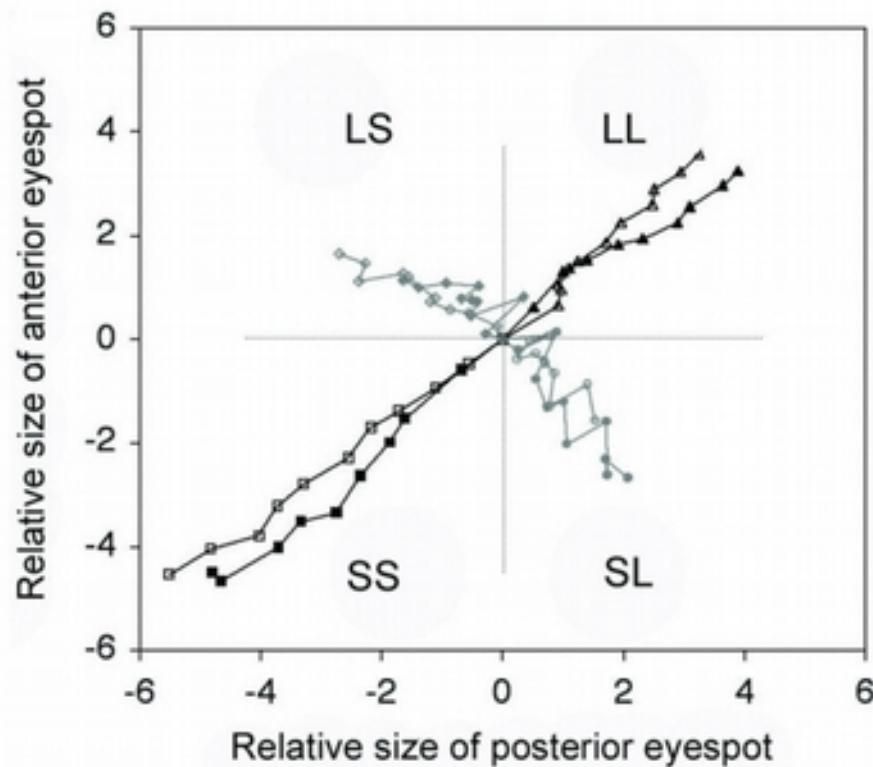


Selection for more black/more gold



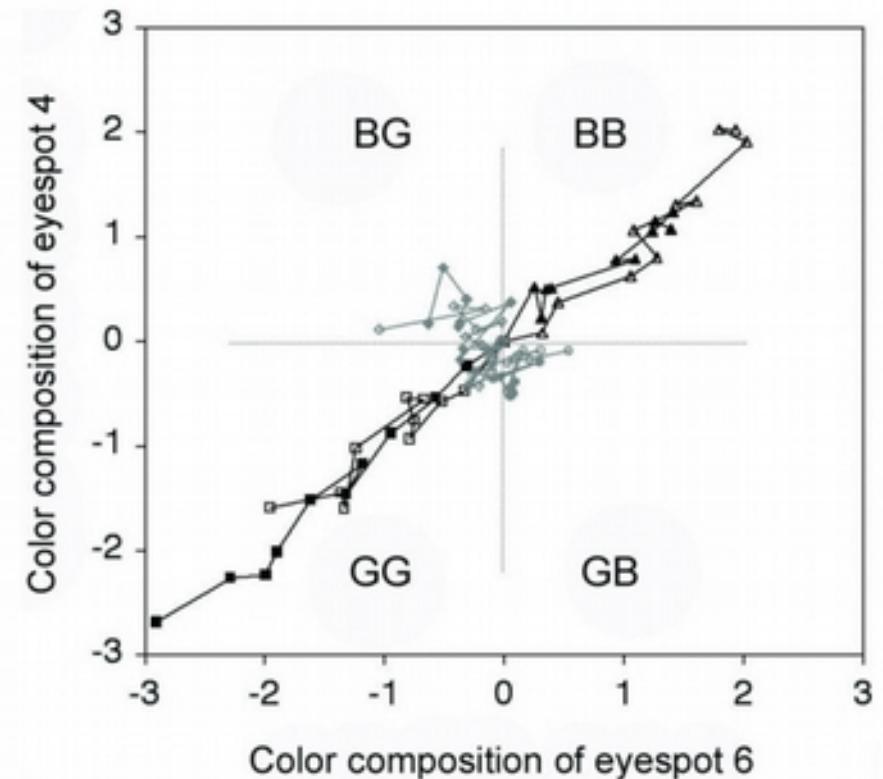
Developmental bias

Eyespot size



L = Large
S = Small

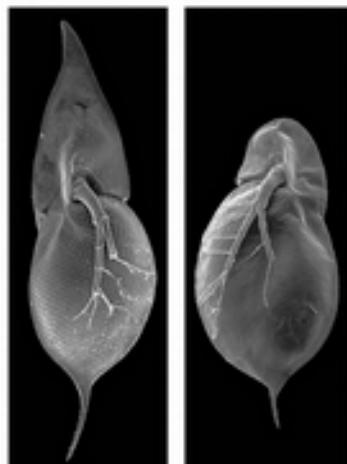
Eyespot colour composition



B = Black
G = Gold

Plasticity

Daphnia



with helmet without helmet

Nemoria arizonaria caterpillars



spring: caterpillars feed on catkins

summer: caterpillars feed on leaves

Water crowfoot plant



leaves growing above water

leaves growing below water

Desert locusts



solitary



gregarious

Commodore butterfly



winter



summer

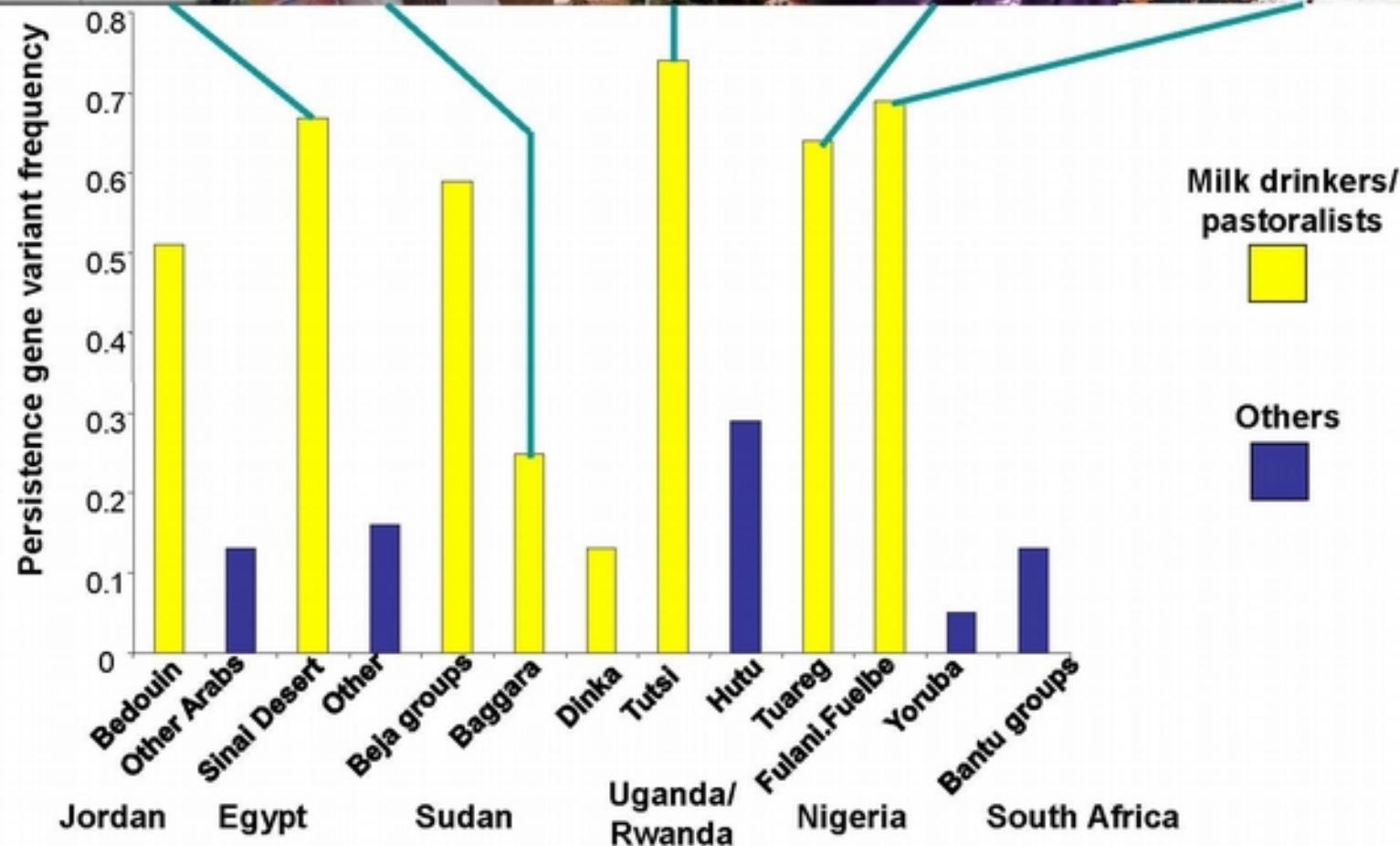
Commodore butterfly: Michael Wild, CC-BY-SA-3.0 (winter), Svdmolen, CC-BY-SA-3.0 (summer)

Daphnia: Agrawal et al (1999)

Nemoria arizonaria caterpillars: Sadava et al (2014)

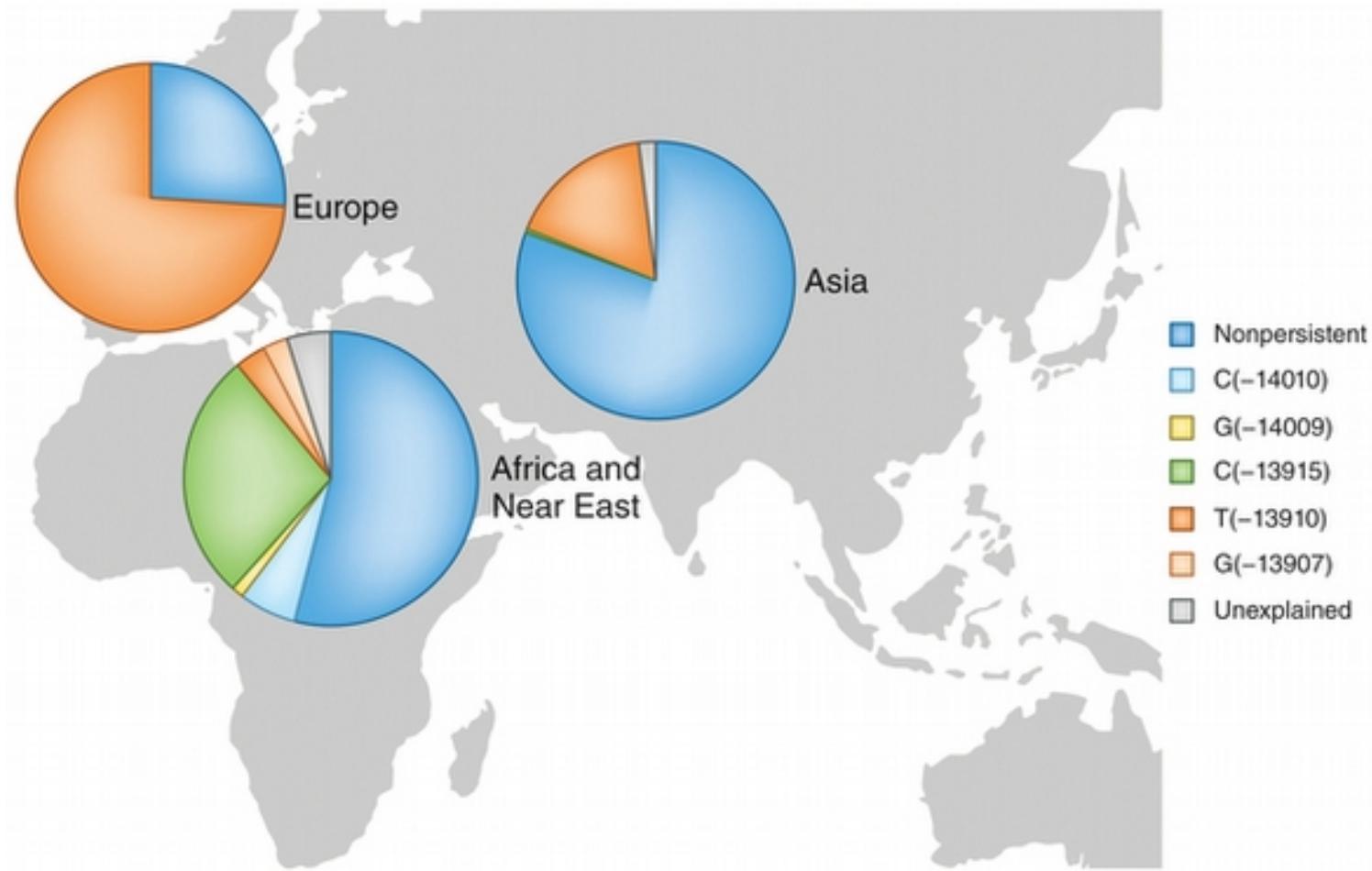
Water crowfoot plant: J R Crellin, CC BY-NC-ND 3.0

Niche construction



Niche construction

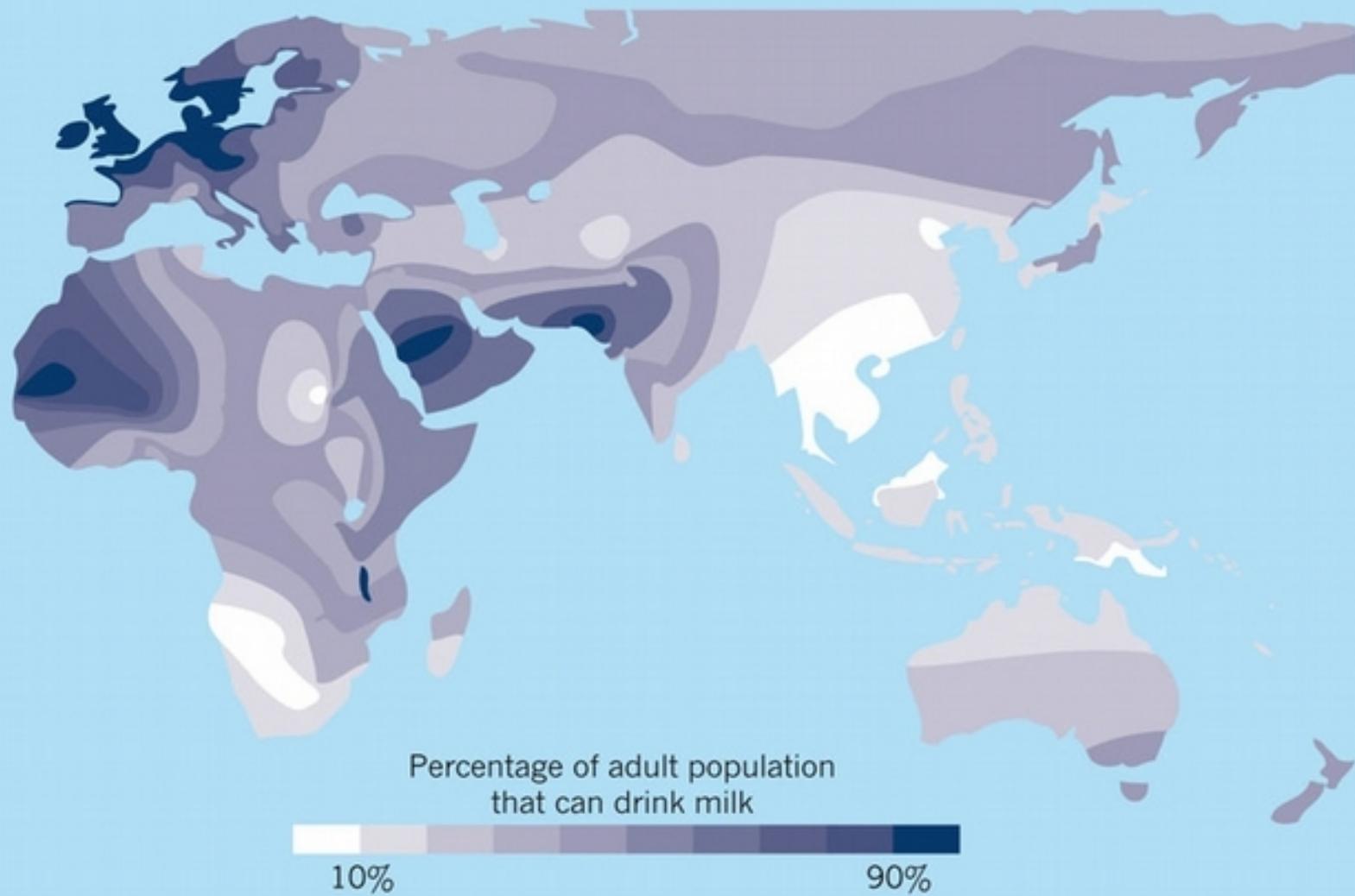
Lactase persistence alleles



Niche construction

LACTASE HOTSPOTS

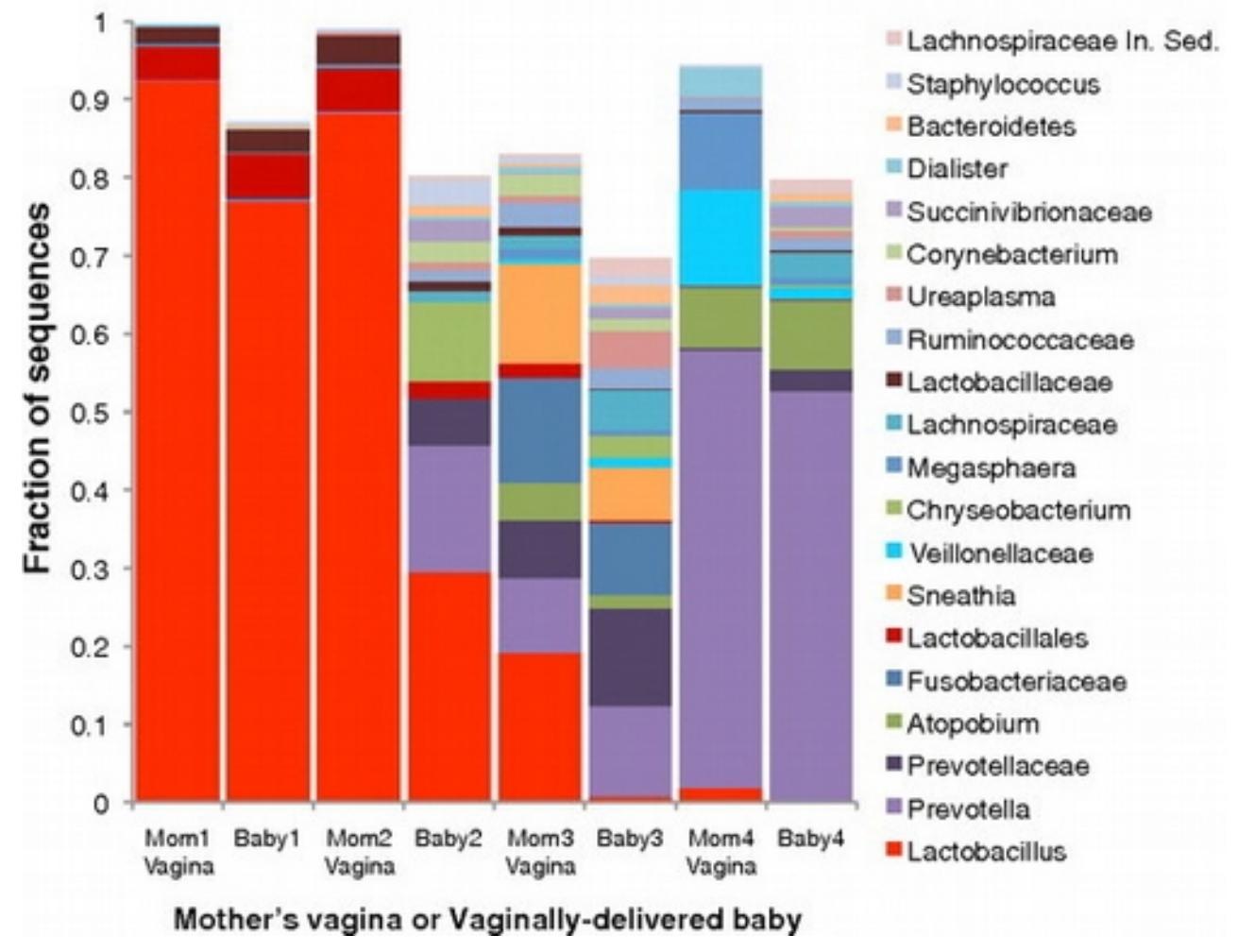
Only one-third of people produce the lactase enzyme during adulthood, which enables them to drink milk.



