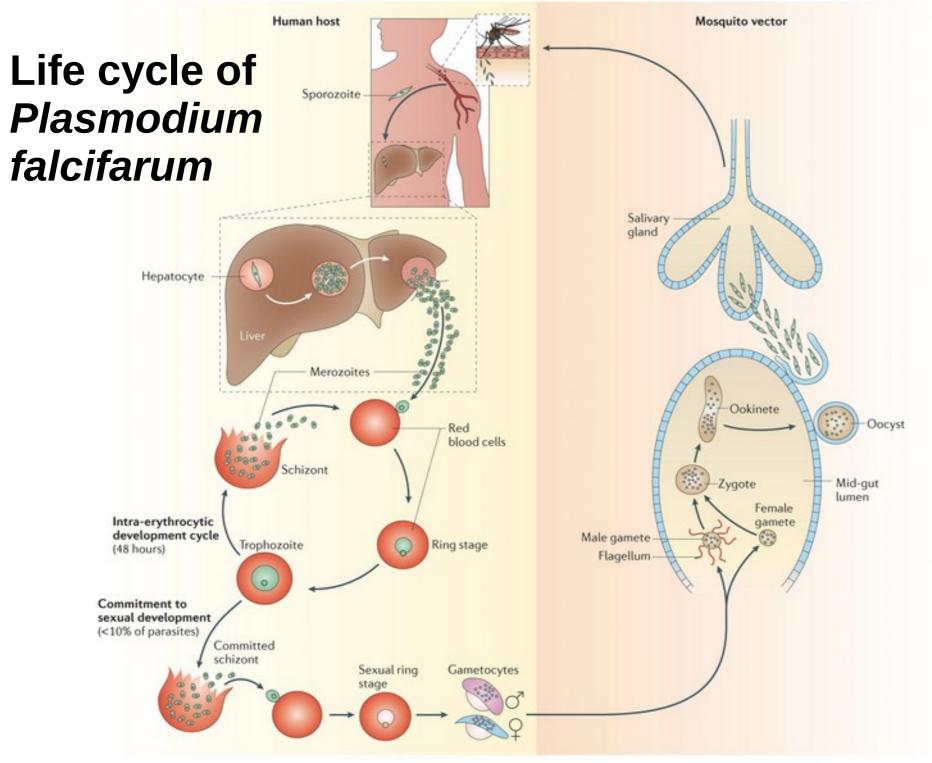
Micro-evo-devo: Genotype-phenotype relationships, convergences, environmental factors

Virginie Courtier-Orgogozo Institut Jacques Monod, Paris

What is evo-devo?

- Development
- Evolution
- Evo-devo





https://www.nature.com/articles/nrmicro3519/figures/1

Nature Reviews | Microbiology

Life cycle of Sacculina carcini



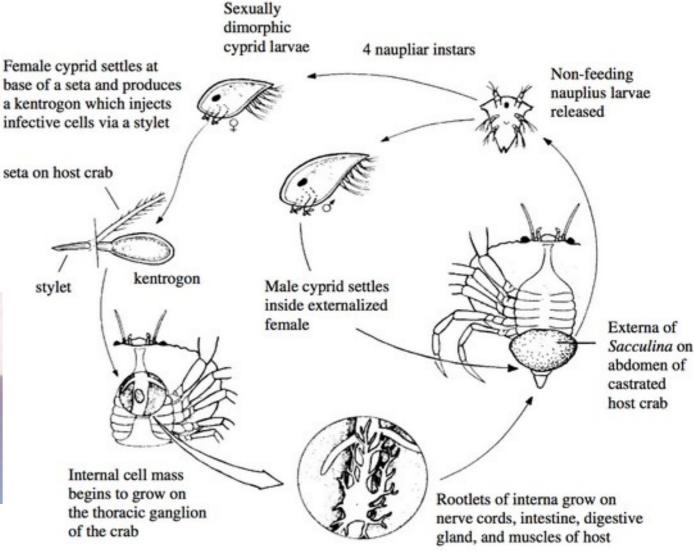


Figure 1. Life cycle of Sacculina carcini modified from Hickman et al. (1988).

Development

process through which a single cell (often a fertilized egg) gives rise to a complex multicellular organism

Frontiers:

Regeneration Unicellular organisms Asexual reproduction Aging Cell death



Examples of subfields: Morphogenesis Organogenesis Cell differentiation

Pradeu, T., Laplane, L., Prévot, K., Hoquet, T., Reynaud, V., Fusco, G., ... & Vervoort, M. (2016). Defining "development". In Current topics in developmental biology (Vol. 117, pp. 171-183). Academic Press.

What is Evolution?

The peppered moth

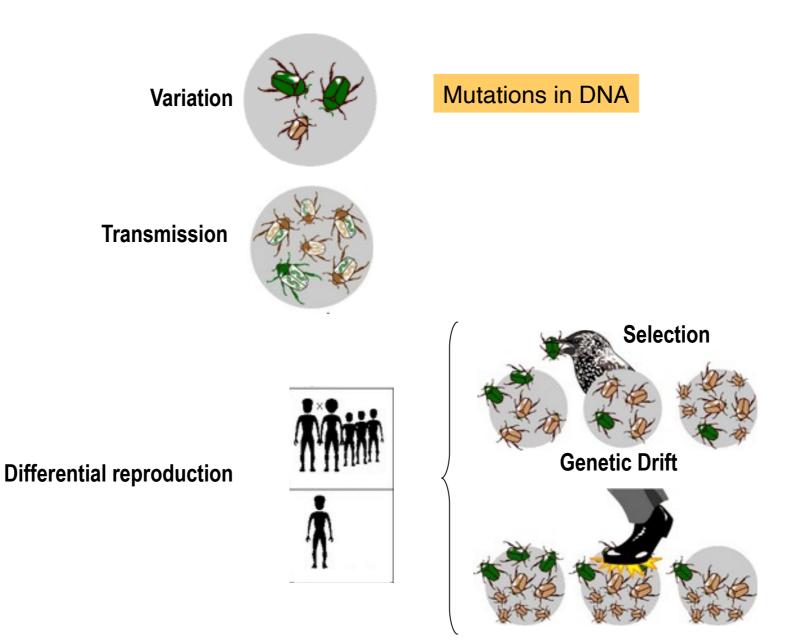
In Manchester, in the early 1800s, nearly all were light coloured:



By 1895, nearly all were dark:

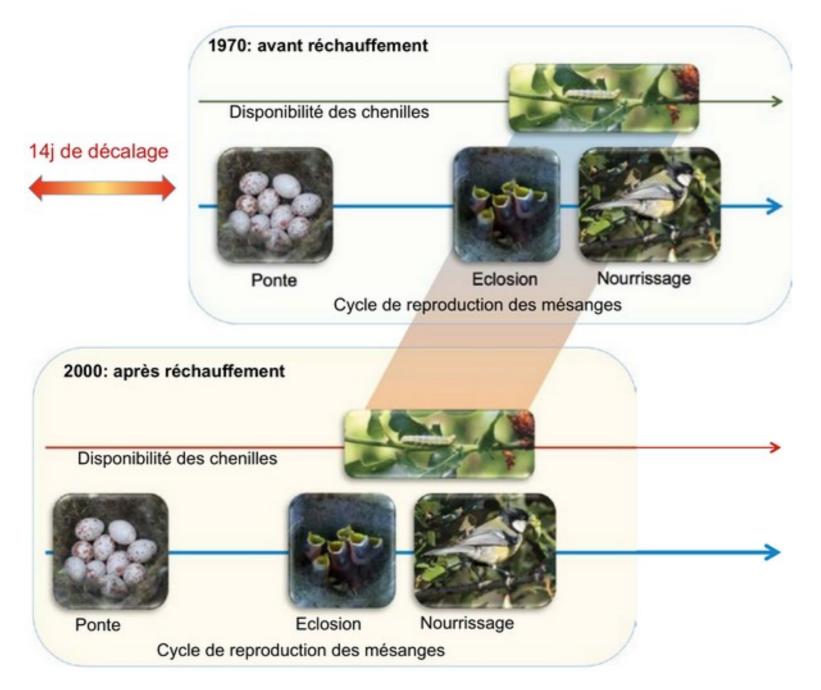


Classical Darwinian Evolution



http://evolution.berkeley.edu

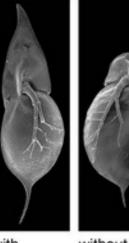
Rapid evolution of great tits



Etonnant vivant : découvertes et promesses du XXIe siècle (2017)