Extra-genetic inheritance

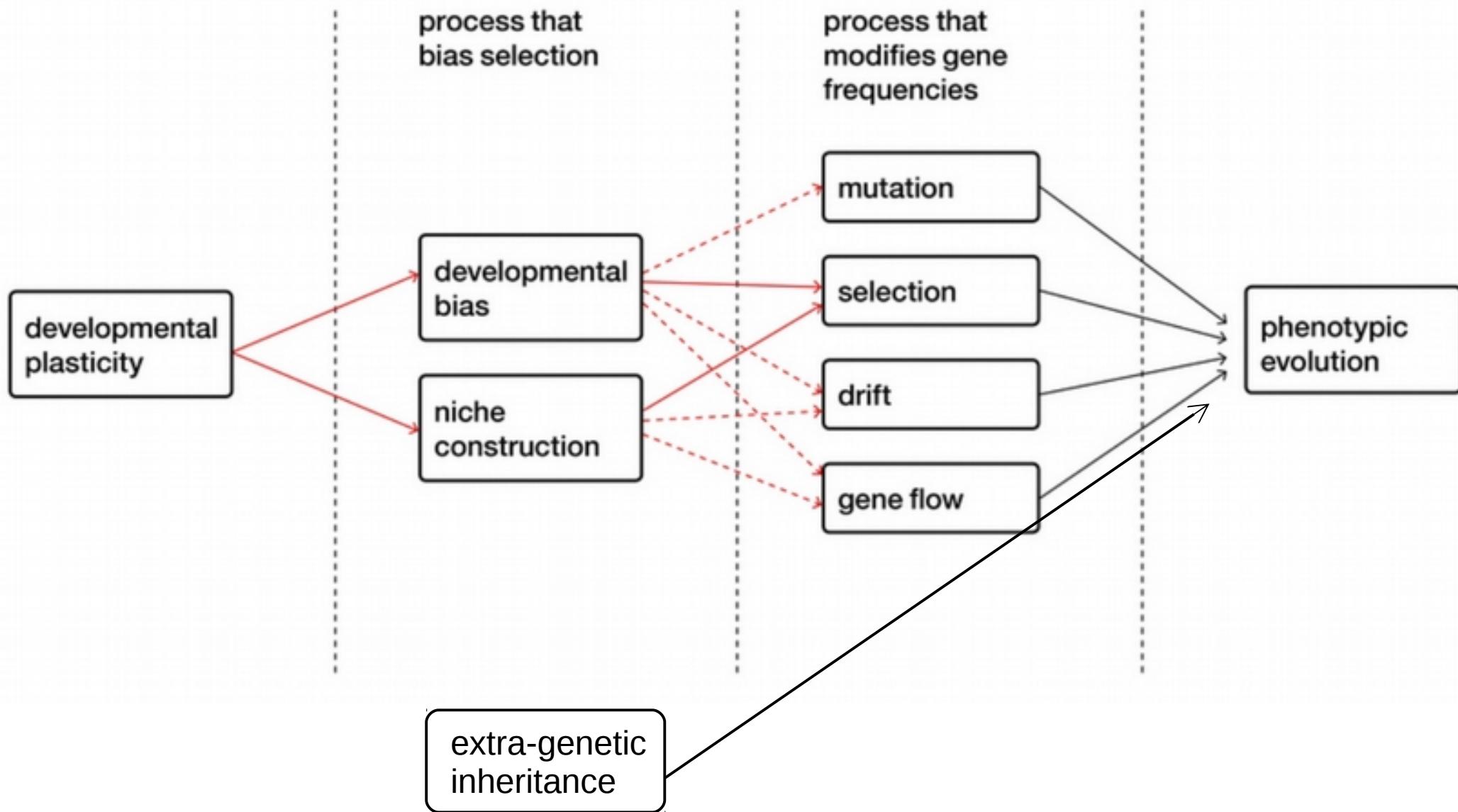
Antibodies from mother to baby's blood
via breast milk and placenta

Microbiome from mother to baby



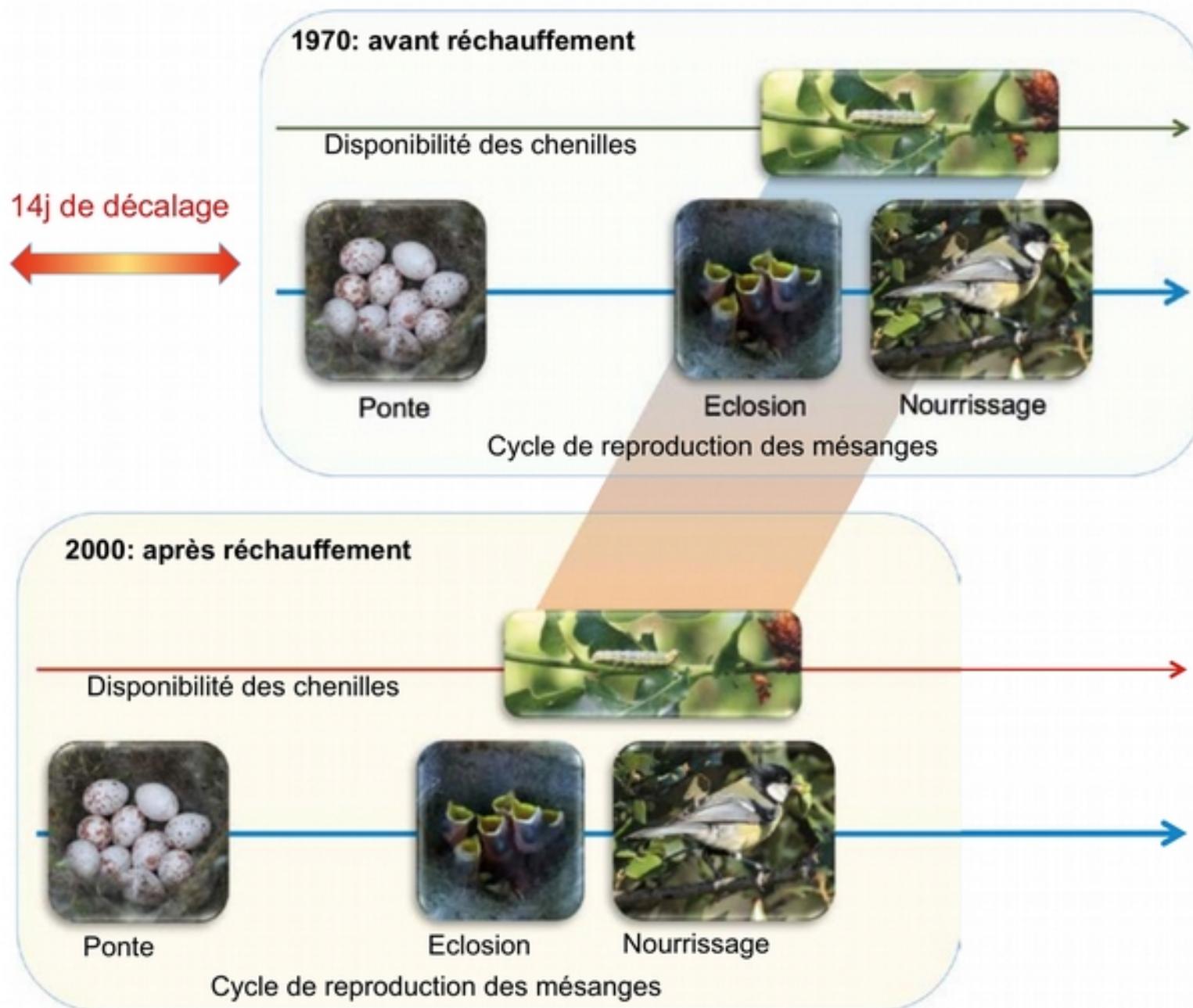
Additions

Classical view



classical MS core assumptions	EES core assumptions
(i) <i>The pre-eminence of natural selection.</i> The major directing or creative influence in evolution is natural selection, which alone explains why the properties of organisms match the properties of their environments (adaptation)	(i) <i>Reciprocal causation</i> (organisms shape, and are shaped by, selective and developmental environments). Developmental processes, operating through developmental bias and niche construction, share with natural selection some responsibility for the direction and rate of evolution and contribute to organism–environment complementarity
(ii) <i>Genetic inheritance.</i> Genes constitute the only general inheritance system. Acquired characters are not inherited	(ii) <i>Inclusive inheritance.</i> Inheritance extends beyond genes to encompass (transgenerational) epigenetic inheritance, physiological inheritance, ecological inheritance, social (behavioural) transmission and cultural inheritance. Acquired characters can play evolutionary roles by biasing phenotypic variants subject to selection, modifying environments and contributing to heritability
(iii) <i>Random genetic variation.</i> There is no relationship between the direction in which mutations occur—and hence the supply of phenotypic variants—and the direction that would lead to enhanced fitness	(iii) <i>Non-random phenotypic variation.</i> Developmental bias, resulting from non-random mutation or phenotypic accommodation, means that some phenotypic variants are more likely than others. Developmental systems facilitate well-integrated, functional phenotypic responses to mutation or environmental induction
(iv) <i>Gradualism.</i> Evolution via mutations of large effects is unlikely because such mutations have disruptive pleiotropic effects. Phenotypic transitions typically occur through multiple small steps, leading to gradual evolutionary change	(iv) <i>Variable rates of change.</i> Variants of large effect are possible, allowing for rapid evolutionary change. Saltation can occur either through mutations in major regulatory control genes expressed in tissue-, module- or compartment-specific manner, or when developmental processes respond to environmental challenges with change in coordinated suites of traits, or through nonlinear threshold effects
(v) <i>Gene-centred perspective.</i> Evolution requires, and is often defined as, <i>change in gene frequencies.</i> Populations evolve through changes in gene frequencies brought about through natural selection, drift, mutation and gene flow	(v) <i>Organism-centred perspective.</i> Developmental systems can facilitate adaptive variation and modify selective environments. Evolution redefined as a <i>transgenerational change in the distribution of heritable traits of a population.</i> There is a broadened notion of evolutionary process and inheritance
(vi) <i>Macro-evolution.</i> Macro-evolutionary patterns are explained by micro-evolutionary processes of selection, drift, mutation and gene flow	(vi) <i>Macro-evolution.</i> Additional evolutionary processes, including developmental bias and ecological inheritance, help explain macro-evolutionary patterns and contribute to evolvability

Evolution rapide des mésanges charbonnières



Our research approach

The genetic program is dead

Evolution is not as random as previously thought

The tree of life is not a tree

Our research approach

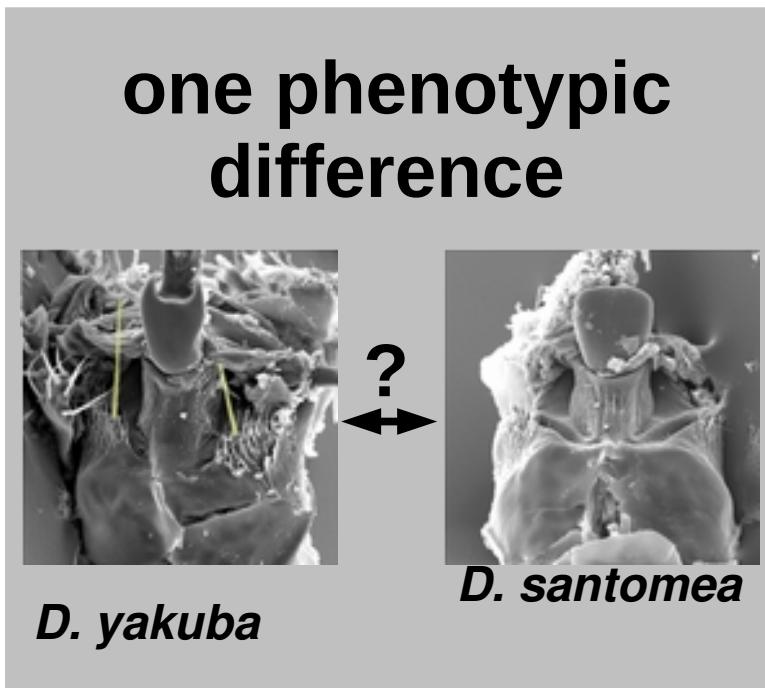
**Finding the mutations
responsible for evolution**



The mutations underlying evolution and diversity of living forms



Our strategy



Causing mutations?

Effect on development?

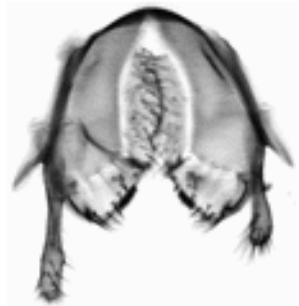
Effect on reproductive success?

Population dynamics?



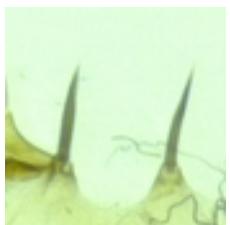
How D. pachea became dependent on its host cactus

Lang et al. 2012 Science



How and why asymmetric genitals evolved

Lang and Orgogozo 2012 Cont Zool
Rhebergen et al. 2016 BMC Evol Biol
unpublished



How a new bristle pattern evolved despite pleiotropy

unpublished



Gephebase: The Loci of Evolution

unpublished



Evolution is partly predictable

Orgogozo 2015 Interface Focus

Collaborators

Gephebase
Arnaud Martin
Baptiste Morizot

Fly anatomy
Jean David

Morphometrics
François Graner

Genotyping
David L. Stern

Fly crosses
Daniel Matute

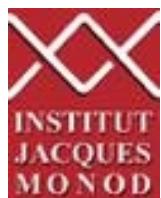
Sex bristles
Alexis Matamoro-Vidal
Isabelle Nuez
Olga Nagy
Alexandre Peluffo

Ventral branches
Alexandre Peluffo

Posterior lobe
Alexis Lalouette

Left-right asymmetry
Michael Lang
Andrea Acurio

Gephebase
Stéphane Prigent
Laurent Arnoult





The Database of Genotype-Phenotype Relationships

Search Gephebase for genes, phenotypes, taxa, mutations, articles:



keyword [search completion]



↓ User Login/Password

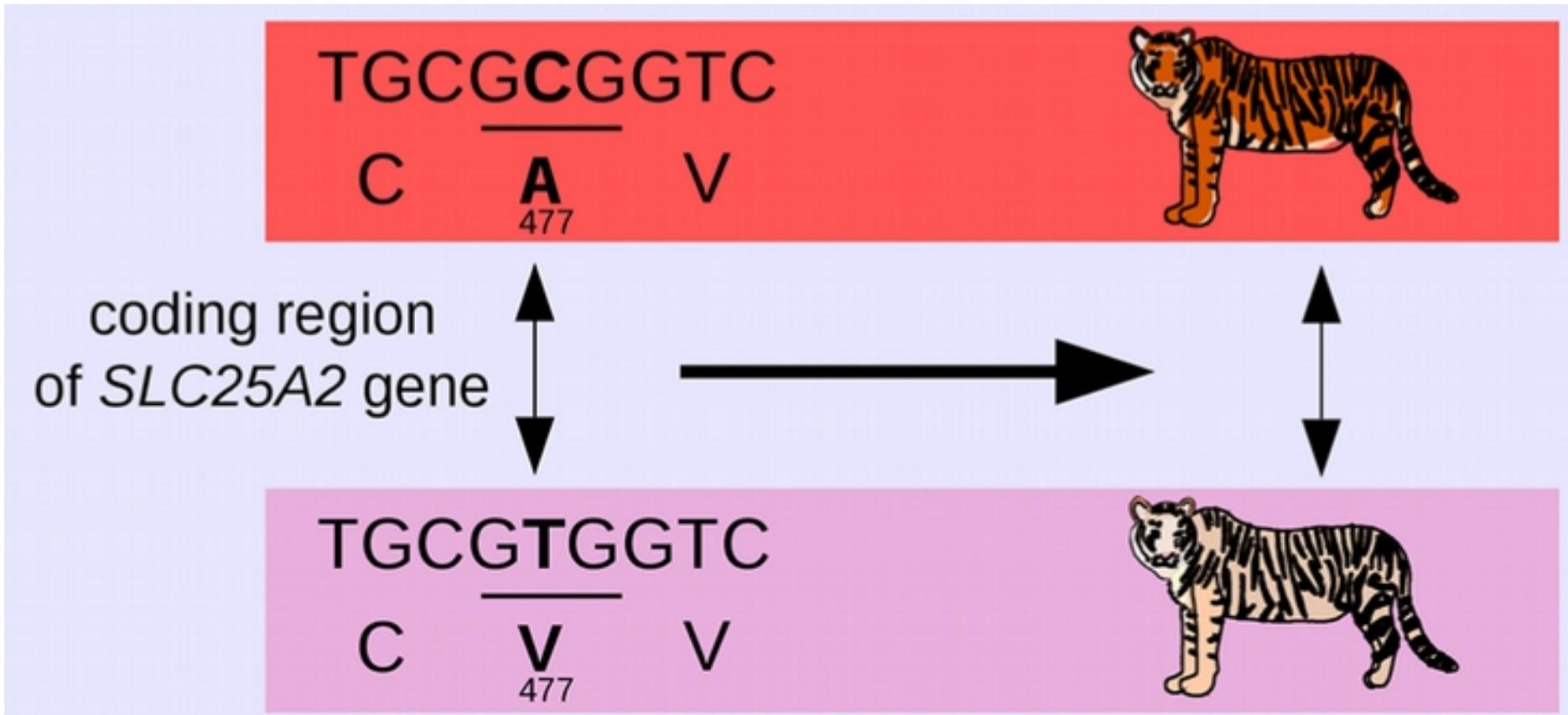
>1400 cases of evolutionary changes
associated to a gene locus
in animals and plants

Martin and Orgogozo 2013 Evolution
Ongoing project





The genotype-phenotype connection



Xu et al 2013 Current Biology
Orgogozo et al 2015 Frontiers Genetics

The genotype-phenotype connection

Report of the National Academy of Sciences on gene drive, 2016

Genotype = “the genetic makeup of an organism that determines a specific phenotype (trait), from one generation to the next, and potentially throughout the population”.

NO!

The genetic program is dead

Cunz, 2008, BMC Computational Biology:

“The genetic program encoding the morphological identity of a single dendrite remains a mystery”.

The phenotype is not caused by genes only



The phenotype is not caused by genes only

No carotenoid

Carotenoid restricted diet



Causes of these differences ?

Genetic



Environment



$$\text{Phenotype} = G + E + GxE$$

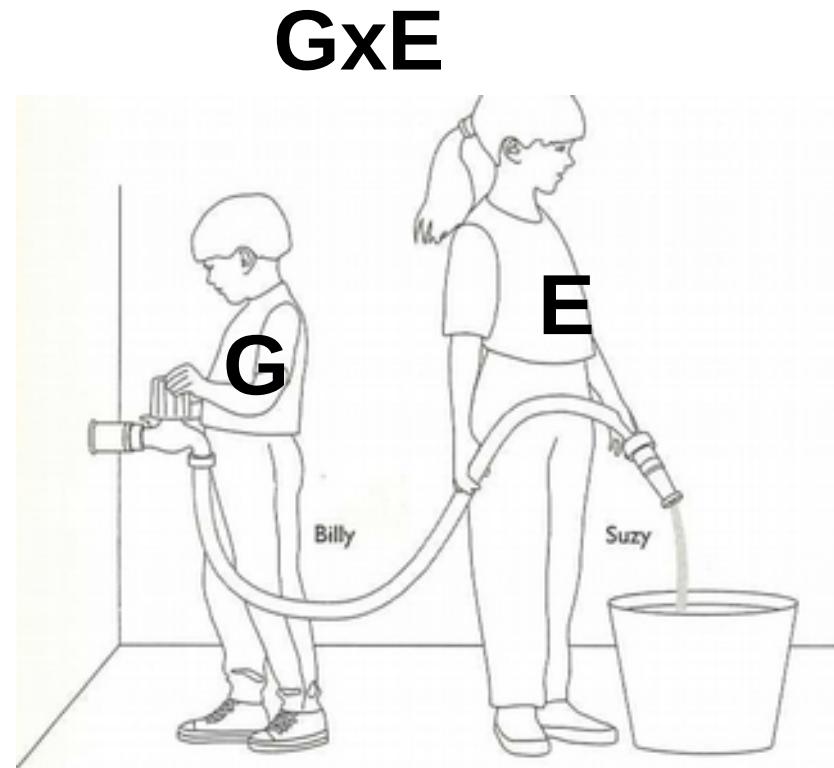
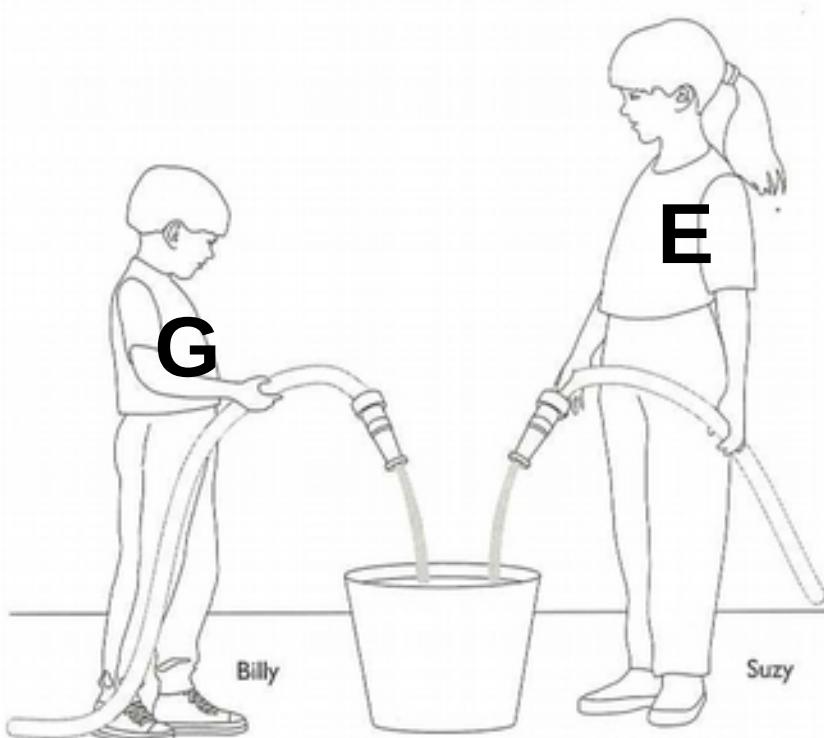
The Siamese cat

An example of GxE

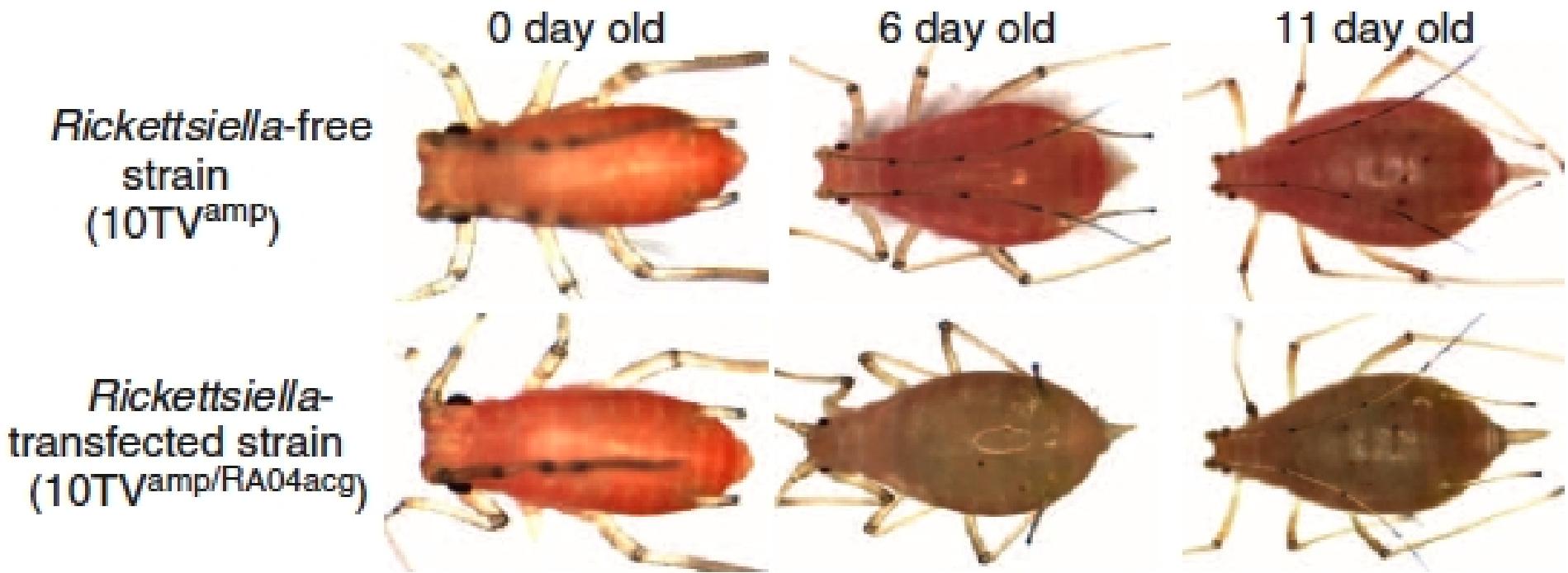


Mutation in *tyrosinase*
Heat-sensitive enzyme
No production of
melanin in warm body
parts

Contributions of the genotype (G) and the environment (E) to phenotypic variation

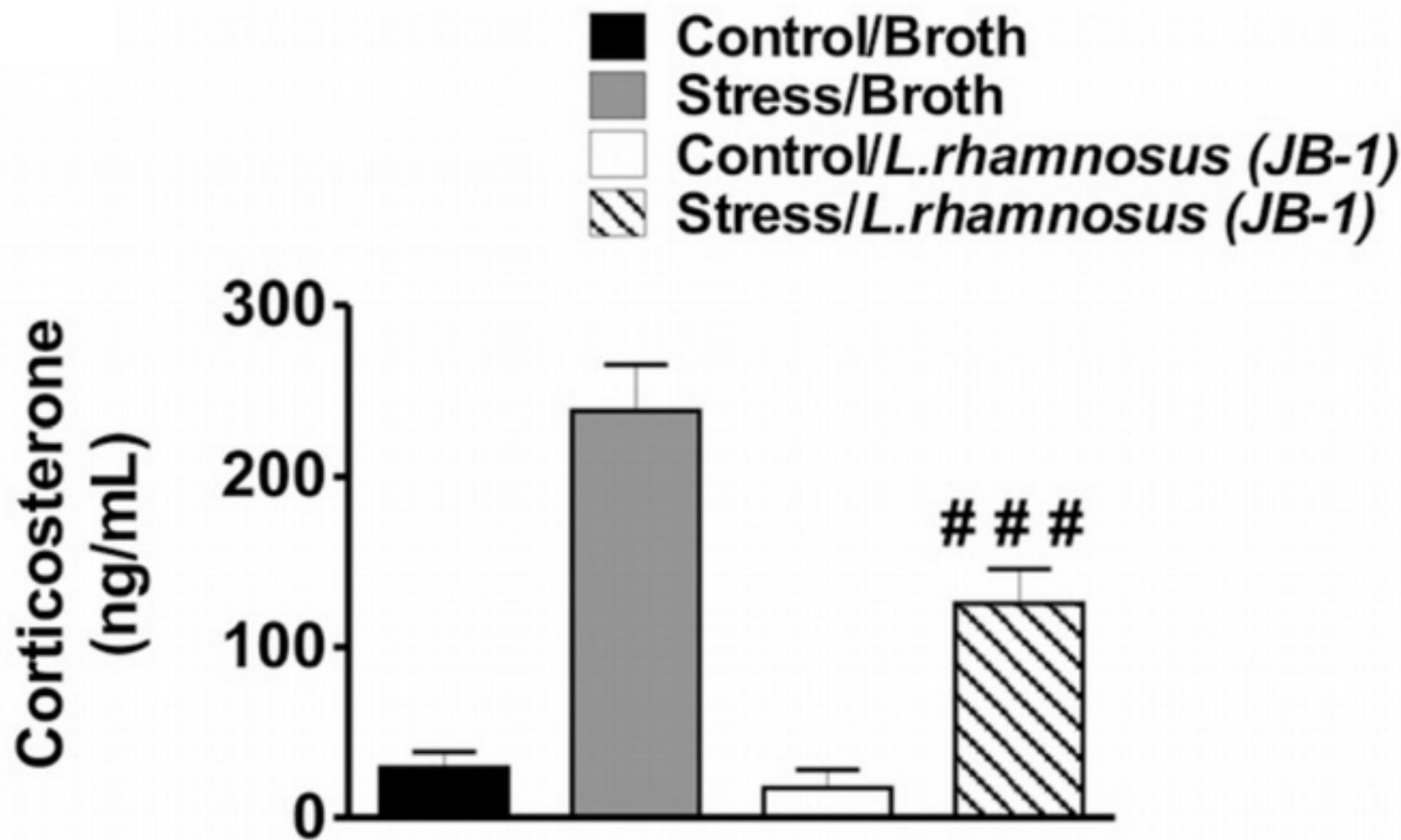


Pea aphid color variation

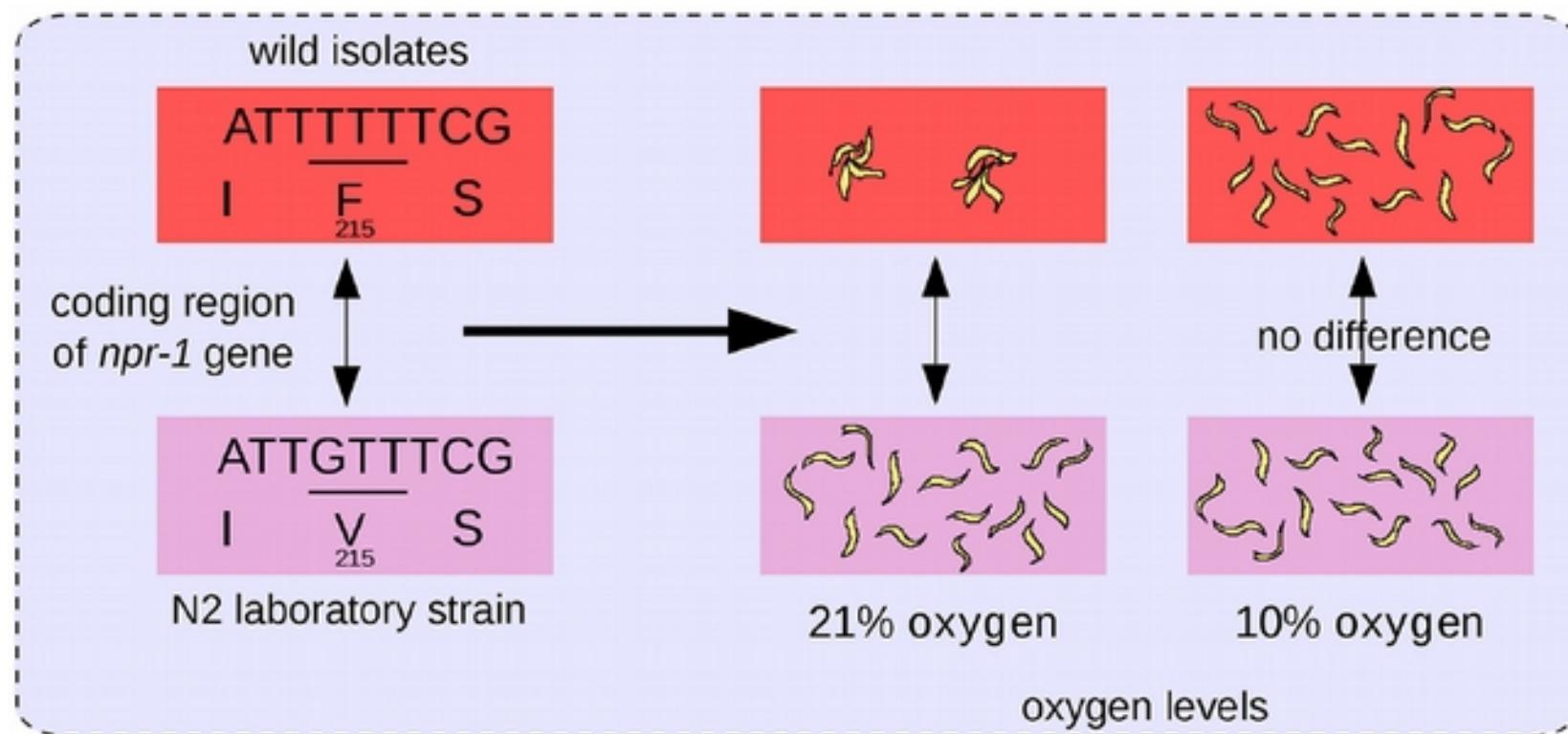


Without *Rickettsiella*, **red** aphid newborns become **red** adults.
With *Rickettsiella*, **red** aphid newborns become **green** adults.

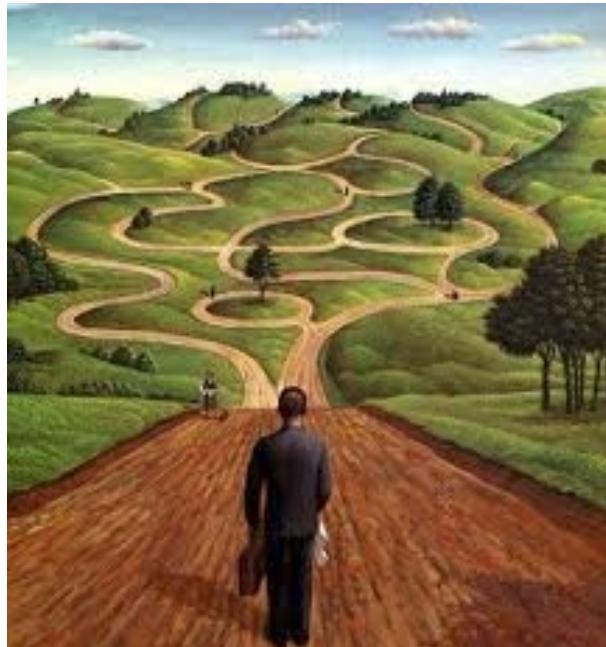
Mouse emotional behavior modified by presence of *Lactobacillus* bacteria



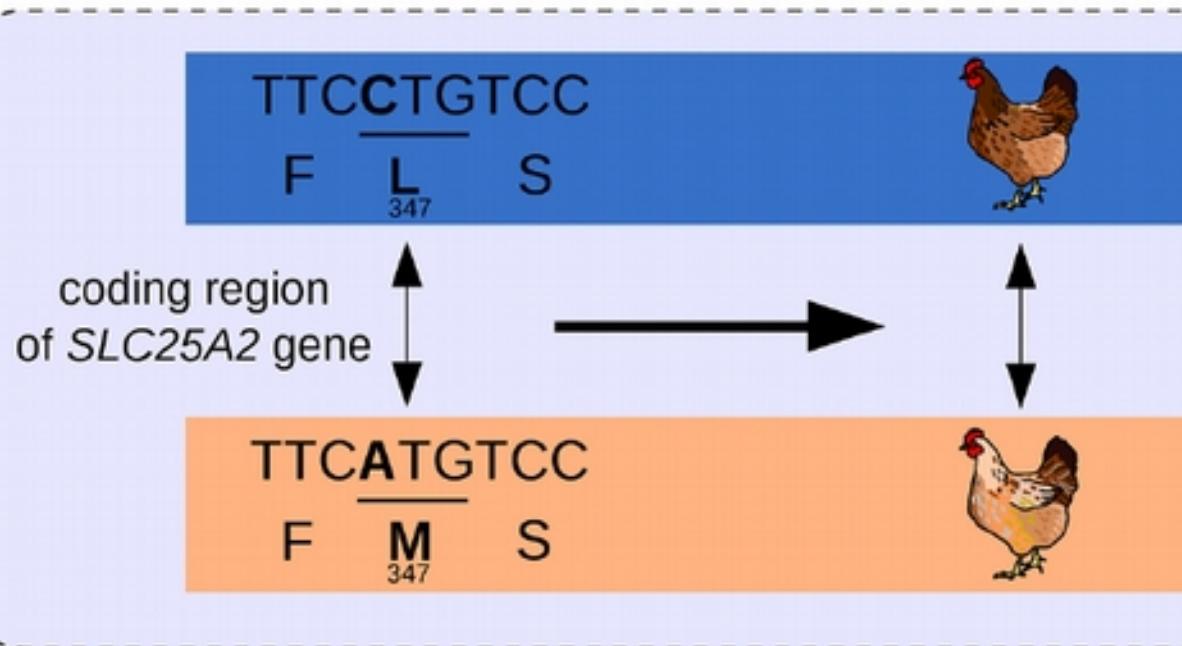
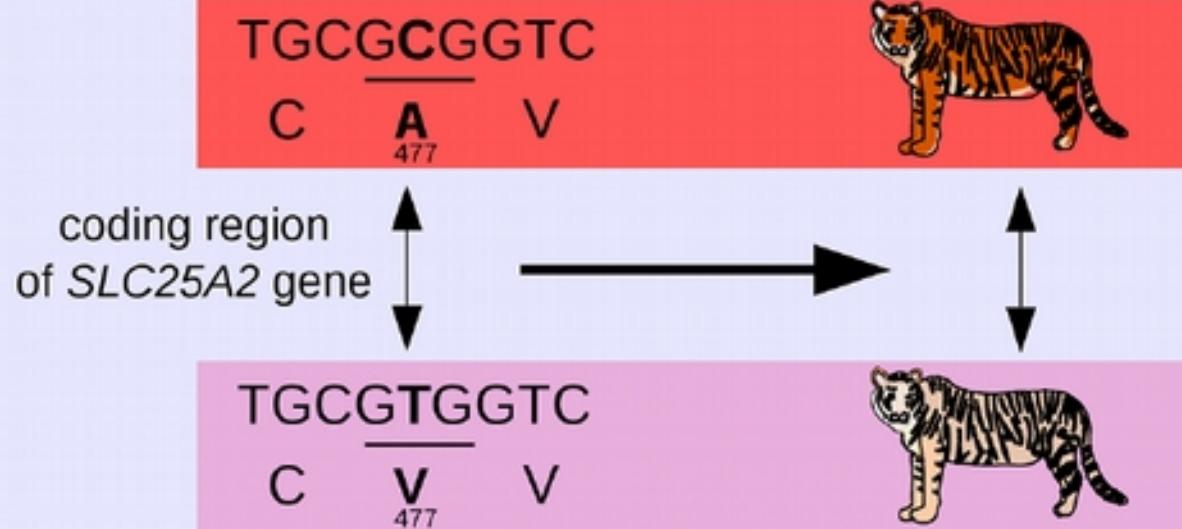
GxE: a difference within a difference



Evolution is not as random as previously thought



Repeated evolution



Repeats 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

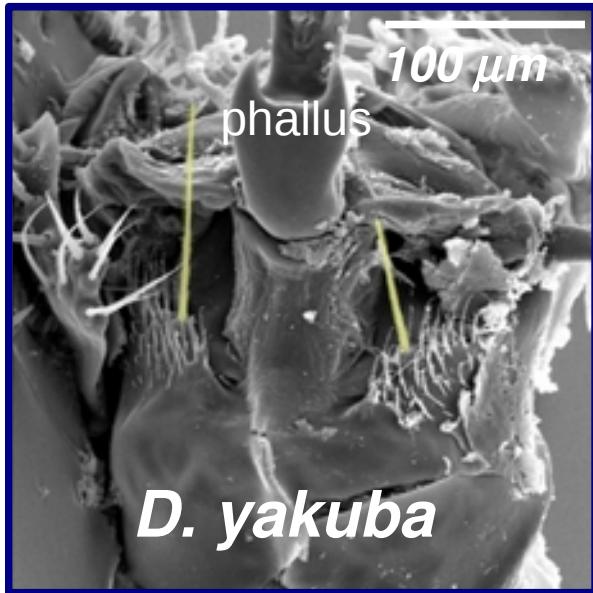
Ex : *sulfate transporter SUL1* in yeasts in low sulfate
(Gresham et al 2008 PloS Genetics)

.. the phenotypes evolving in certain environments

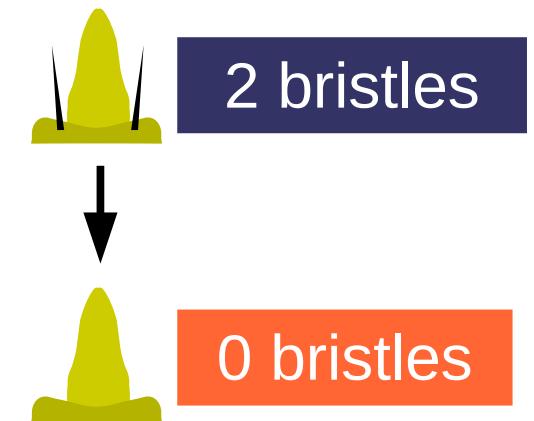
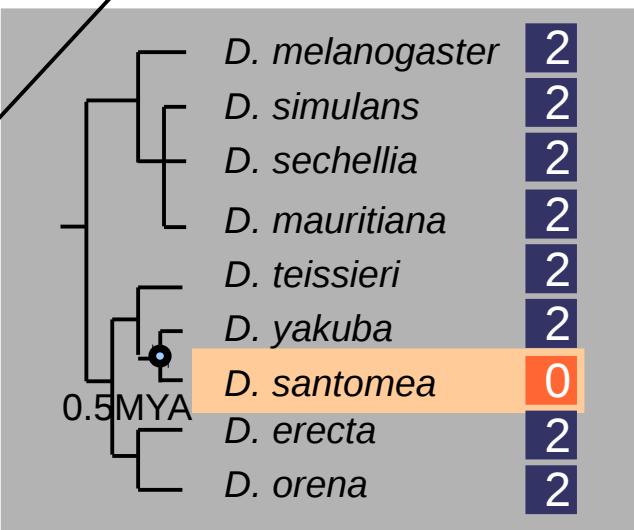
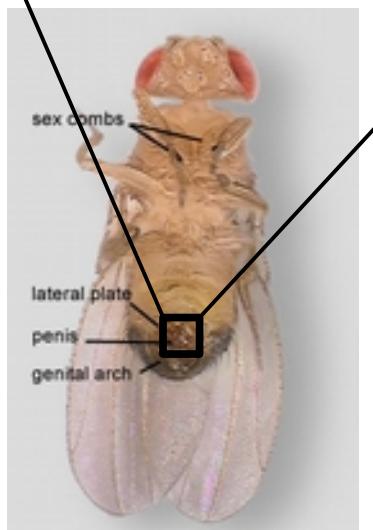
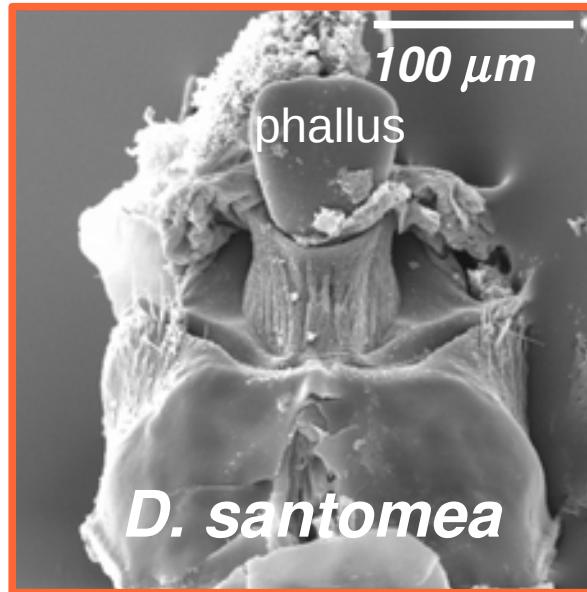


Ex : flying marsupial phalanger and placental flying squirrel

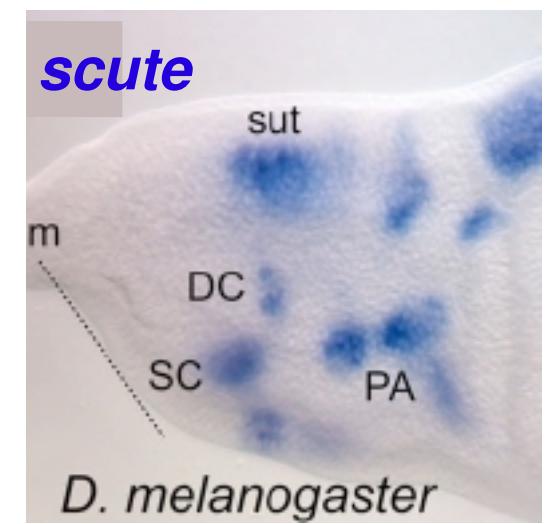
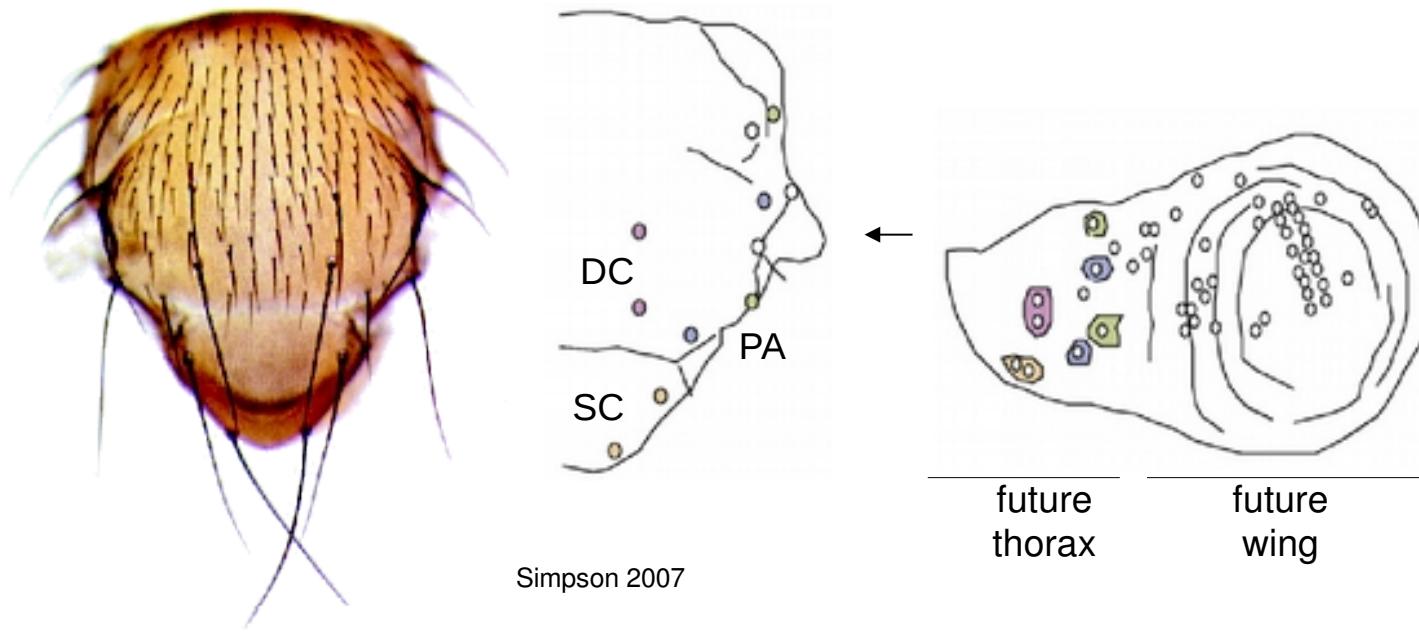
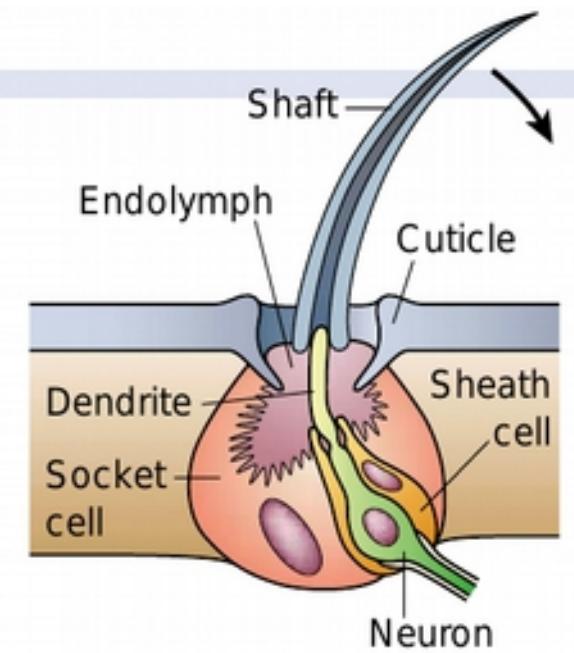
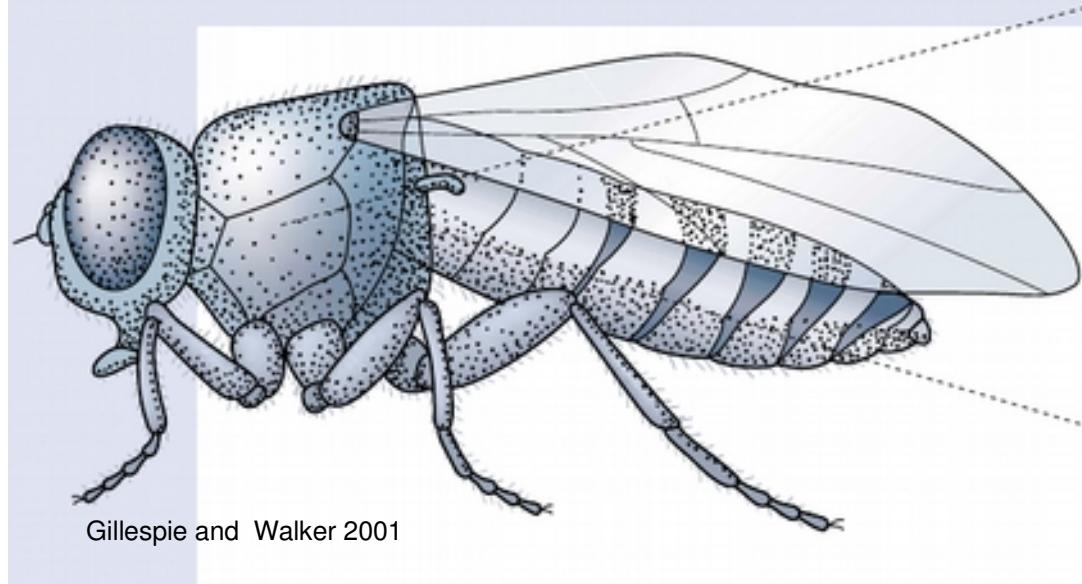
What are the mutations responsible for the loss of sex bristles in *D. santomea*?