Plasticity: one genotype \rightarrow several phenotypes

Daphnia



with

helmet

without helmet

Nemoria arizonaria caterillars



spring: caterpillars feed on catkins



summer: caterpillars feed on leaves

Water crowfoot plant



leaves growing above water

leaves growing below water

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

Daphnia: Agrawal et al (1999)

Nemoria arizonaria caterillars: Sadava et al (2014)

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

Desert locusts



solitary



gregarious

Commodore butterfly

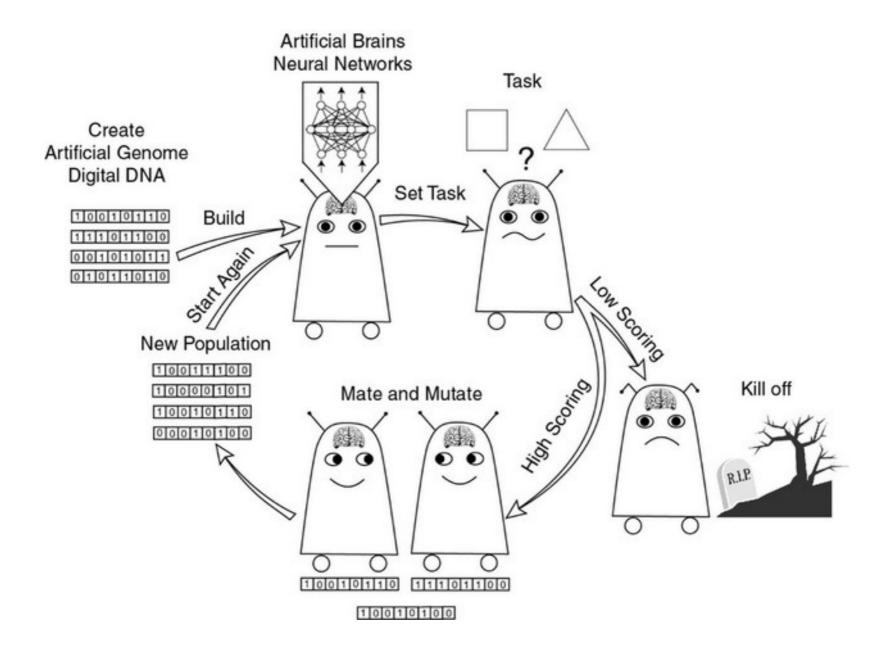




winter

summer





Evolution

process through which the characteristics of populations or species change over successive generations

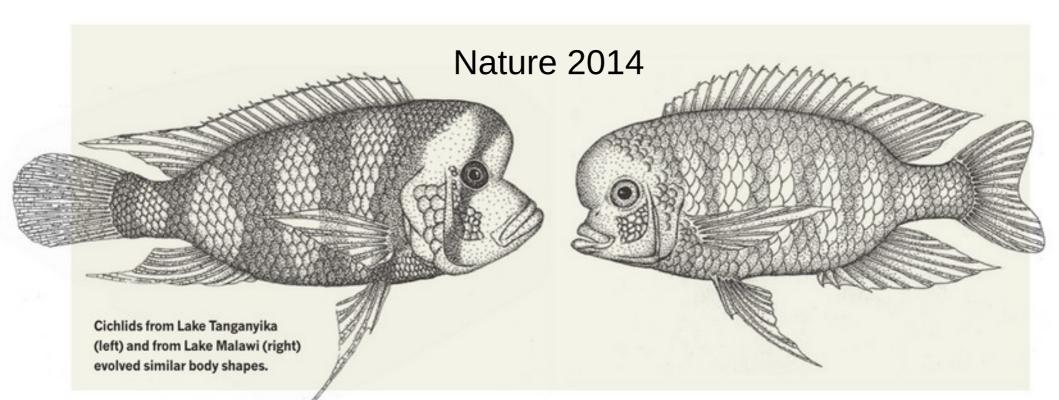
Frontiers:

Origin of Life Evolutionary robotics Plasticity

Examples of subfields:

Paleontology Experimental evolution Modelling of Evolution





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.