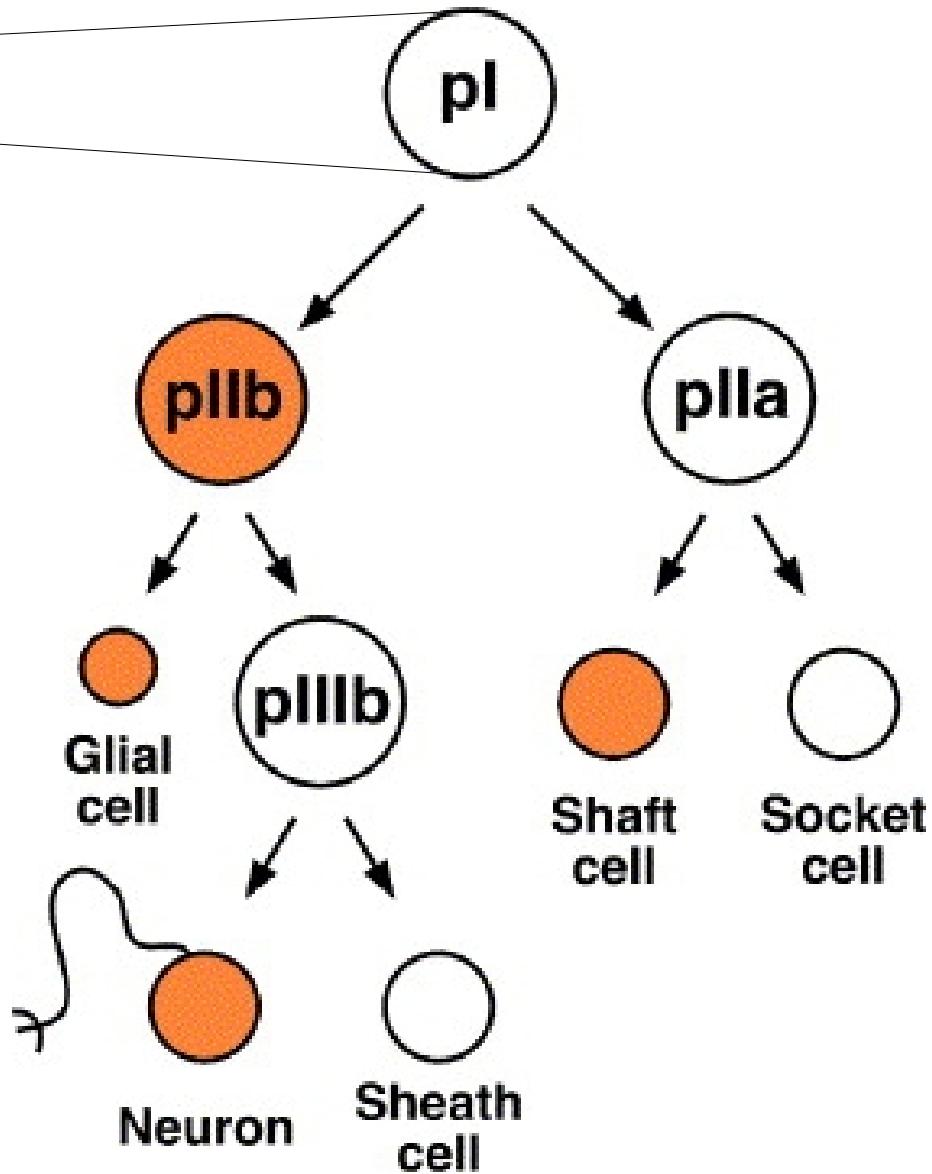
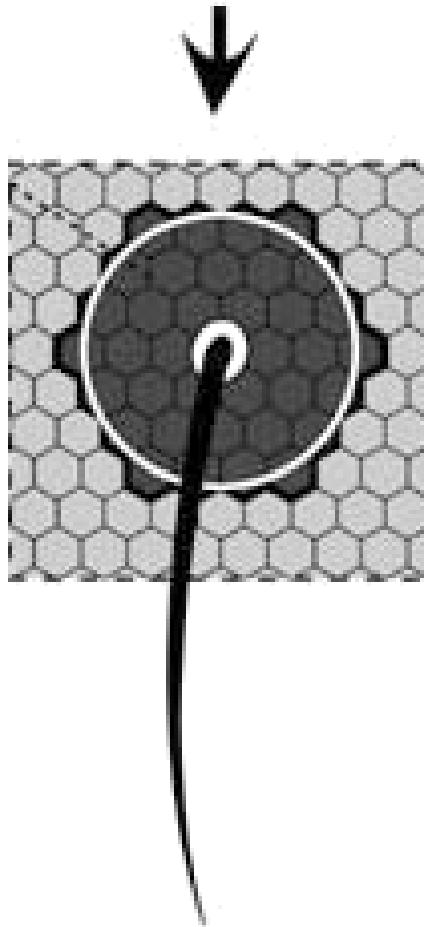
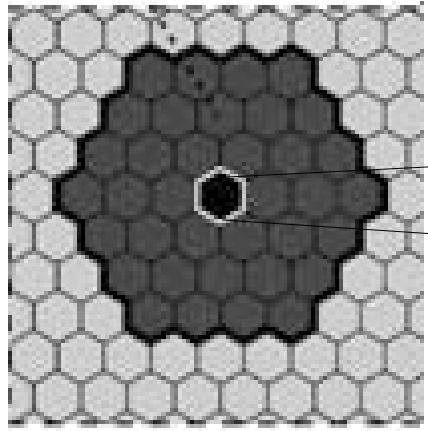


mutation(s)?

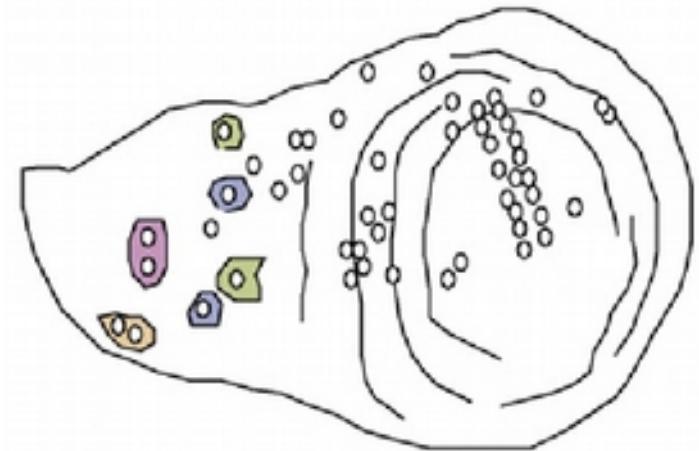
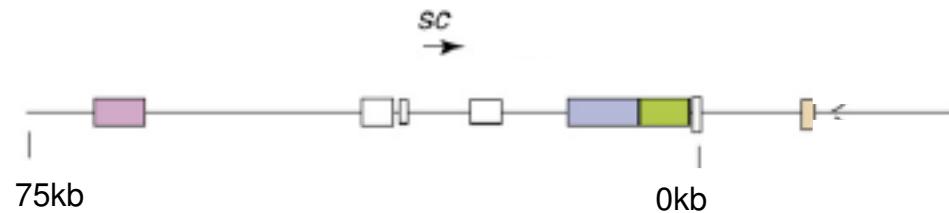


Bristle development

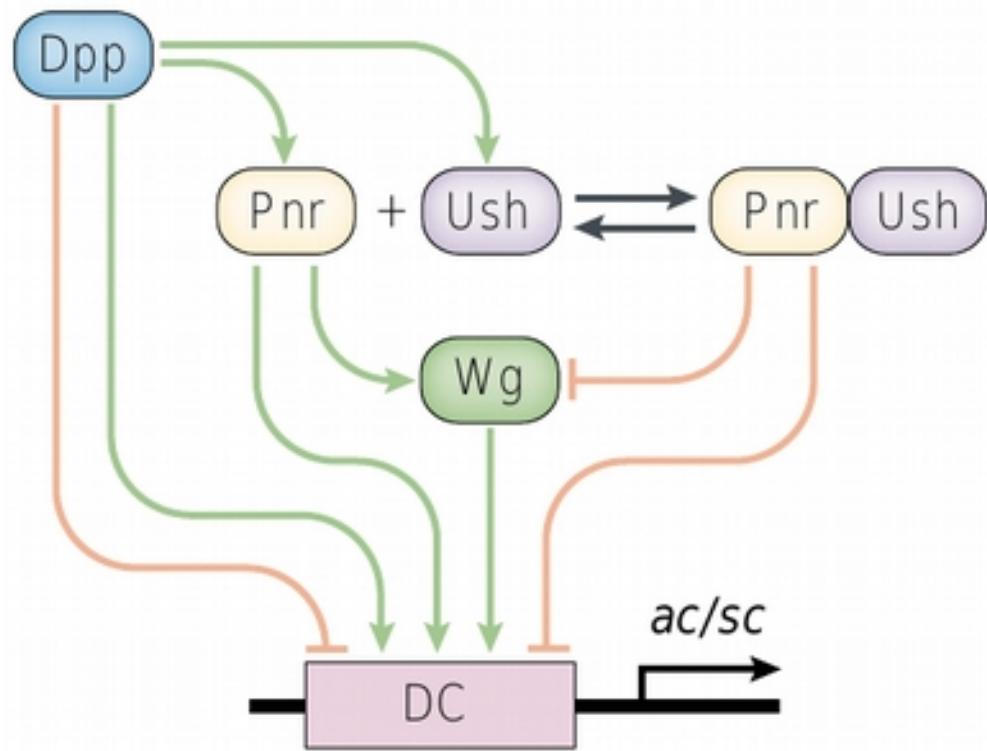




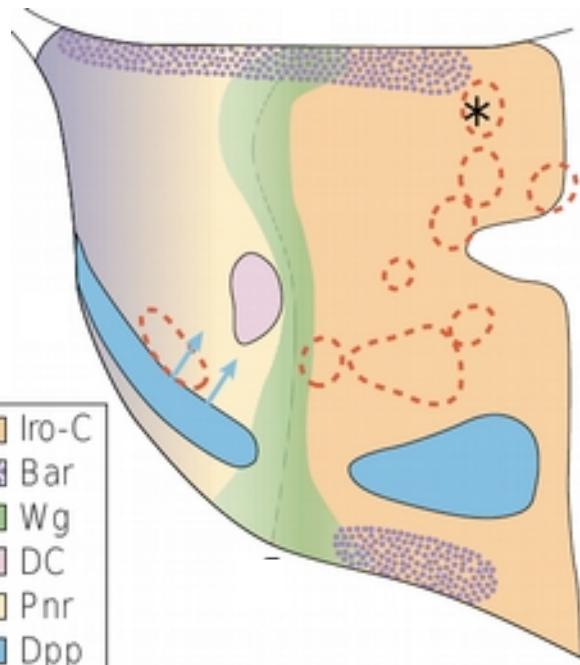
scute cis-regulatory elements are “master switches”



Simpson 2007



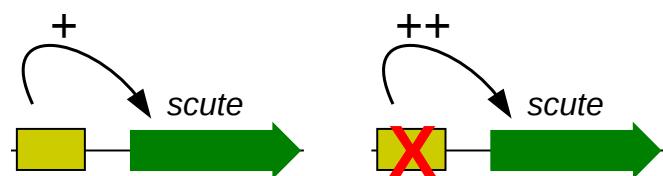
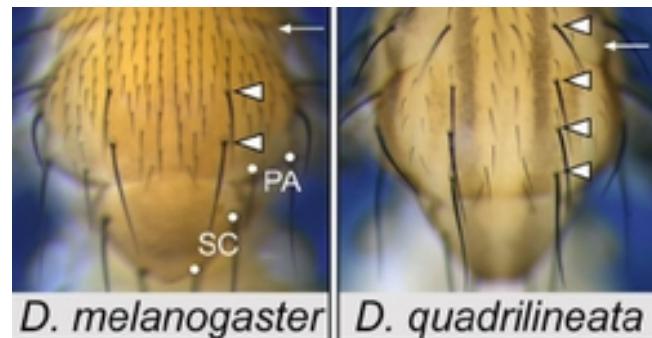
Iro-C
Bar
Wg
DC
Pnr
Dpp
Ush



Gómez-Skarmeta 2003

Evolution of fly bristle pattern

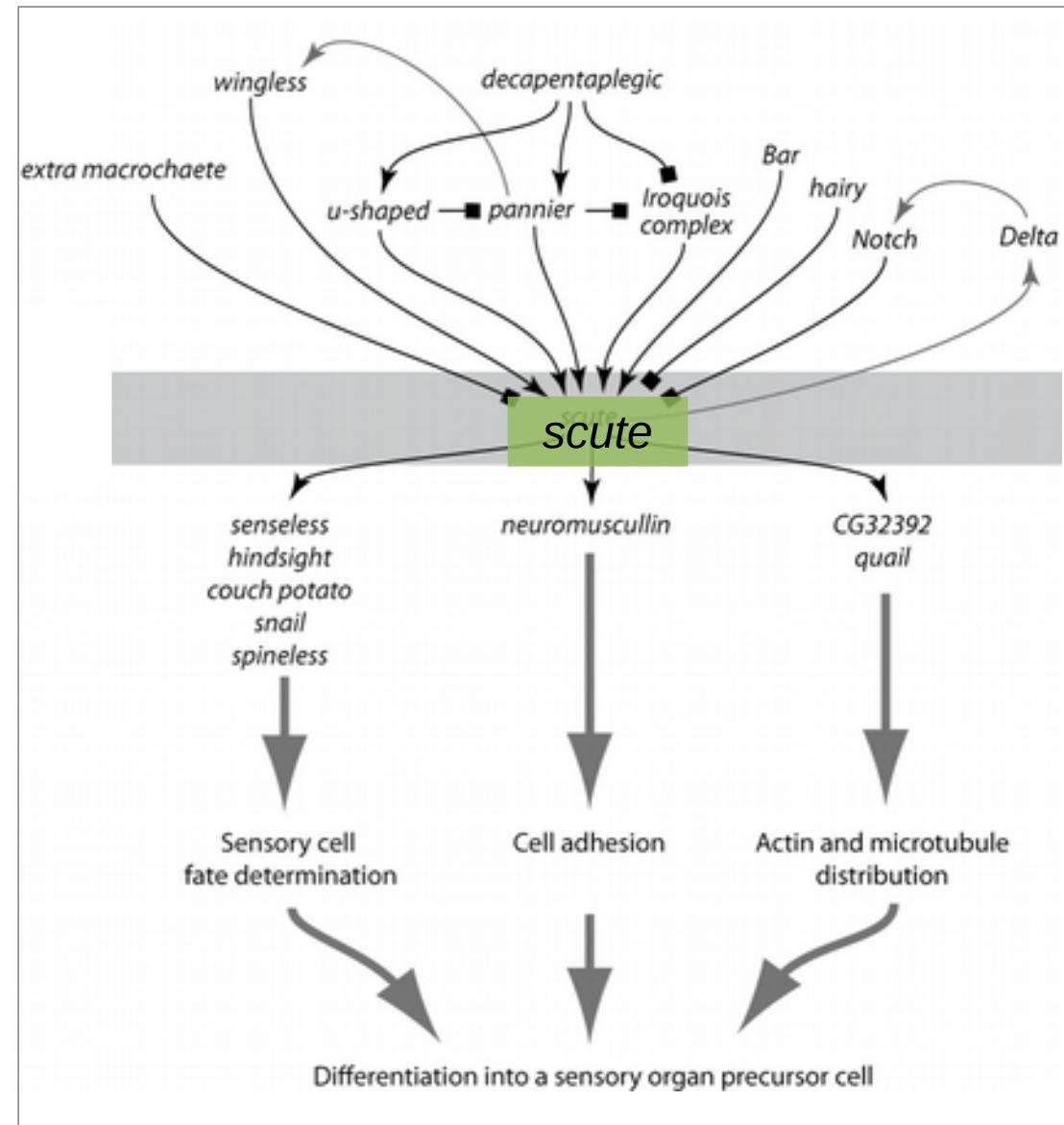
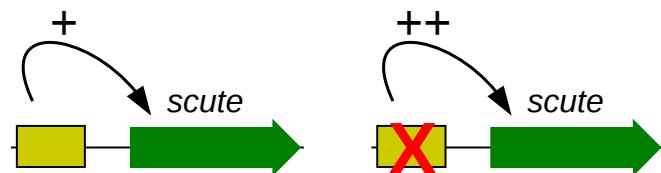
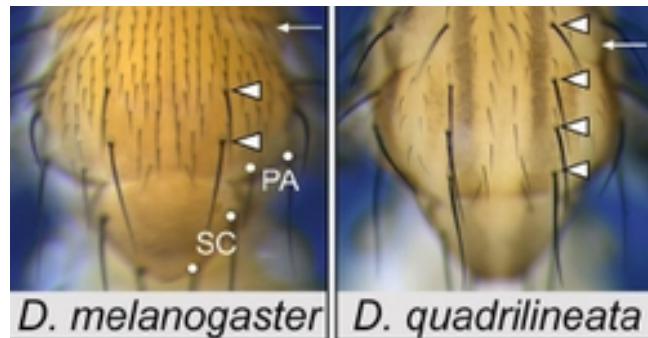
Cis-regulatory element (CRE) in *scute*



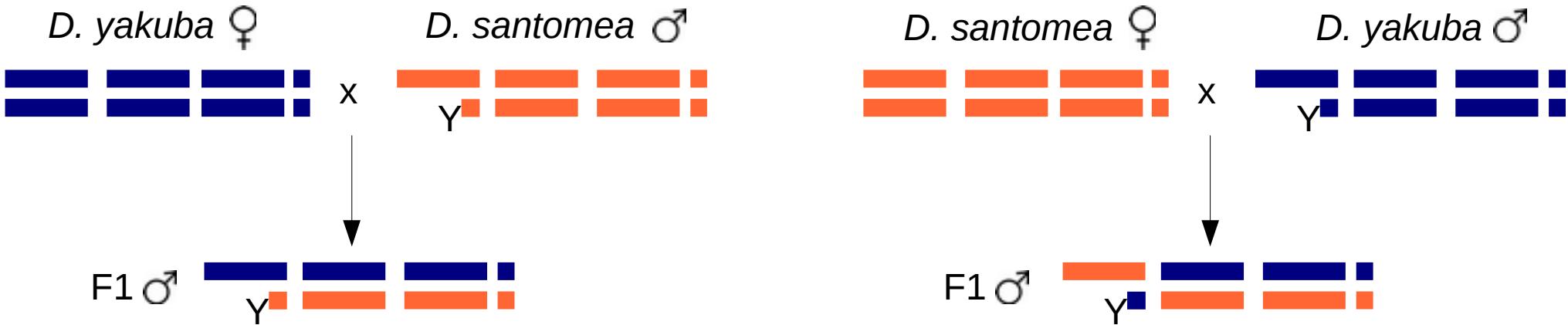
Genetic evolution is predictable

Evolution of fly bristle pattern

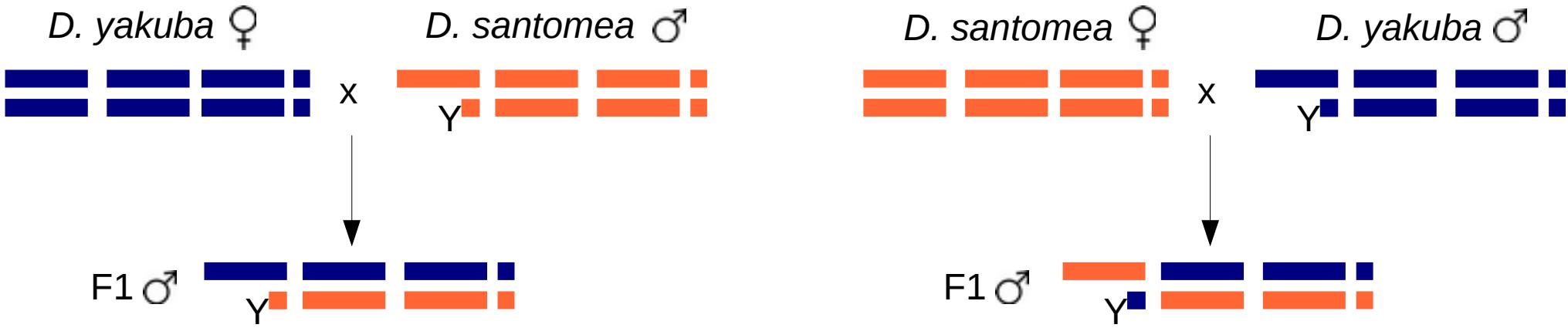
Cis-regulatory element (CRE) in *scute*



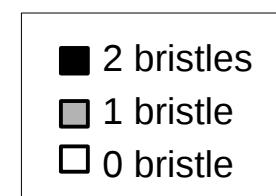
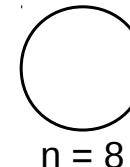
Is the causing mutation X-linked?



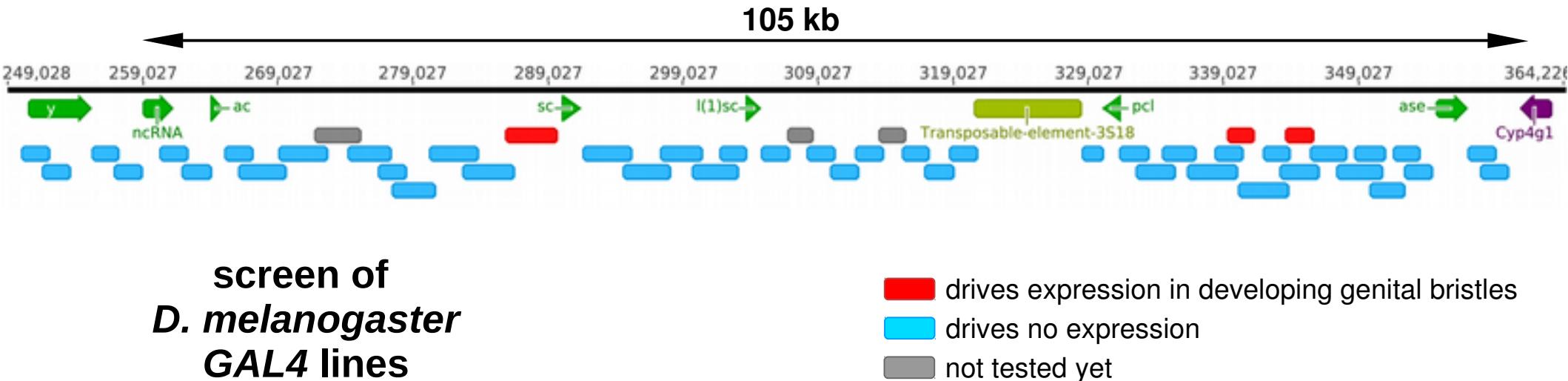
Is the causing mutation X-linked?