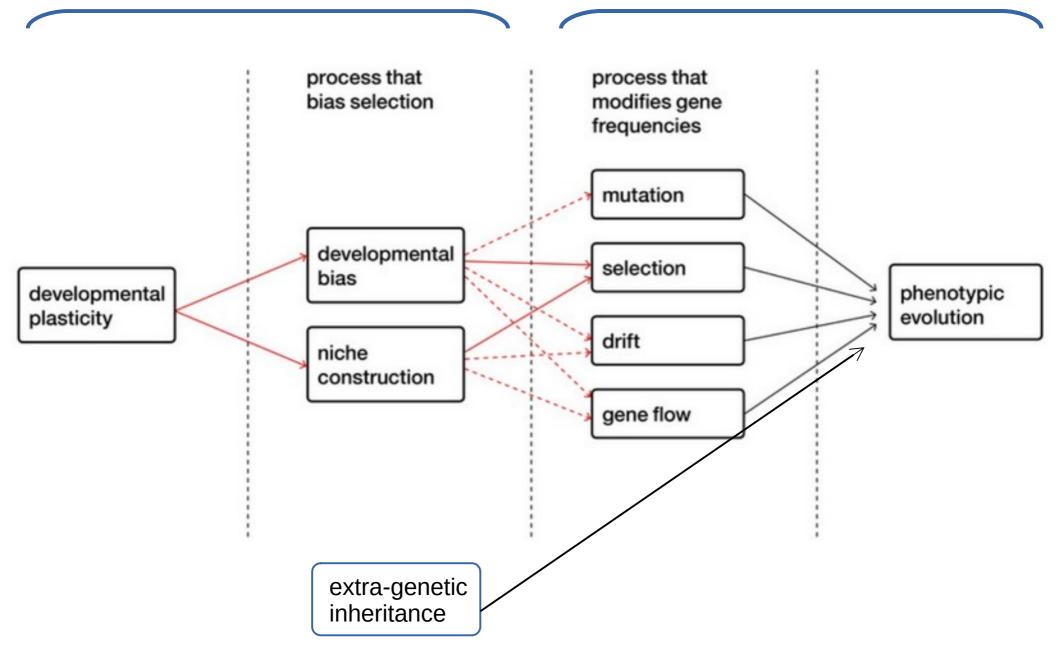
COUNTERPOINT No, all is well

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

Extended evolutionary synthesis

Additions

Classical view



Developmental Bias

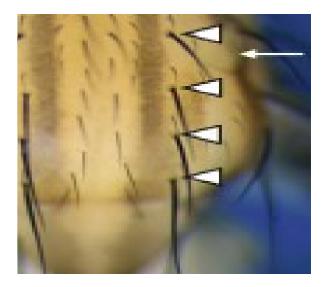
Standing variation

D. melanogaster

► variation no variation

Natural evolution

D. quadrilineata

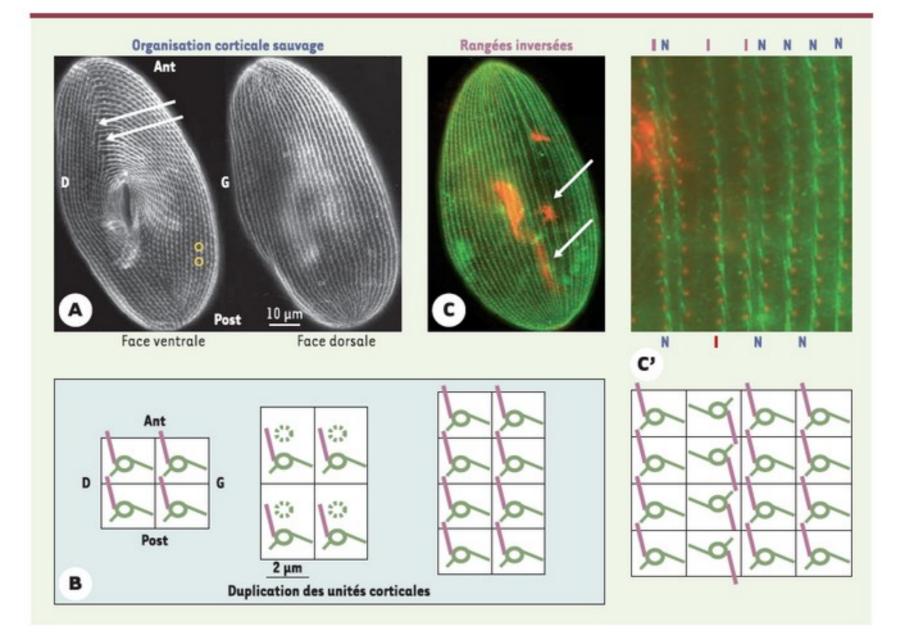


Marcellini et al 2006 PloS Biol

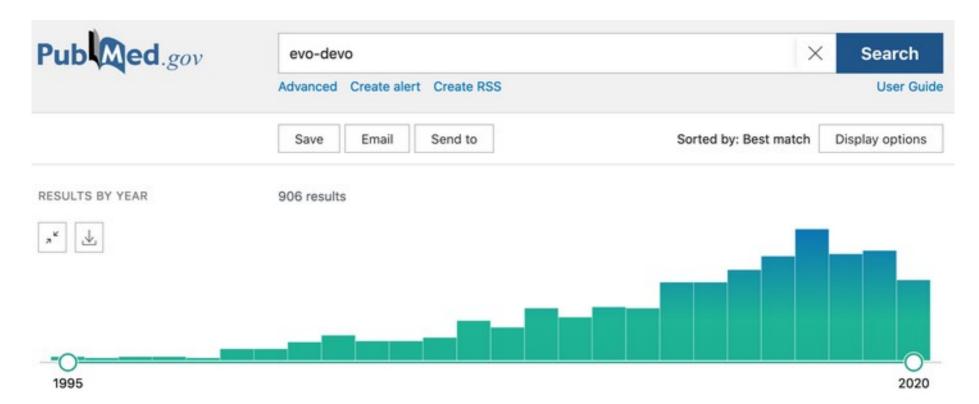




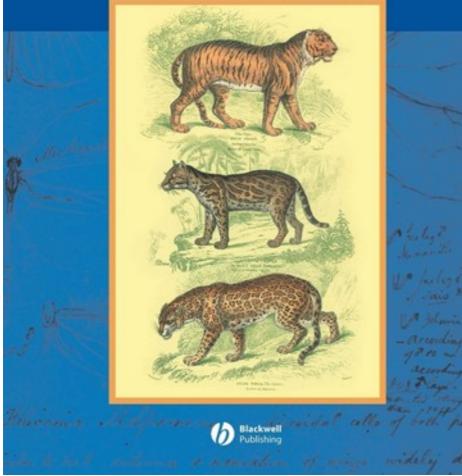
Cortical heredity in Paramecium

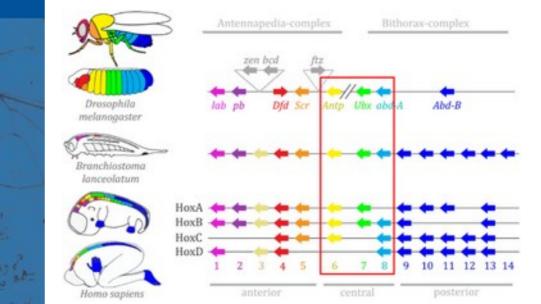


What is Evo-Devo?