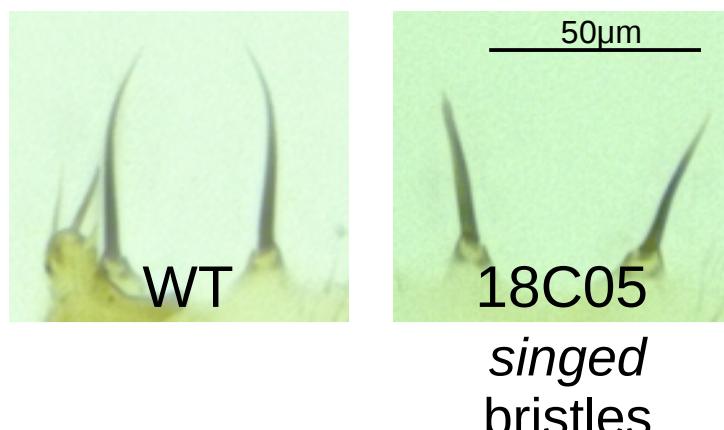
The
causing
mutation is
X-linked



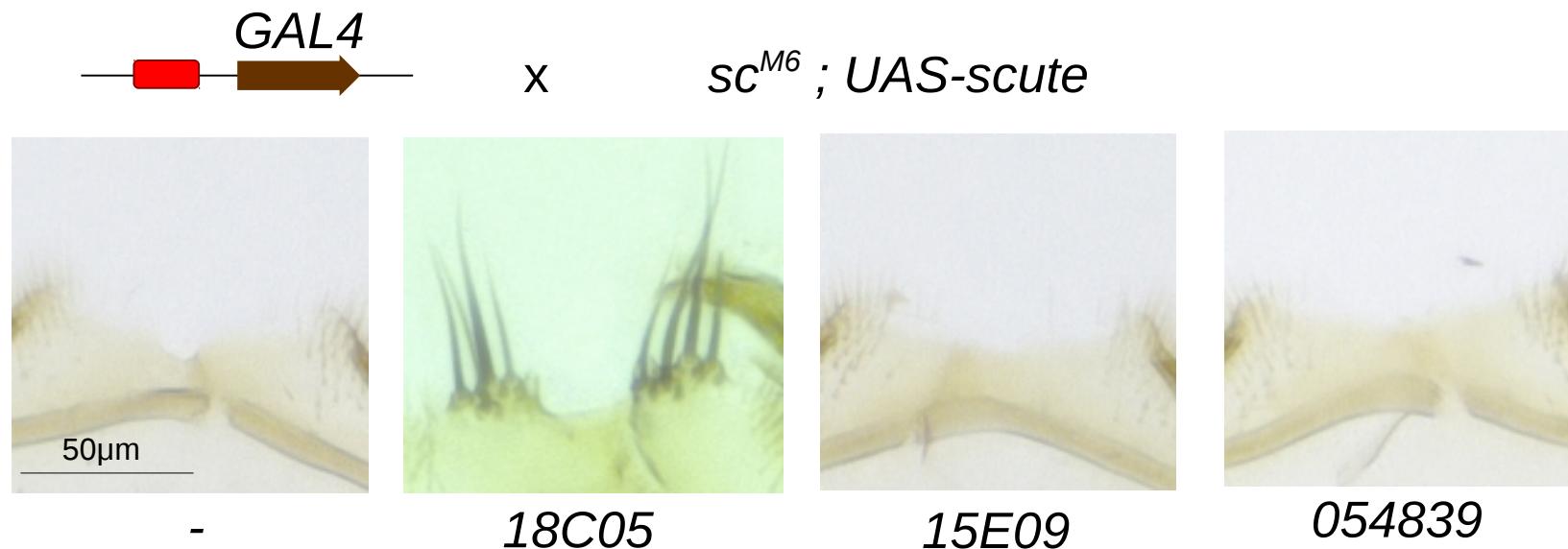
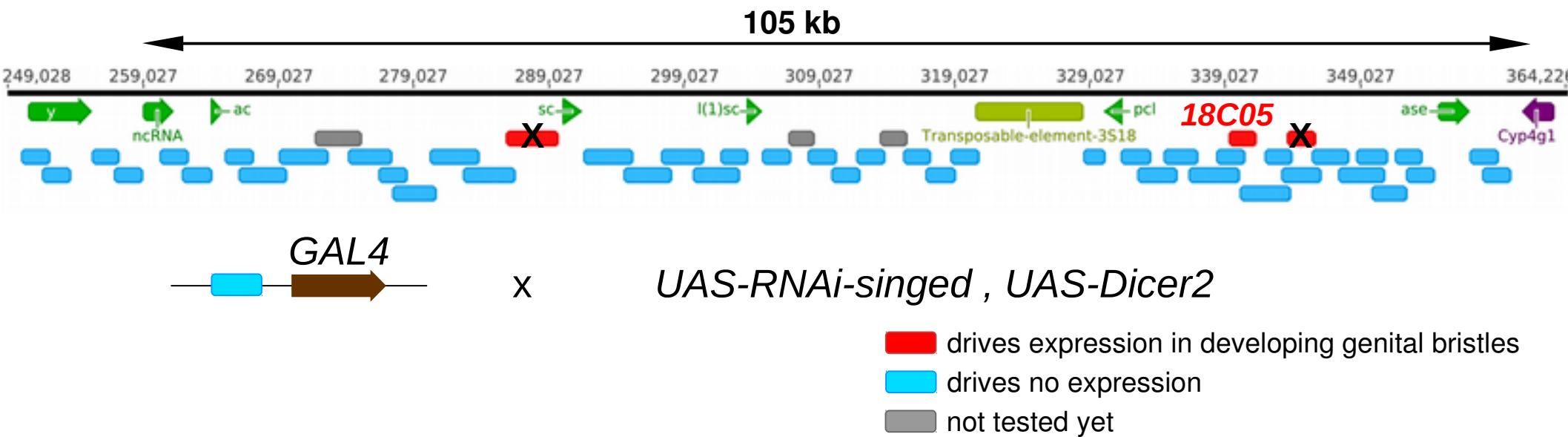
Screening a 100-kb region



GAL4 X *UAS-RNAi-singed , UAS-Dicer2*

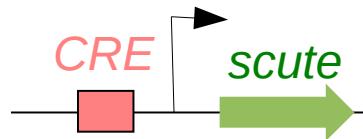


18C05 drives expression independently of scute



Mutation(s) in 18C05 cause loss of bristles

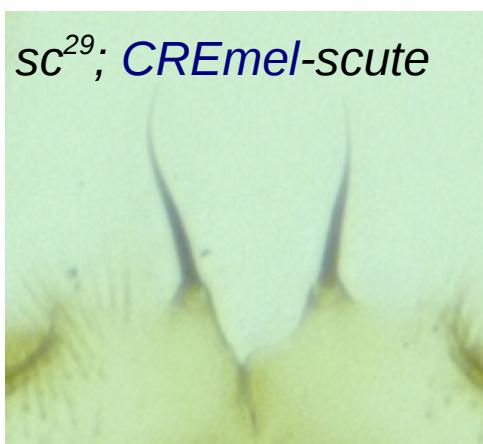
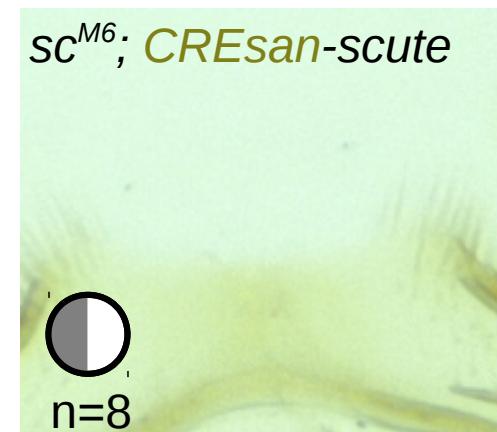
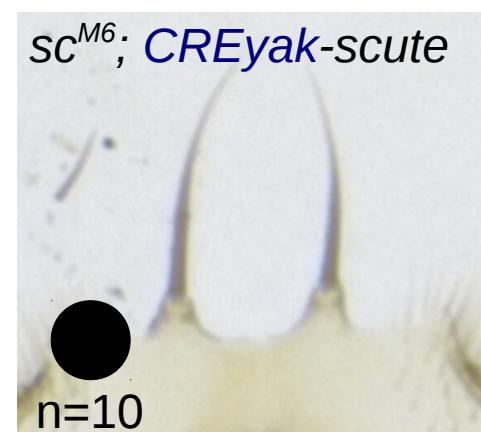
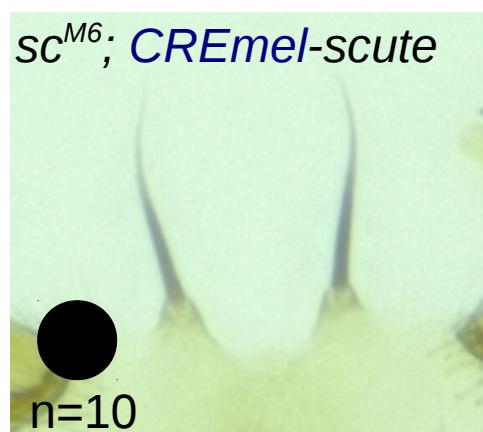
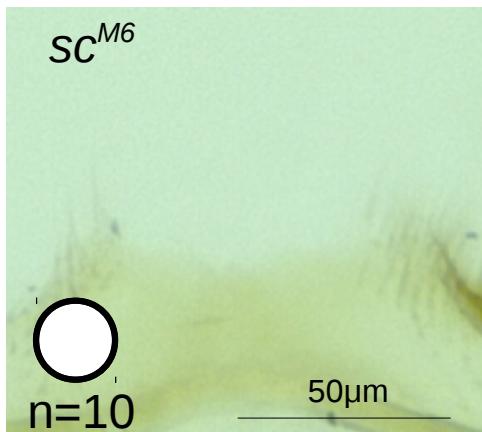
D. yakuba



D. santomea



D. melanogaster transgenics



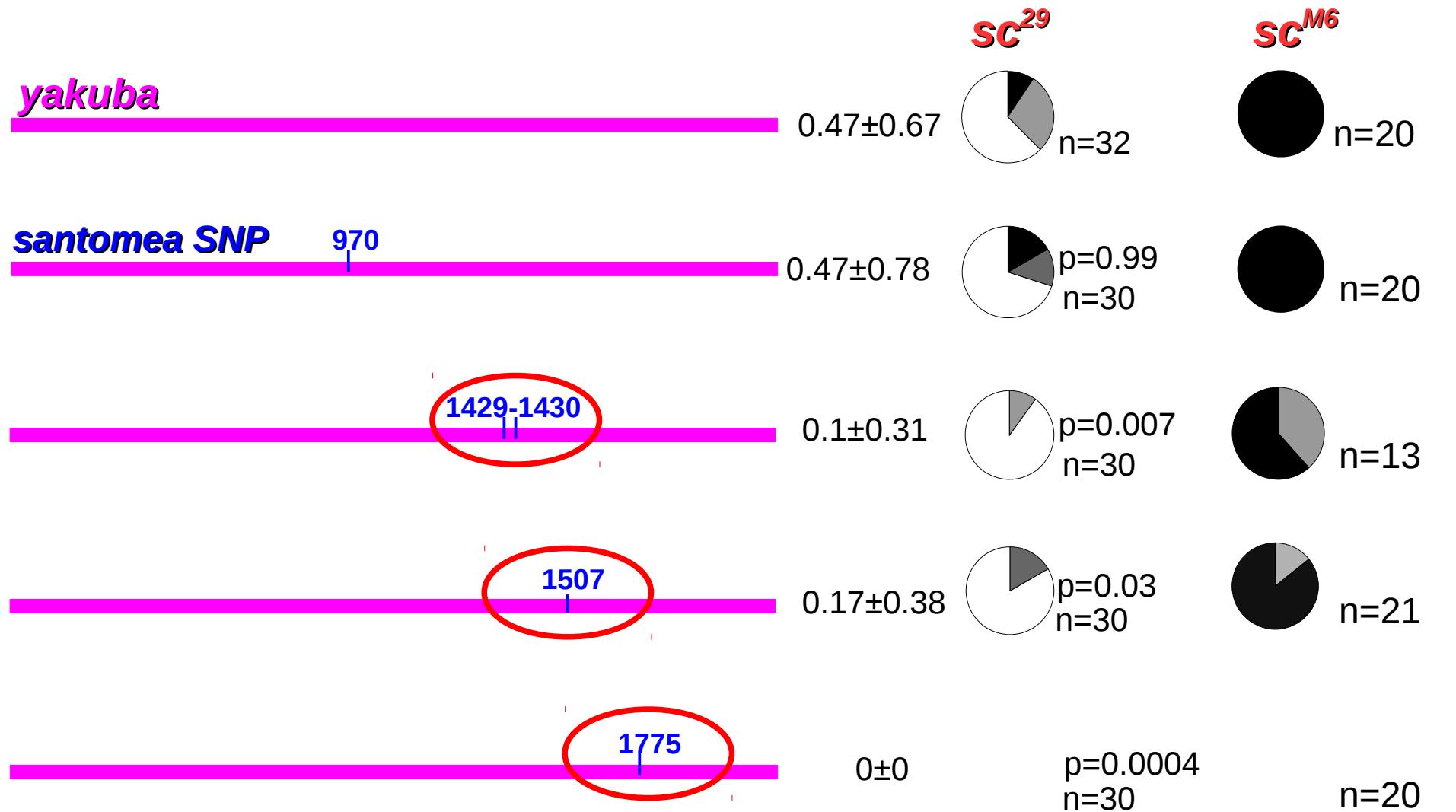
$SC^{29}; CRE_{yak}-scute$



27 SNPs and 3 indels between *D. santomea* and *D. yakuba*



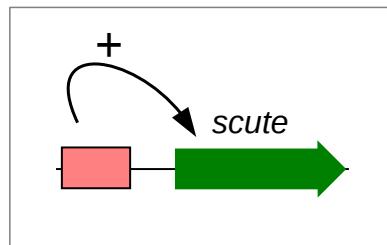
At least 3 mutations affect bristle number



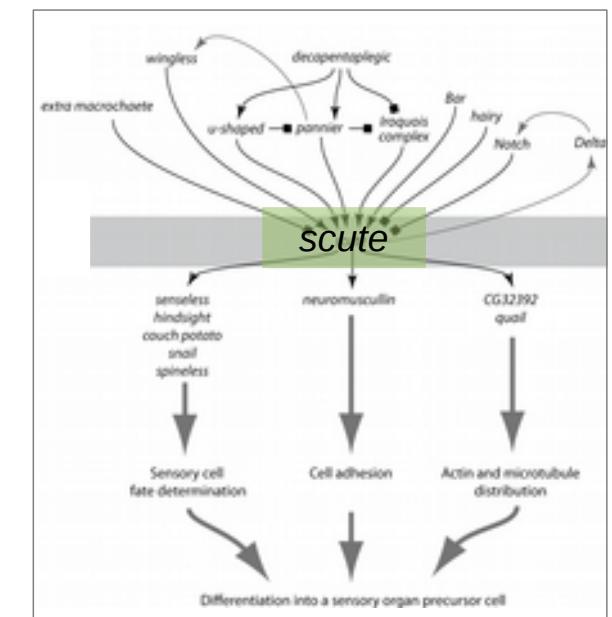
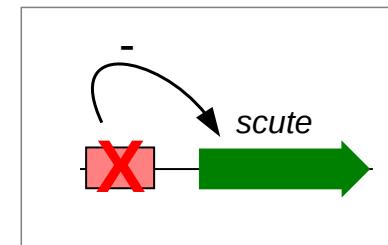
D. santomea lost genital bristles via mutations in *scute*



Ancestral state



Derived state



**Why is the set of genes
causing evolution limited?**

There are specialized genes in a genome

Steroid hormone biosynthesis

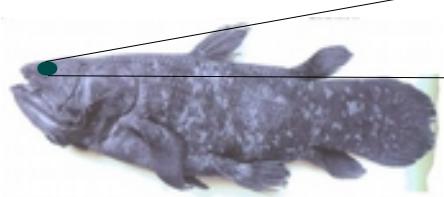


a specialized tissue
specialized enzymes

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**

Specialized loci in the genome

Proteins that interact with external molecules

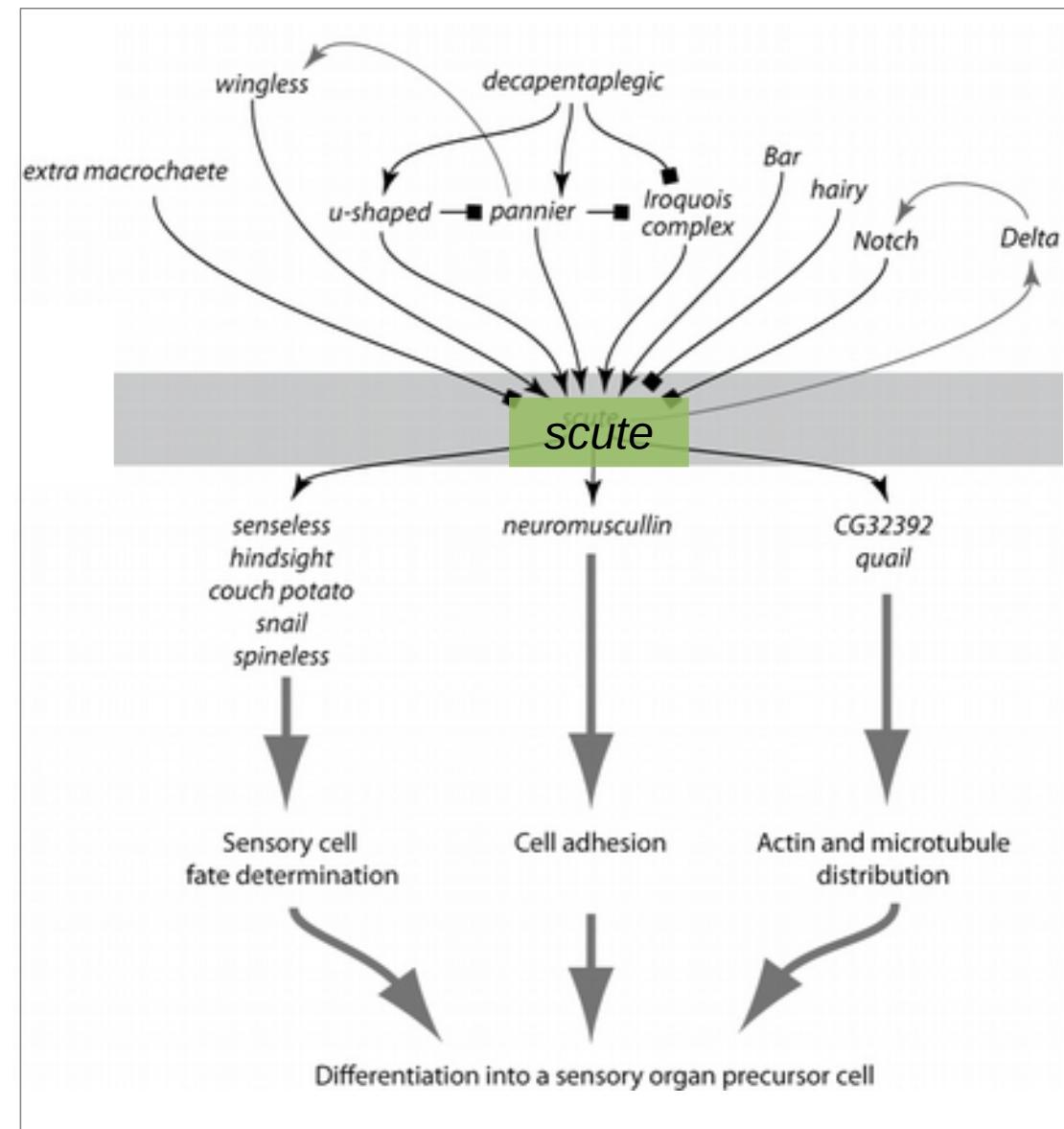
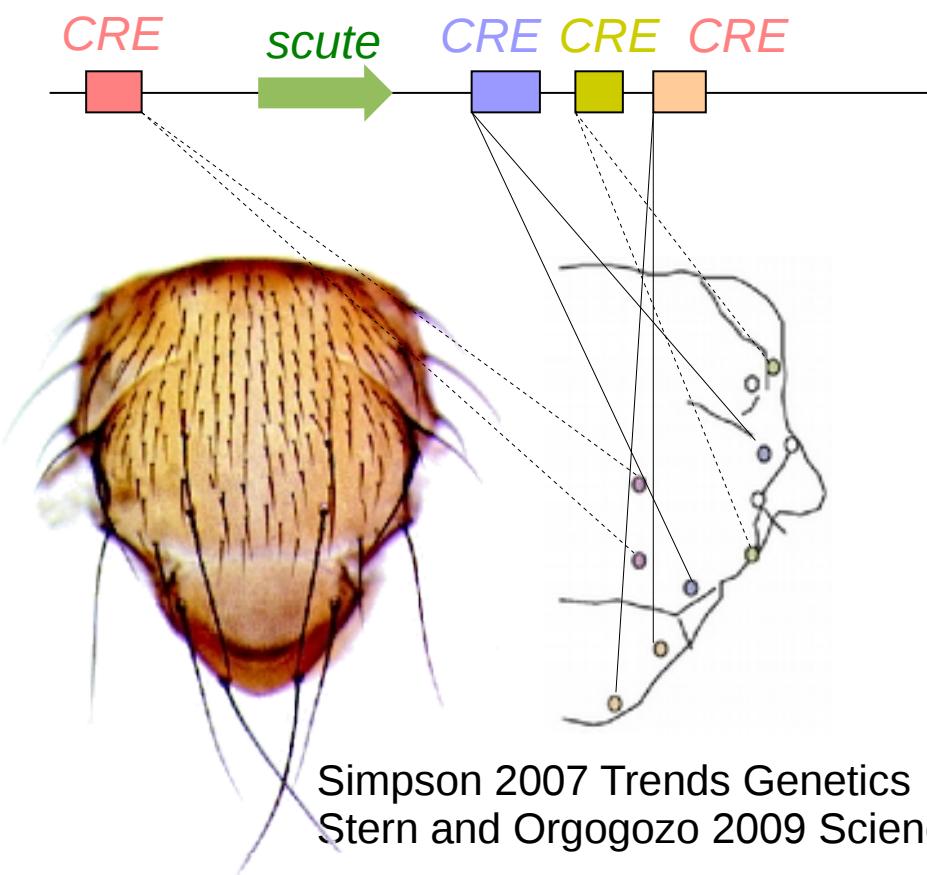
oxygen, photons, insecticide, cholesterol...

Specialized loci in the genome

Proteins that interact with external molecules

oxygen, photons, insecticide, cholesterol...