SEAN B. CARROLL JENNIFER K. GRENIER SCOTT D. WEATHERBEE FROM DNA TO DIVERSITY MOLECULAR GENETICS AND THE EVOLUTION OF ANIMAL DESIGN SECOND EDITION





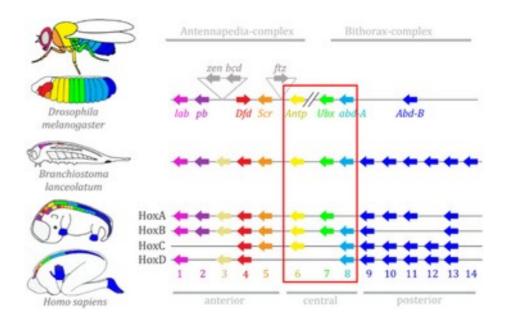
2001

Evo-devo

field of biological research that compares development between species to understand how they evolved

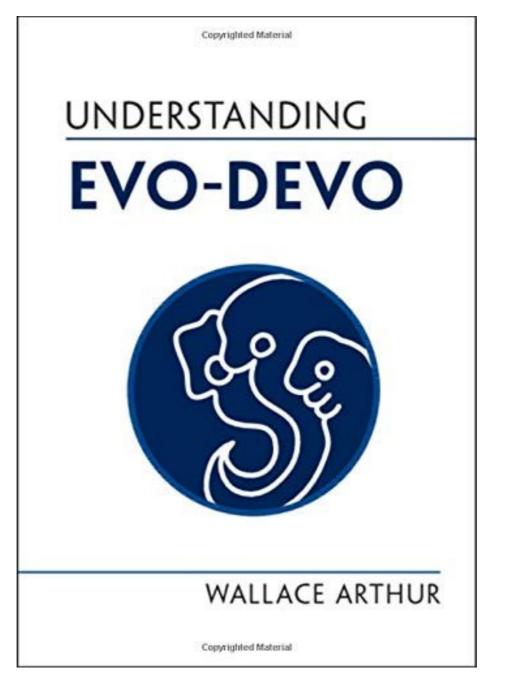
Frontiers:

Same as development and evolution Ecology Physiology, Behavior



subfields:

Macro-evo-devo (distantly related species) Micro-evo-devo (closely related species or populations)



April 2021

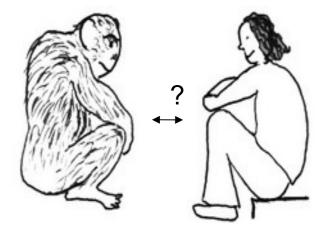
Micro-evo-devo in Drosophila

What makes us different?

between individuals

between species

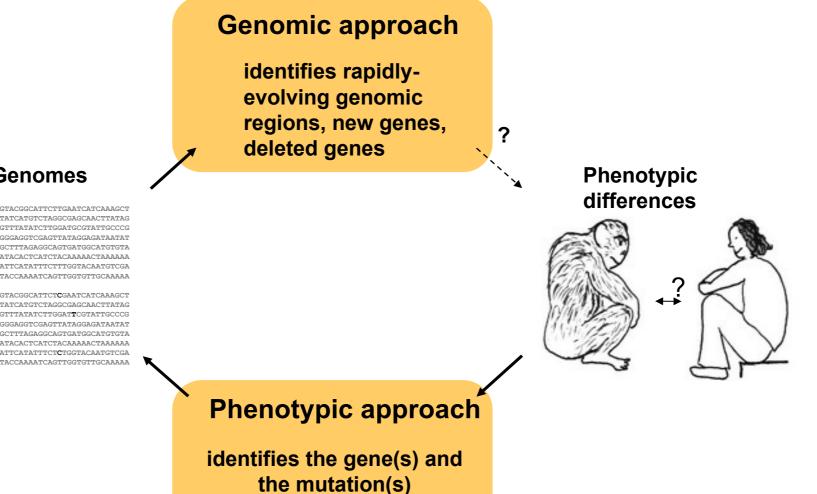




Where do we come from?

What are the mutations responsible for phenotypic differences?