Cis-regulatory elements of “developmental switch genes”



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

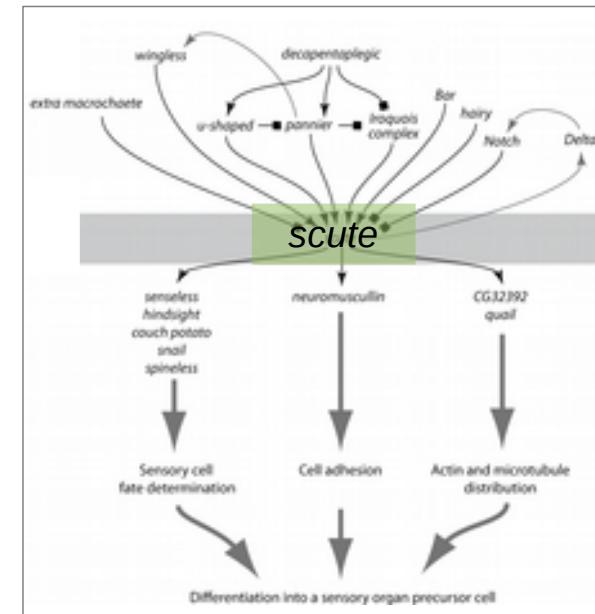
From random processes can emerge predictability

Many unpredictable processes
at a low level



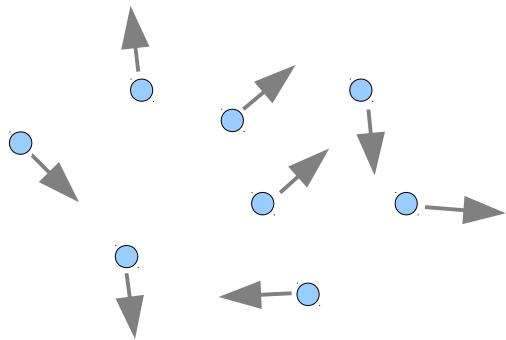
Predictable Evolution
at the genetic level

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



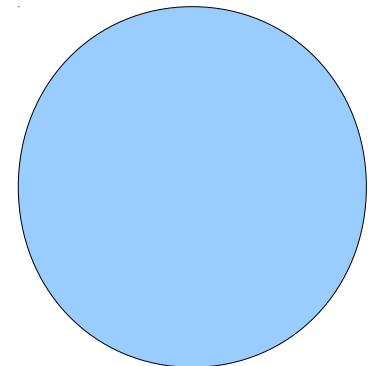
From random processes can emerge predictability

Microscopic world

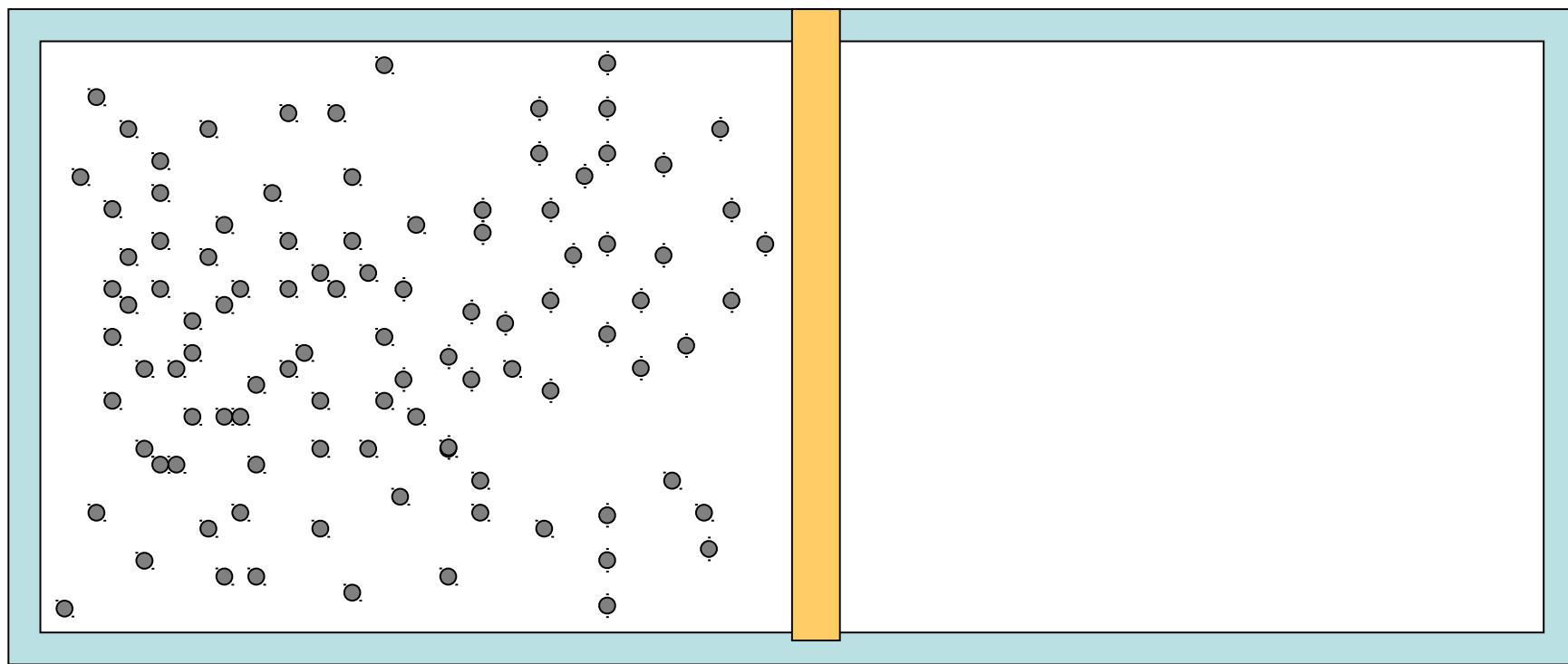


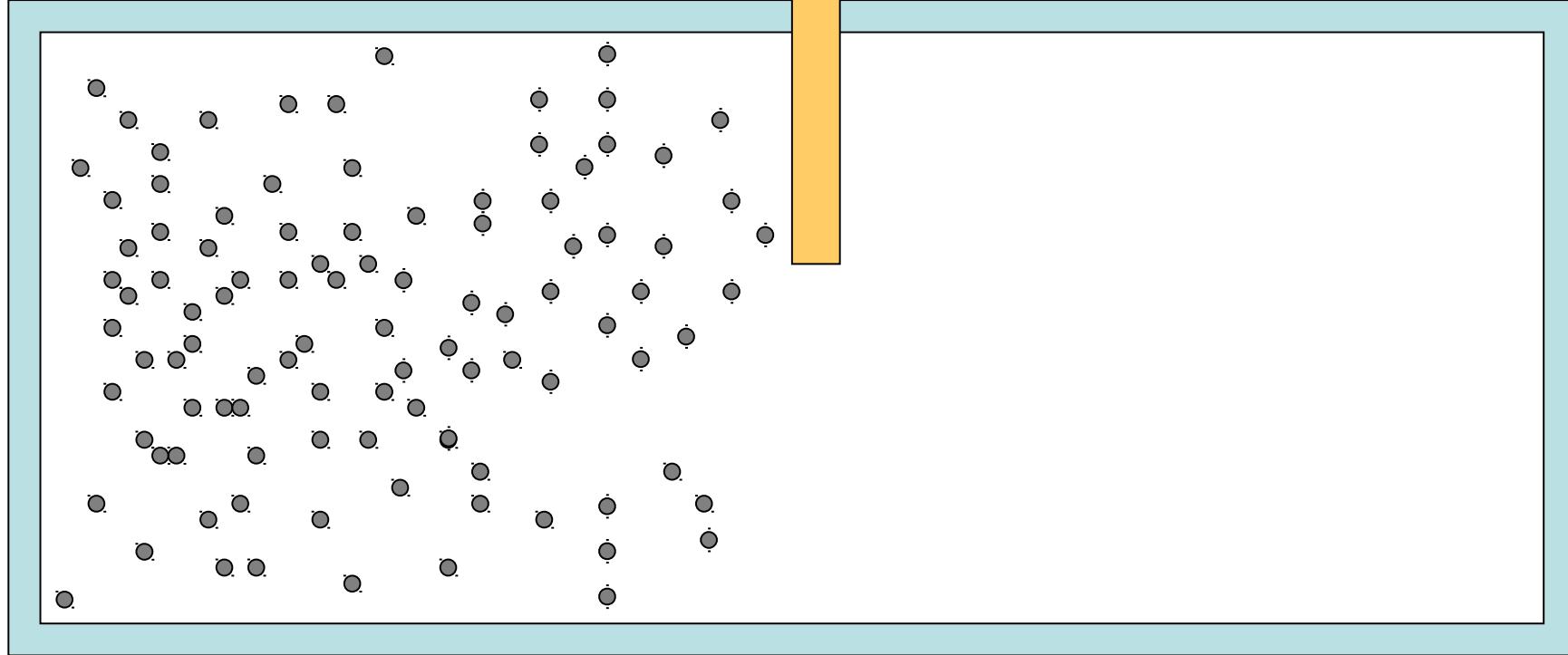
Position, mass, velocity of each particle

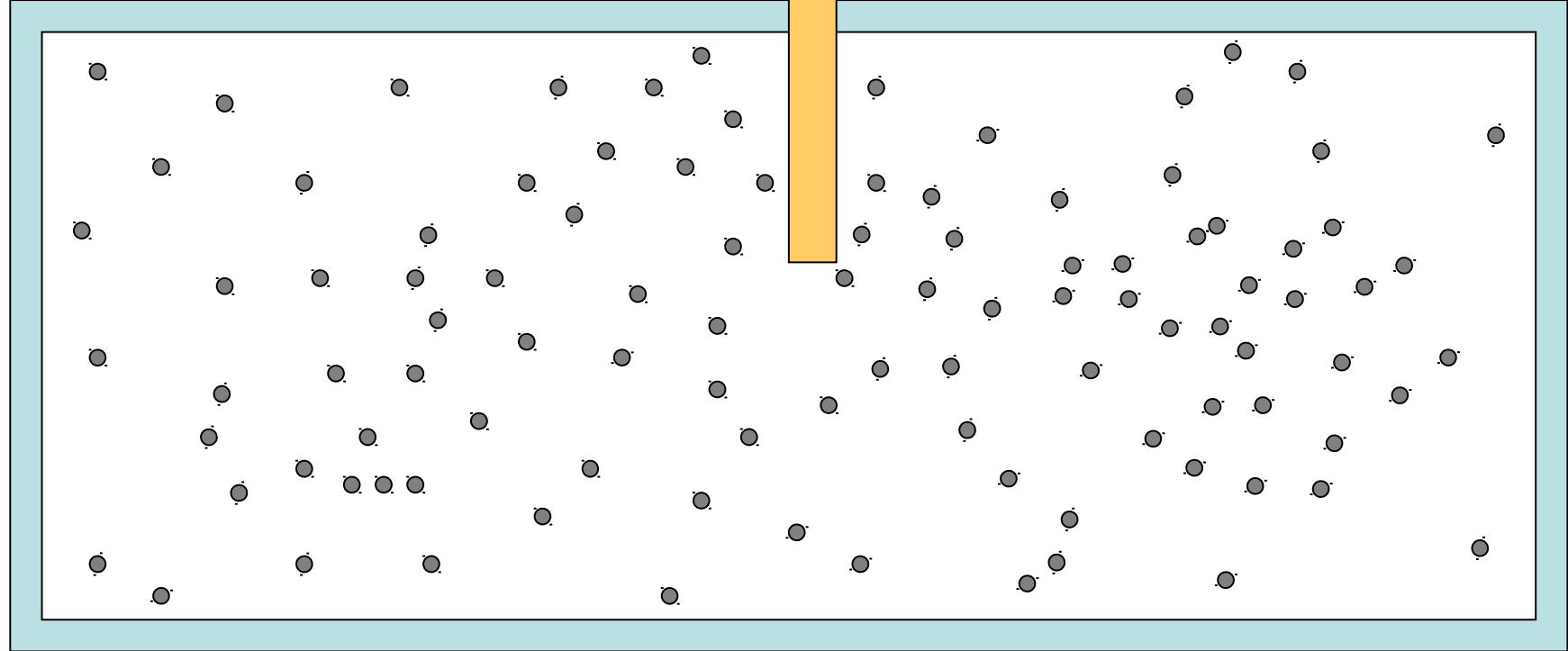
Macroscopic world



Pressure, Volume, Temperature,
Number of moles





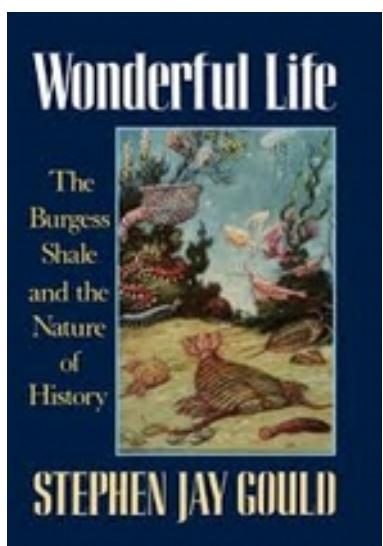


After a few seconds

UNPREDICTABLE ?

Evolution

PREDICTABLE ?



1989

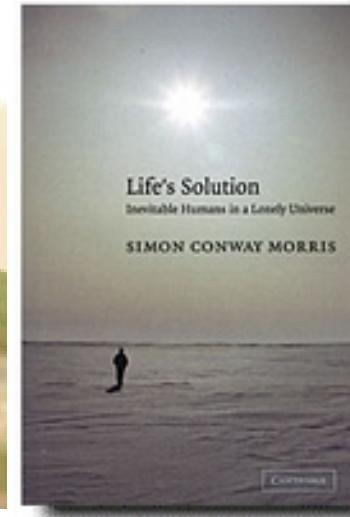
Rare events with important consequences

Evolutionary mechanisms are random

Many possibilities unexplored



S Conway Morris

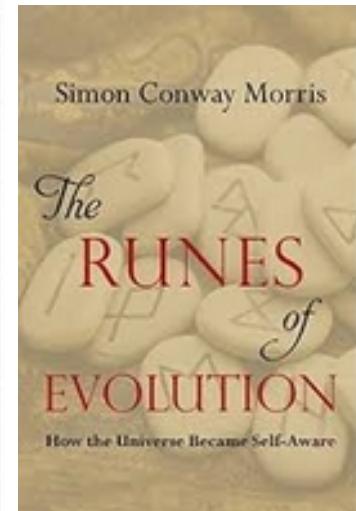


2003

Multiple convergences

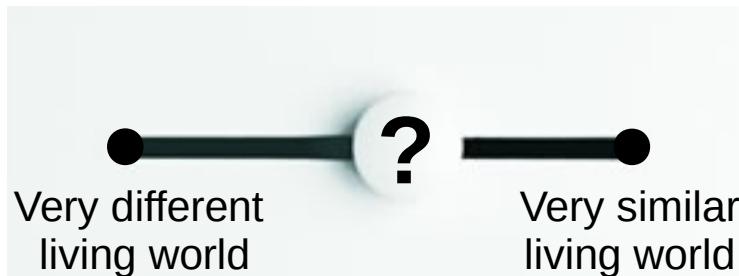
Not that many possibilities

All possibilities explored



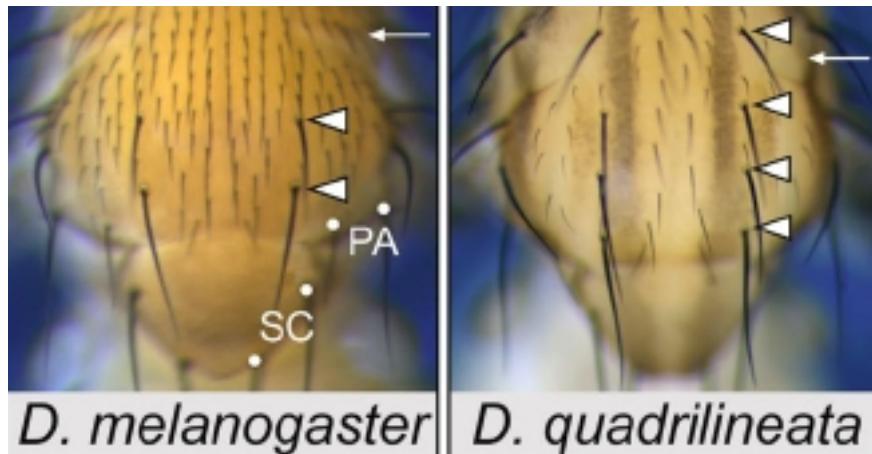
2015

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



Bristle evolution: always *scute* cis-regulatory elements?

1) extra bristles in *D. quadrilineata*

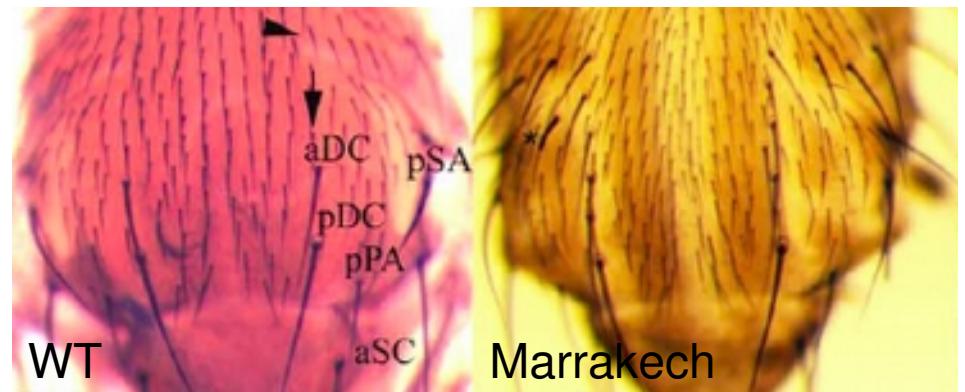


D. melanogaster

D. quadrilineata

Marcelini and Simpson 2006

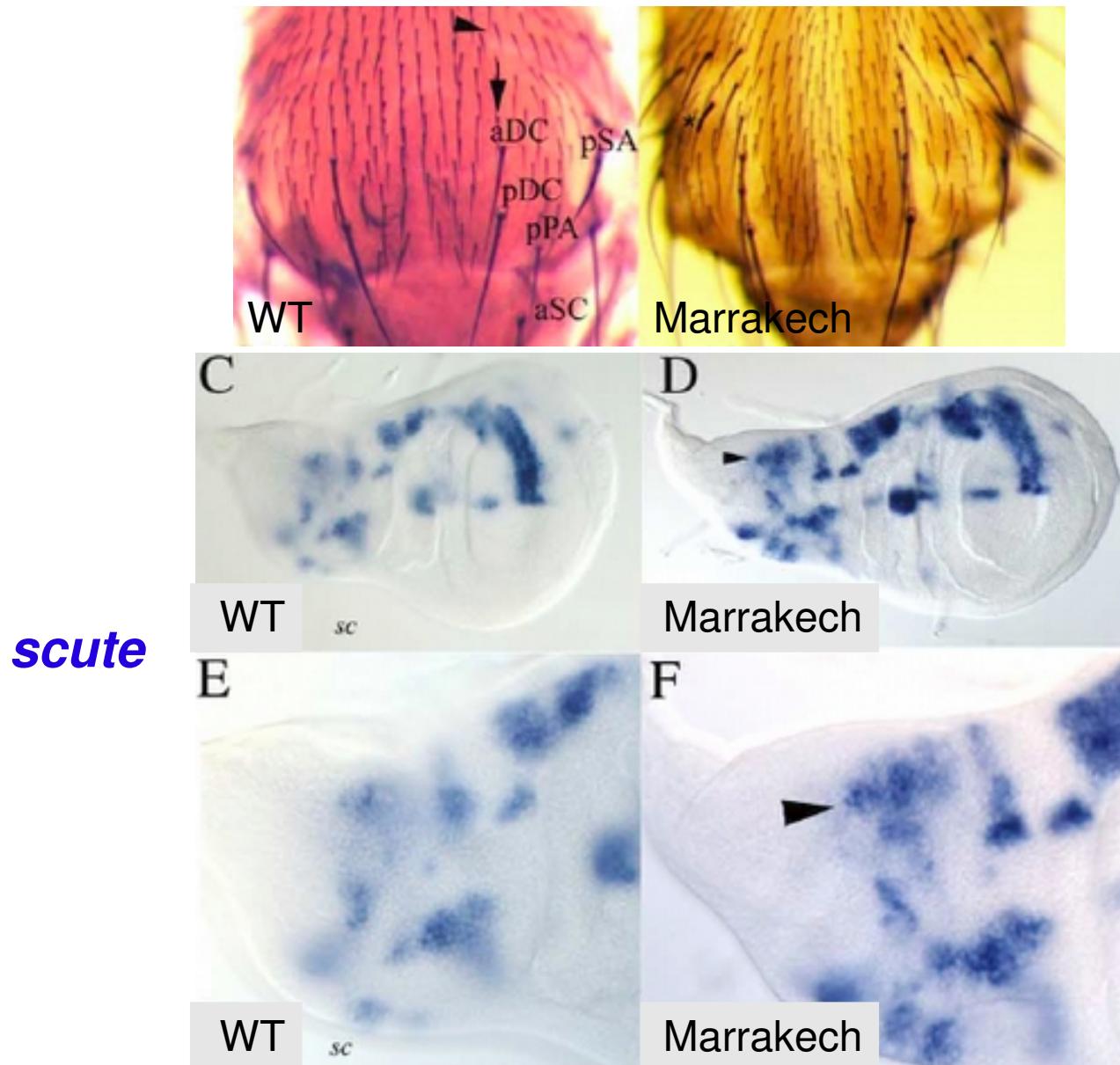
2) extra bristles
in a *D. melanogaster* subpopulation



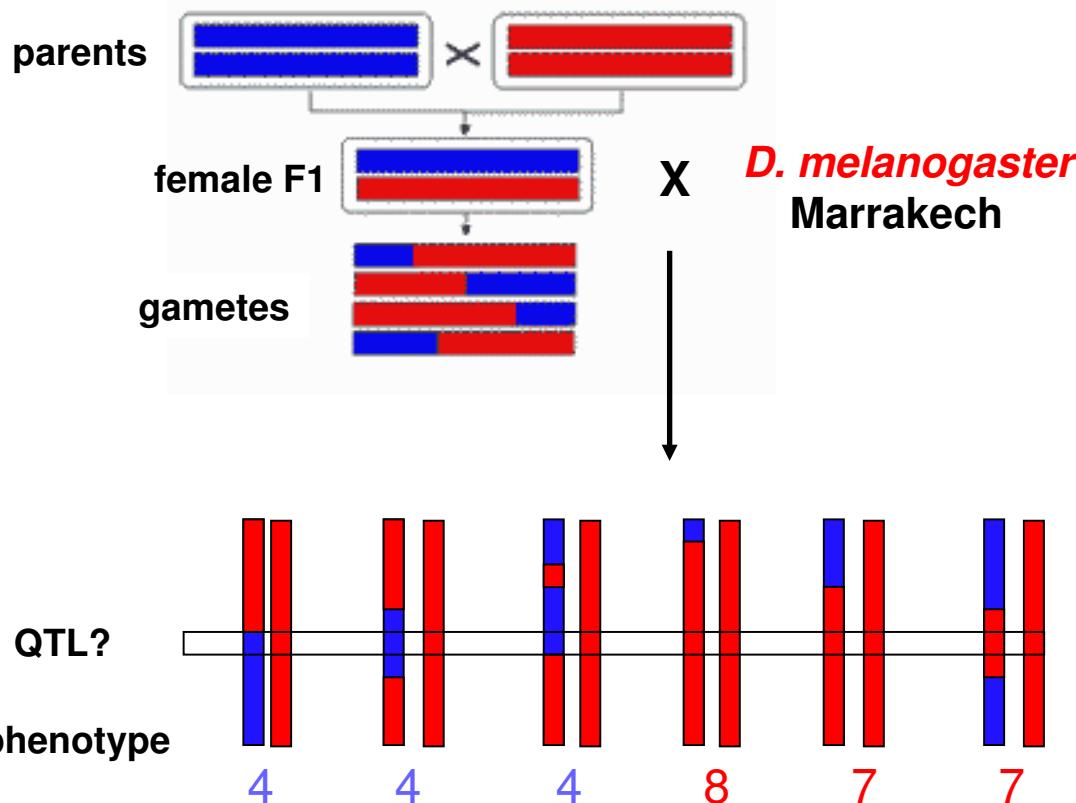
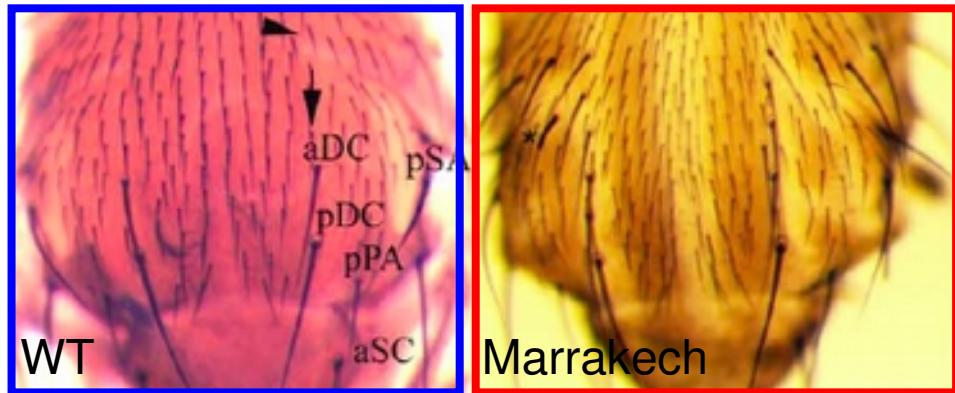
Gibert et al 2005

3) bristle loss in *D. santomea*

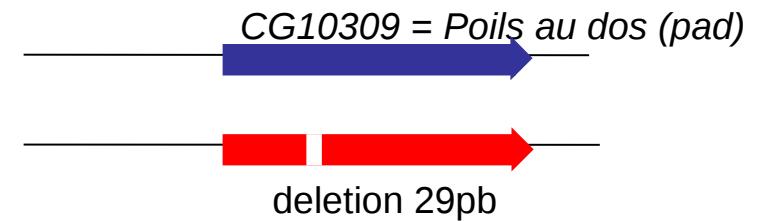
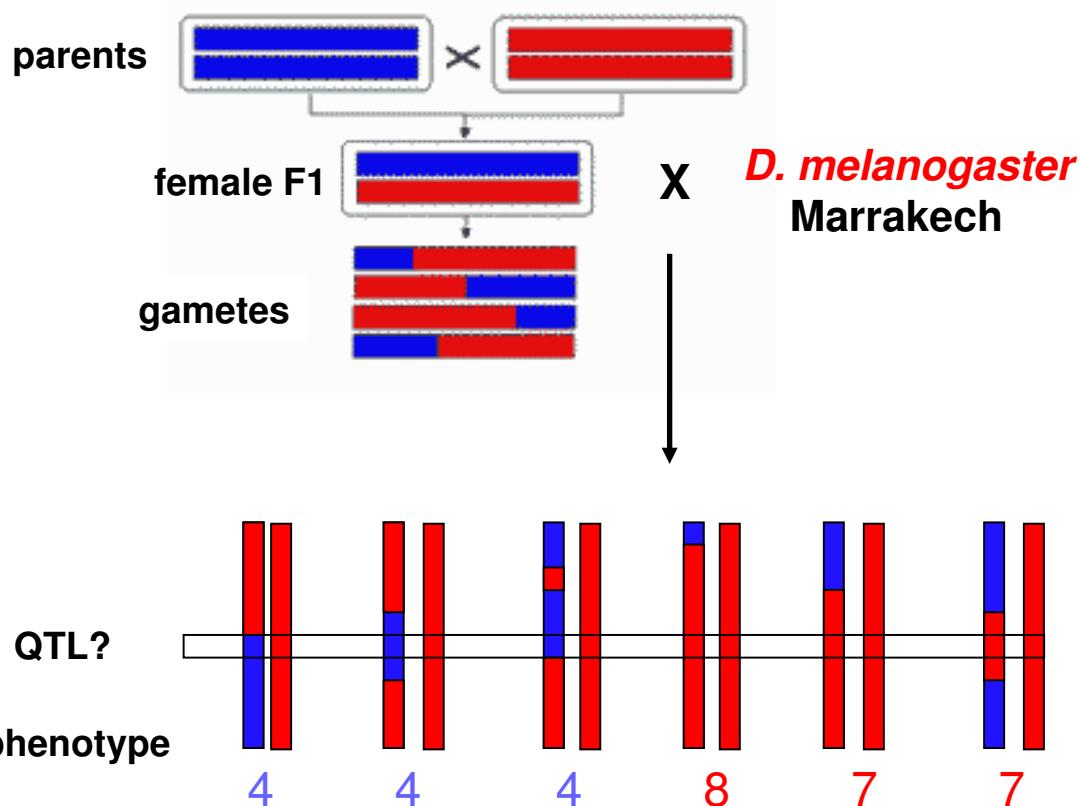
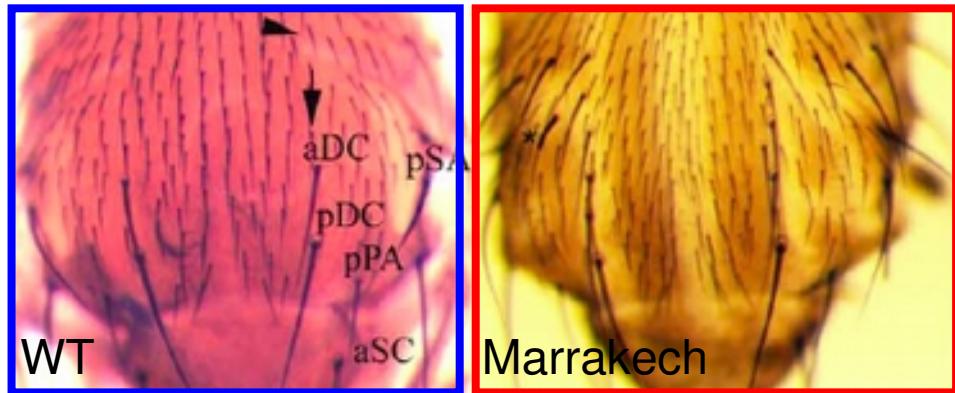
Extra bristles in *D. melanogaster*-Marrakech correlate with larger *scute* expression domain

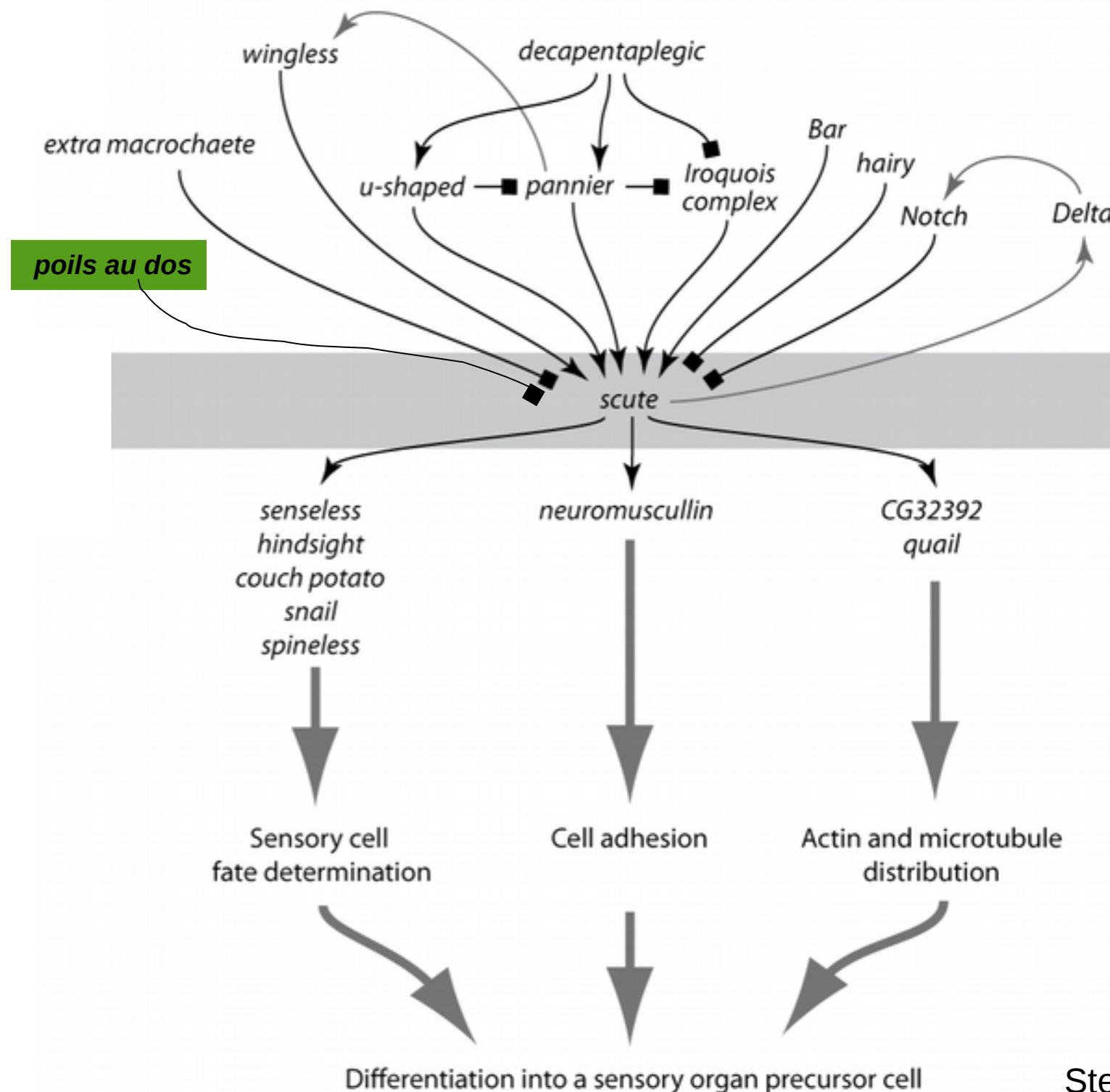


Extra bristles in *D. melanogaster*-Marrakech due to mutation(s) in *poils-au-dos*

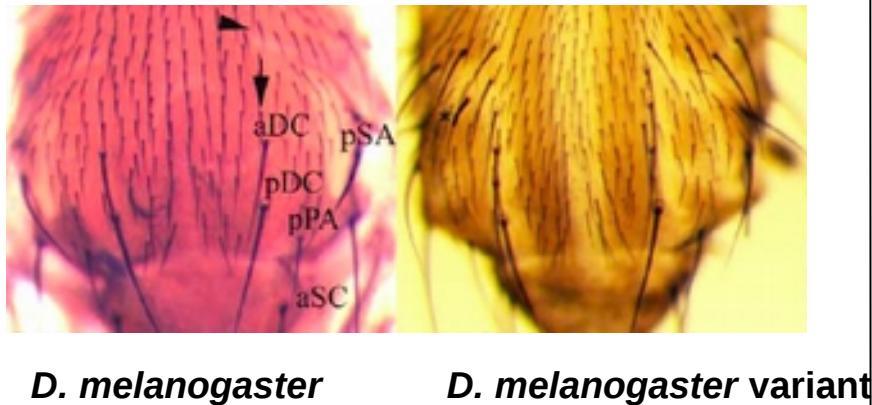


Extra bristles in *D. melanogaster*-Marrakech due to mutation(s) in *poils-au-dos*

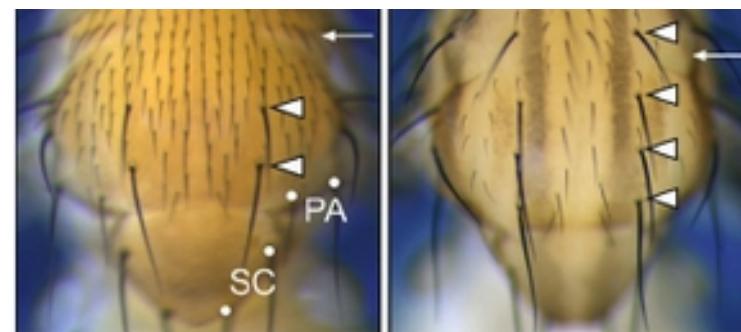




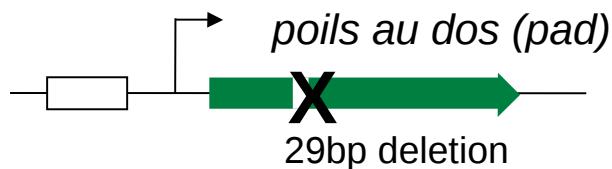
Short-term evolution...



...versus long-term evolution

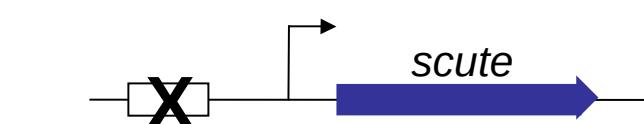


D. melanogaster *D. melanogaster* variant



null mutation in coding region
change in thorax and wing

(Gibert et al., 2005)



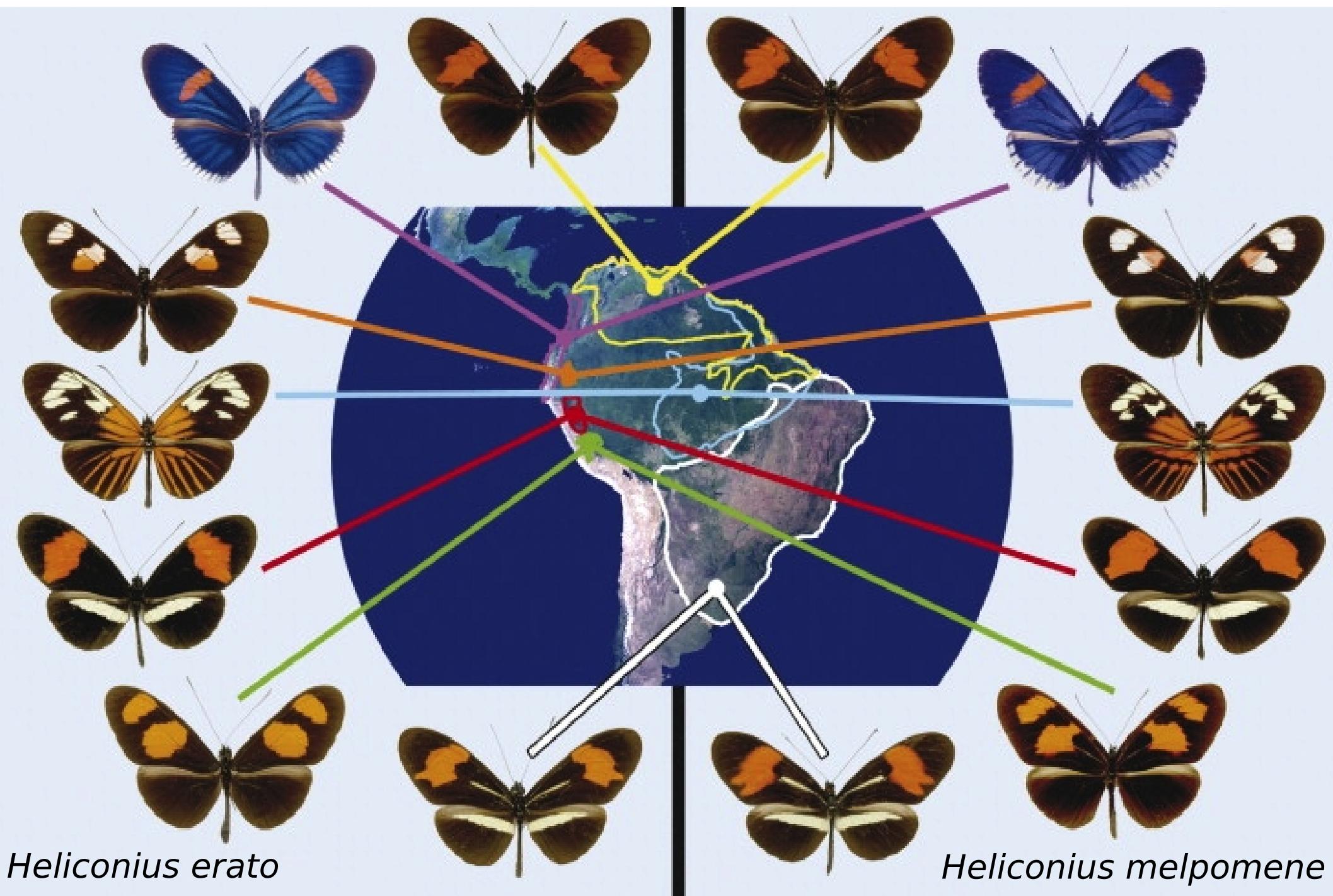
cis-regulatory mutation
change in the thorax only

(Marcellini et al. 2006)



**The tree of life
is not a tree**

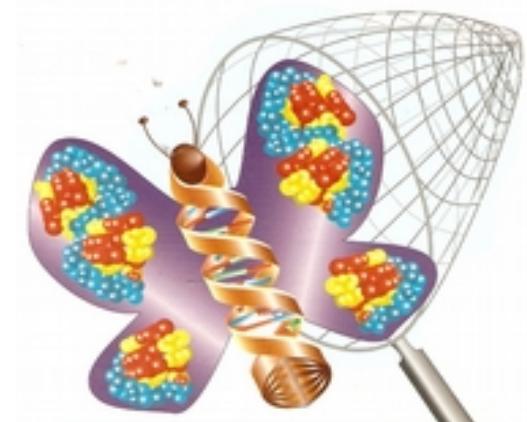
Natural replicates of evolution



Larry Gilbert, UT Austin



**This drawer =
SIBLINGS from a
hybrid cross**



**GENE
HUNT**

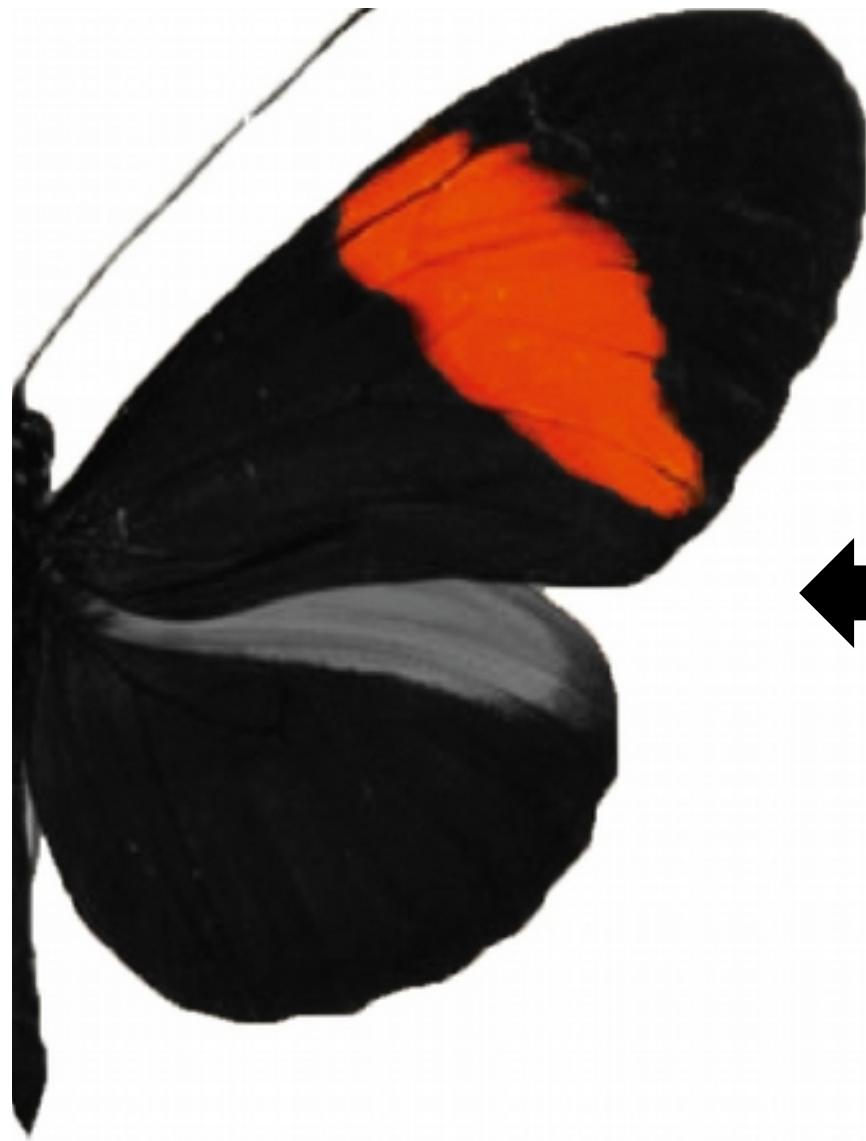


Heliconius melpomene

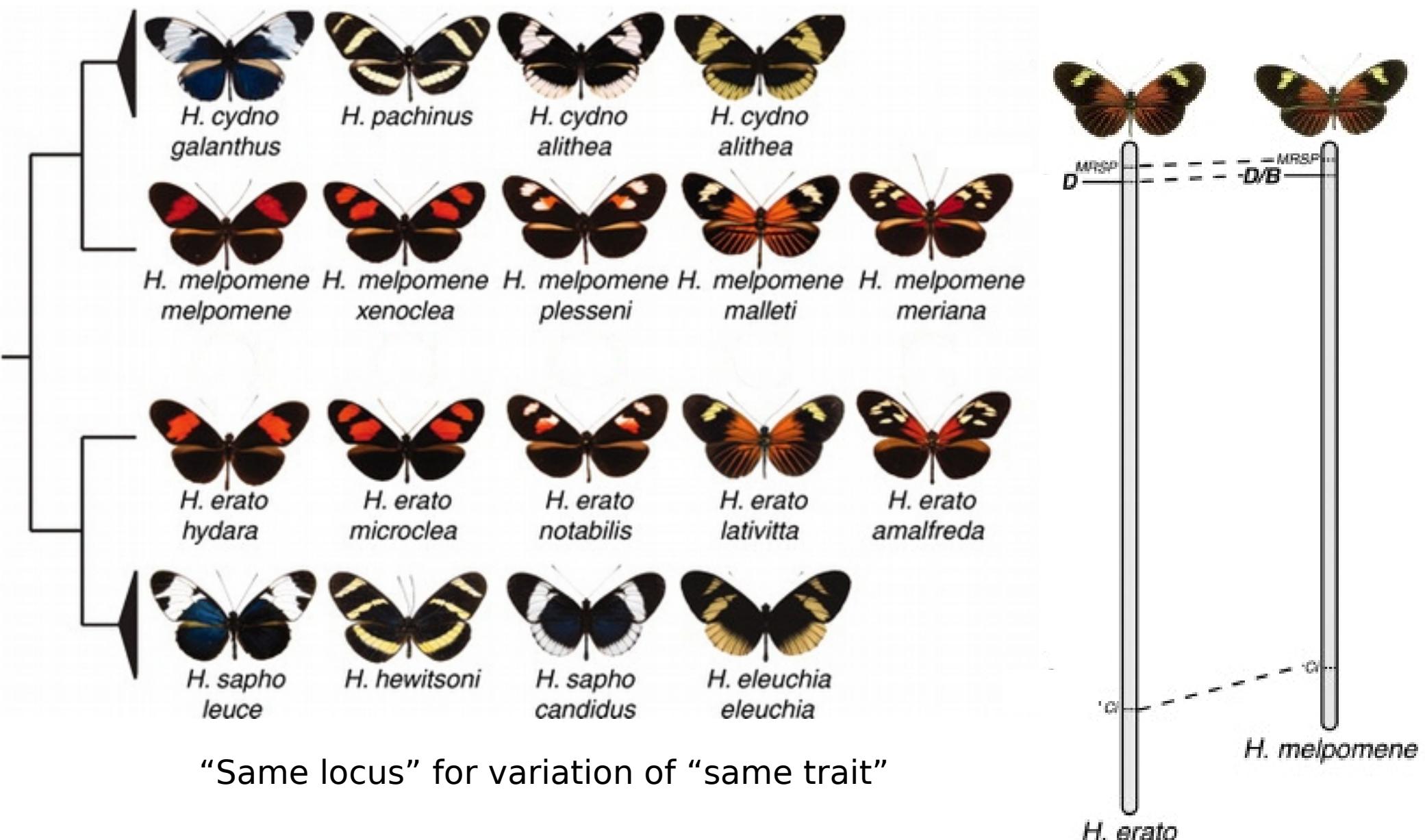


Heliconius erato

Which genes?
Same genes in co-mimics?