responsible for a phenotypic change



Genomes

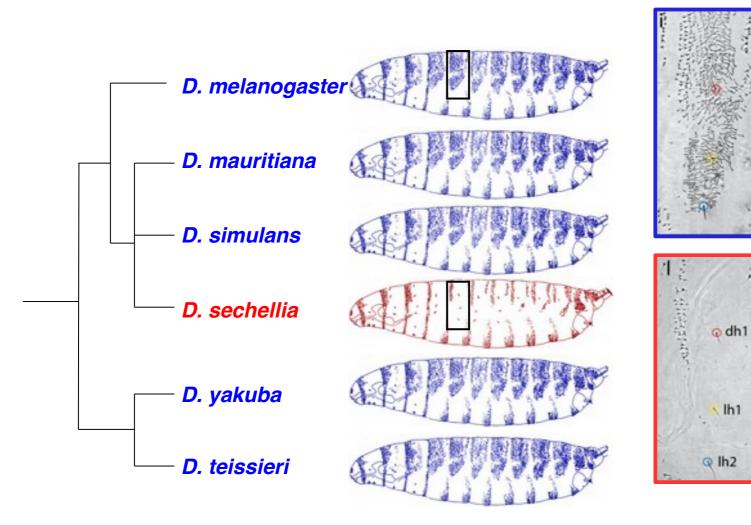
CCTCCTCCATACCCAAATGGATGGTACGGCATTCTTGAATCATCAAAGCI TAGAGCGGGGGAATCGAAGCATATATCATGTCTAGGCGAGCAACTTATAG TGTTCCGTTCCCAAGCTGGTGAAGTTTATATCTTGGATGCGTATTGCCCG CACTTGGGCGCTAATTTGAGTAAGGGAGGTCGAGTTATAGGAGATAATAT TGAATGTCCCTTTCACCACTGGAGCTTTAGAGGCAGTGATGGCATGTGTA CCAATATTCCCTACAGCAGCAATATACACTCATCTACAAAAACTAAAAAA TGGACCTCCACCGAAGTGAATGGATTCATATTTCTTTGGTACAATGTCGA AGAATCTGAAGTTCCGTGGAATATACCAAAATCAGTTGGTGTTGCAAAAA

CCTCCTCCATACCCAAATGGATGGTACGGCATTCTCGAATCATCAAAGCT TAGAGCGGGGGGAATCGAAGCATATATCATGTCTAGGCGAGCAACTTATAG TGTTCCGTTCCCAAGCTCGTGAAGTTTATATCTTGGATTCGTATTGCCCG CACTTGGGCGCTAATTNGAGTAAGGGAGGTCGAGTTATAGGAGATAATAT TGAATGTCCCTTTCACCACTGGAGCTTTAGAGGCAGTGATGGCATGTGTA CCAATATTCCCTACAGCAGCAATATACACTCATCTACAAAAAACTAAAAAA TGGACTTCCACCTAAGTGAATGGATTCATATTTCTCTGGTACAATGTCGA AGAATCTGAAGTTCCGTGGAATATACCAAAATCAGTTGGTGTTGCAAAAA

Hair pattern in flies



Succena and Stern 2000 PNAS McGregor, Orgogozo et al. 2007 Nature Frankel, Wangl and Stern 2012 PNAS

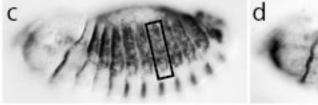


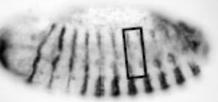
The causing mutation is on the X chromosome

The causing mutation is on the X chromosome

Evolution caused by a change in the svb gene

- **1** Transcription factor that promotes trichome formation
- **2** Correlation between expression pattern and phenotype





D. melanogaster D. sechellia



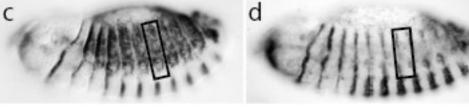
D. melanogaster

D. sechellia

The causing mutation is on the X chromosome

Evolution caused by a change in the svb gene

- Transcription factor that promotes trichome formation
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D. melanogaster D. sechellia