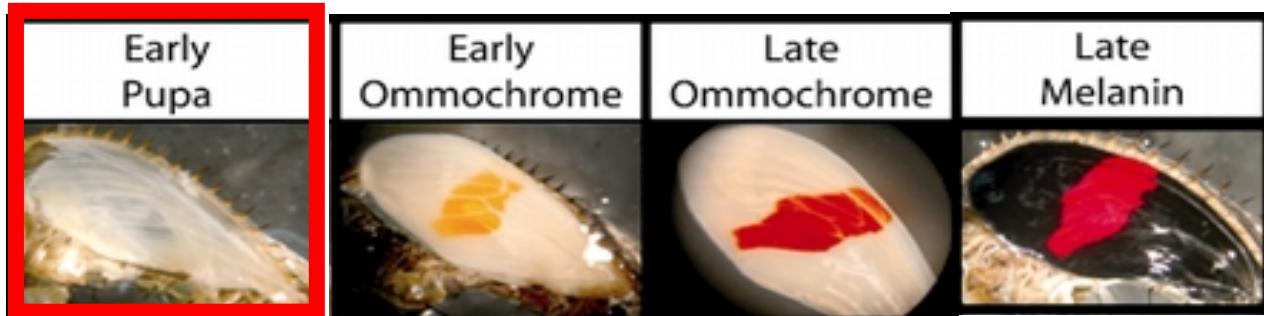
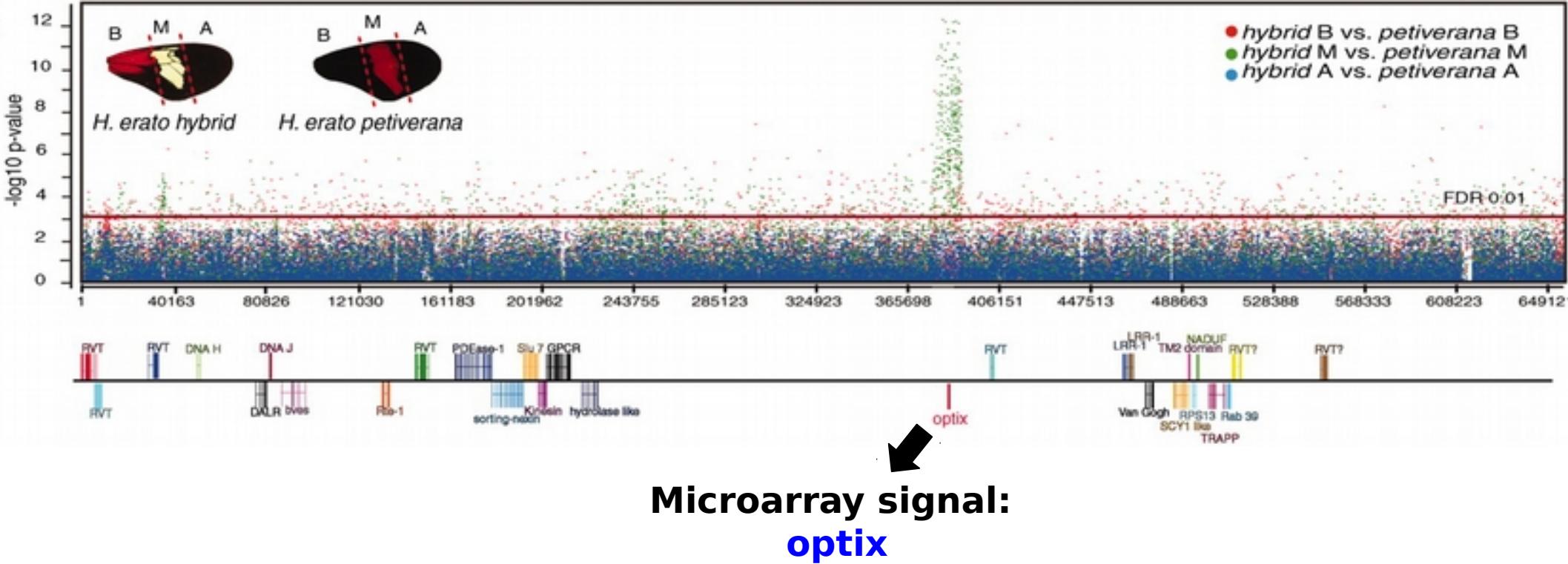


Genetic parallelism in co-mimics

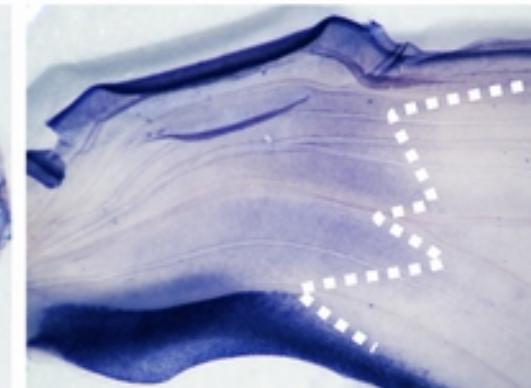
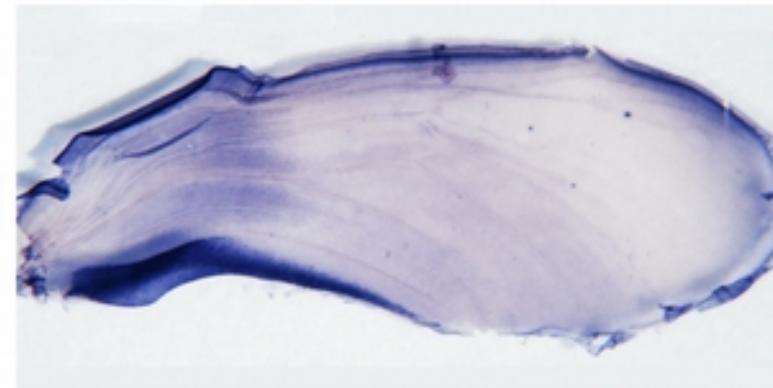


“Same locus” for variation of “same trait”

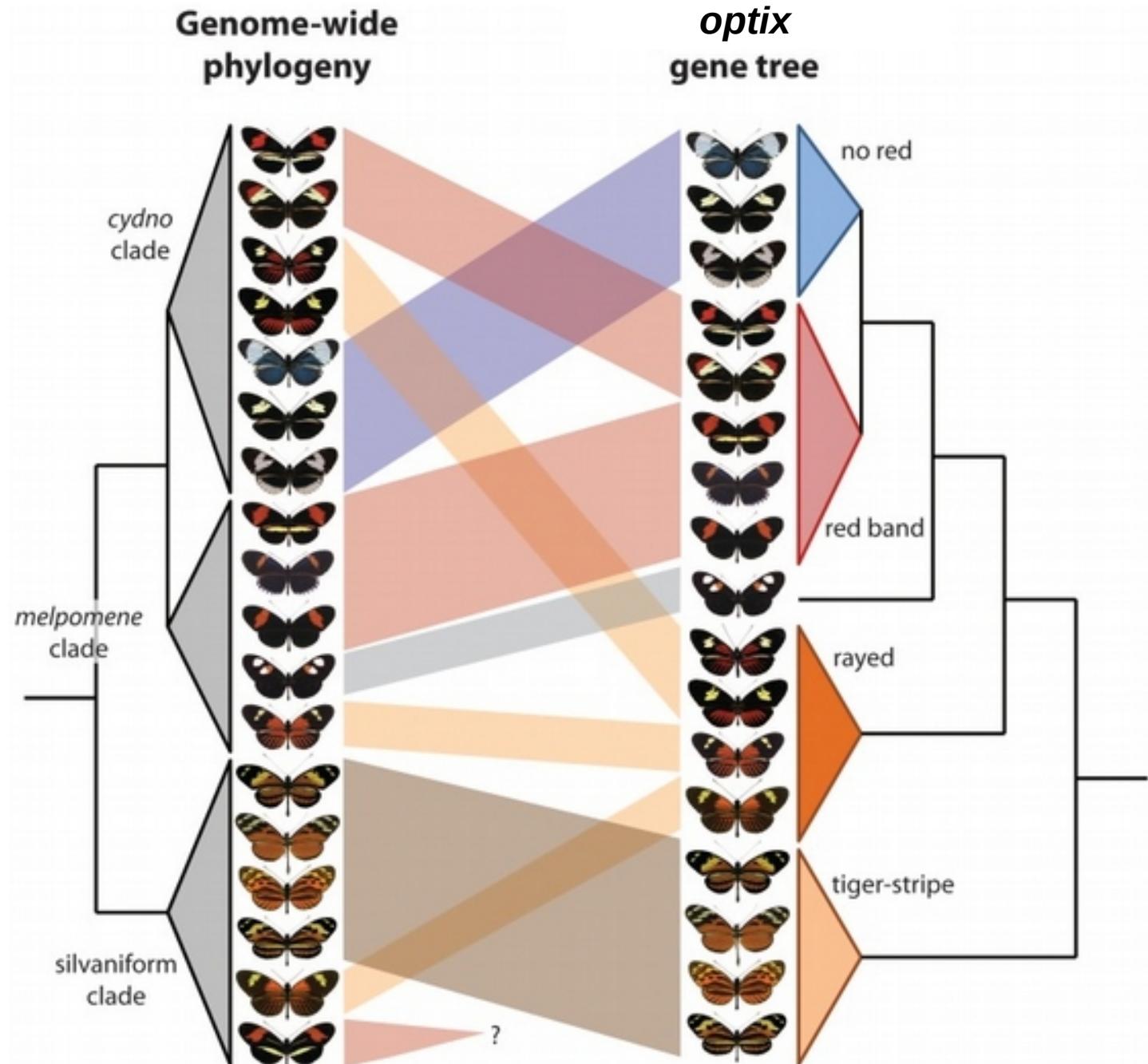
Fine mapping done by the labs of Chris Jiggins and Owen McMillan
Baxter Genetics 2008



Optix marks red scale precursors

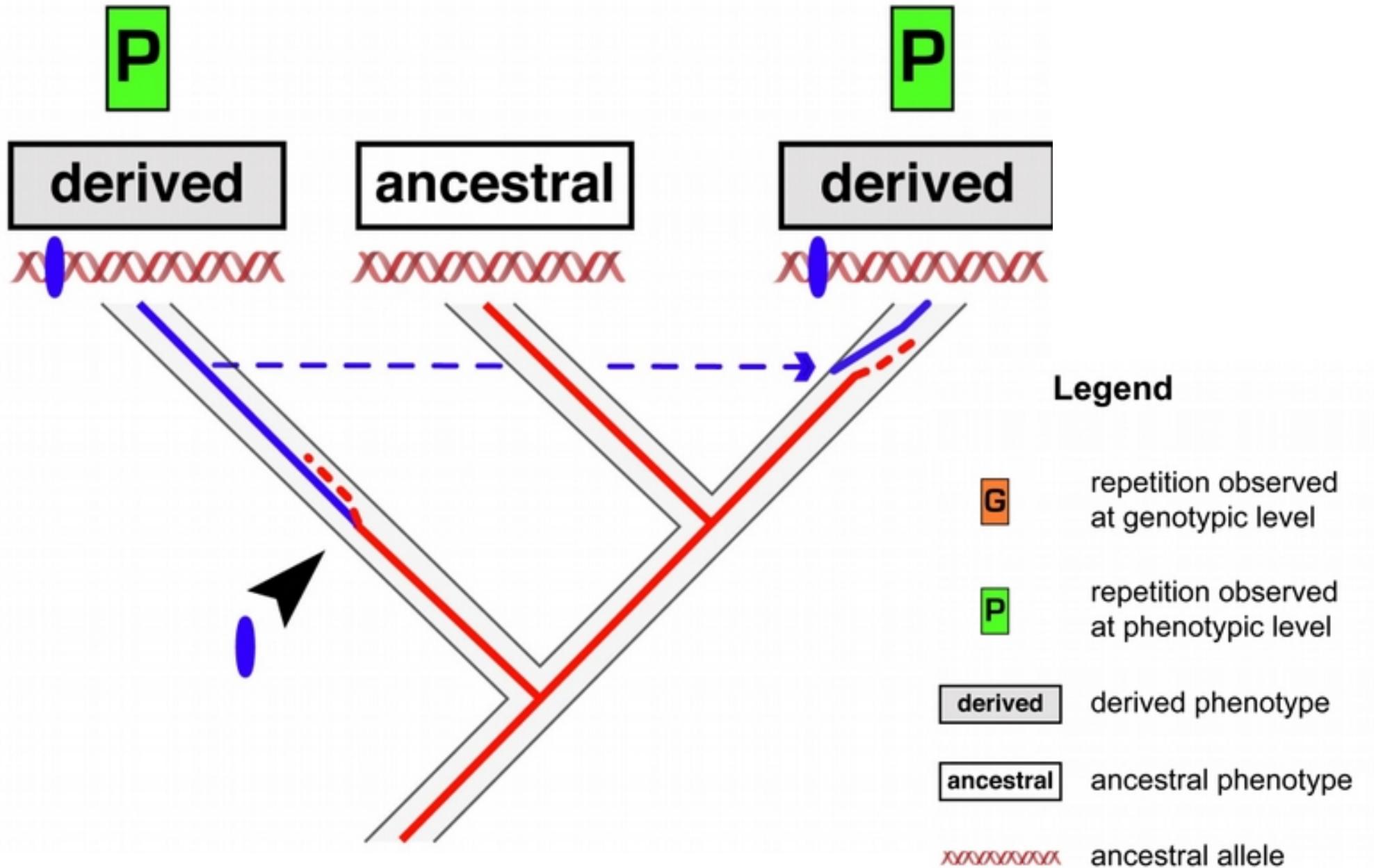


Genetic plagiarism of the red wing color pattern



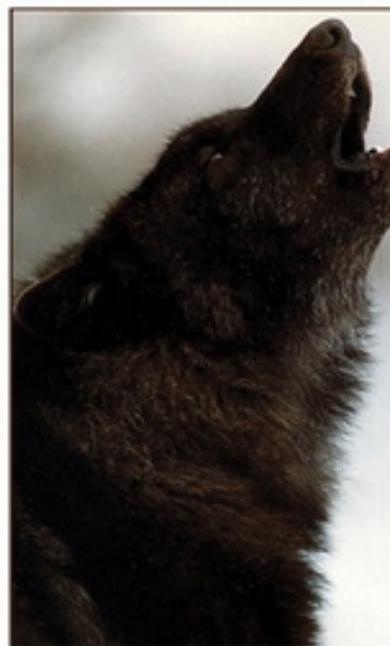
(Reed 2011, Pardo-Diaz 2012,
Heliconius Consortium 2012)

Lateral transfer



Black color in wolves comes from a dog gene

B



10 mo



10 yrs



10 mo

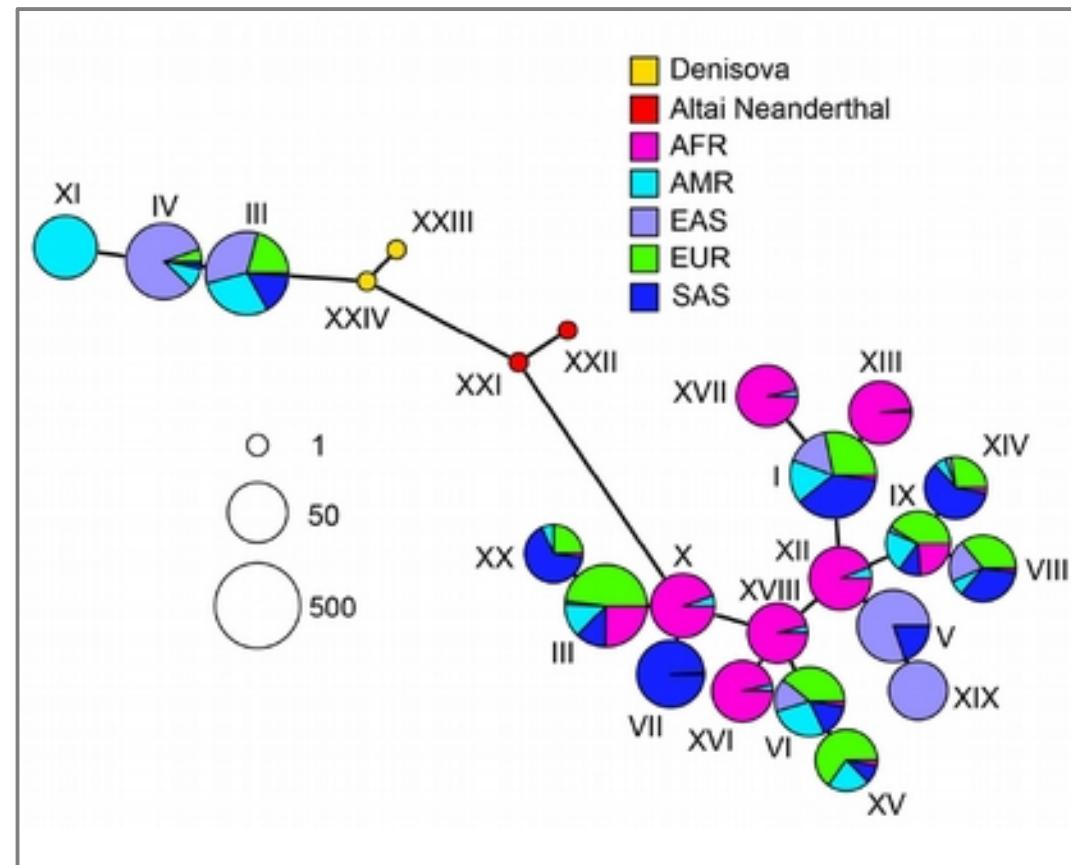
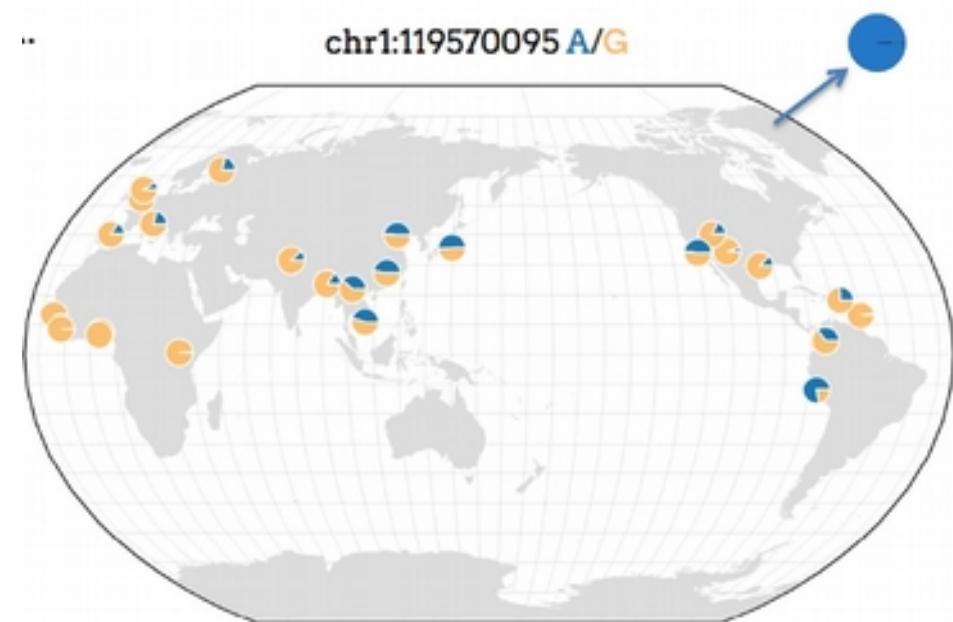


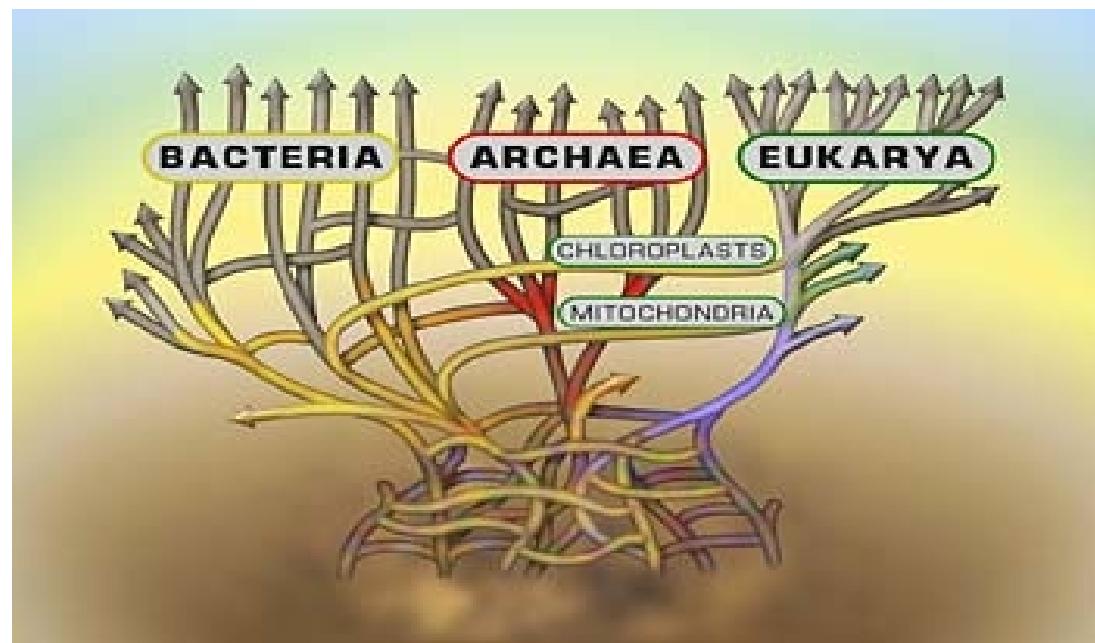
10 yrs

CBD103 ΔG / +
(K^B/k^y)

CBD103 + / +
(k^y/k^y)

Cold Tolerance in Inuits may come from Archaic Denisovan DNA





Conclusion

Our research approach: finding the mutations causing evolution

The genetic program is dead

Evolution is not as random as previously thought

The tree of life is not a tree

Additions

Classical view