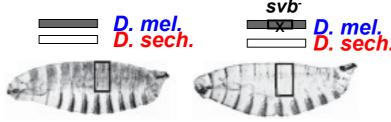
D. melanogaster

D. sechellia

Complementation assay with *D. melanogaster svb^{wt}* or *svb*⁻

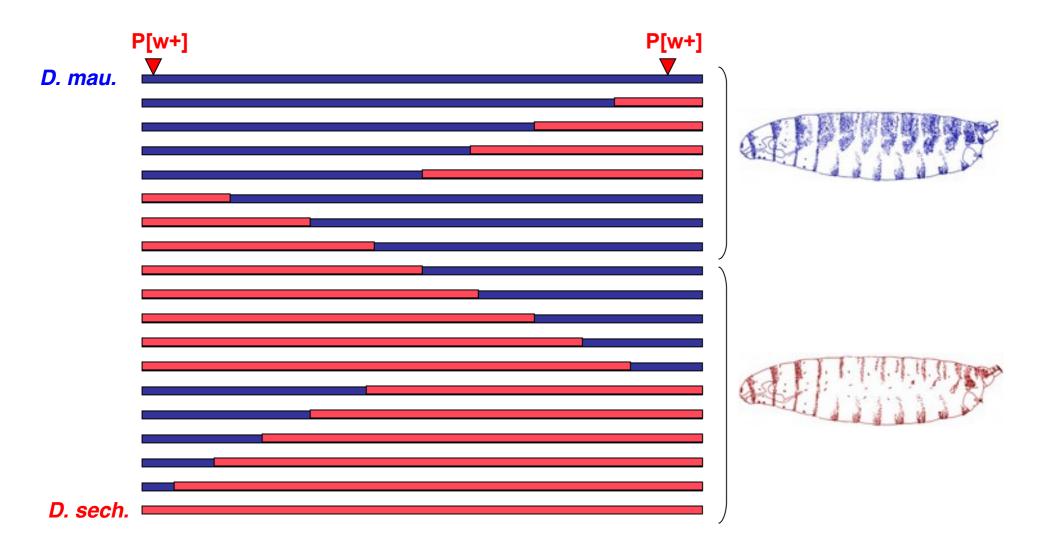
hybrids D. mel./D.sech.

3

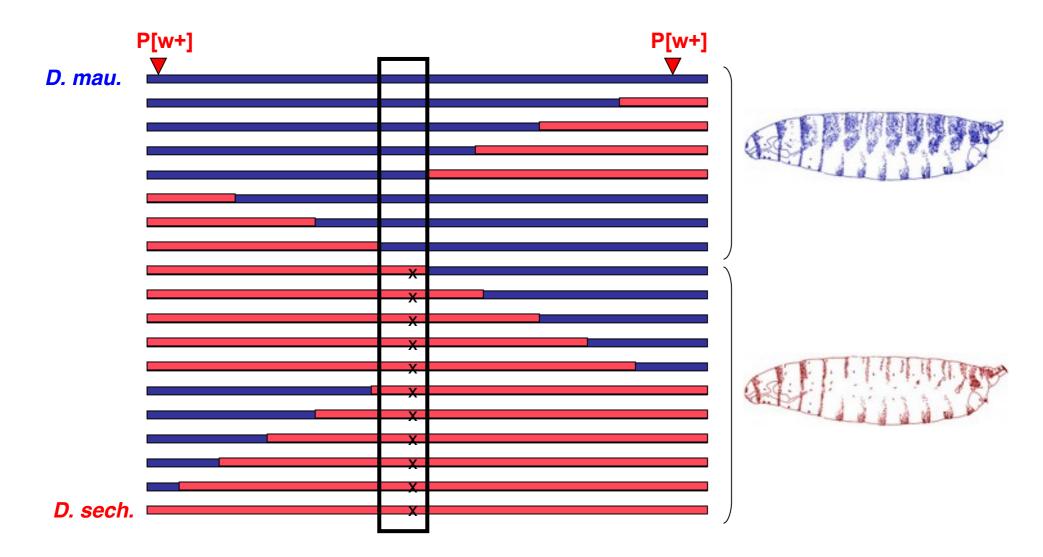


Sucena and Stern, 2000

Where is the mutation ?



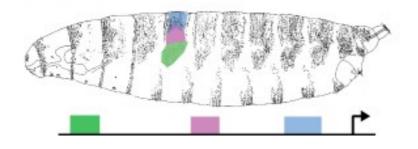
Where is the mutation ?

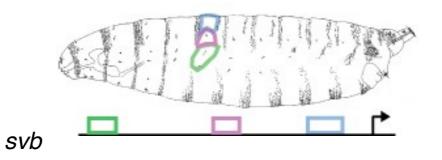


Several mutations in a single gene

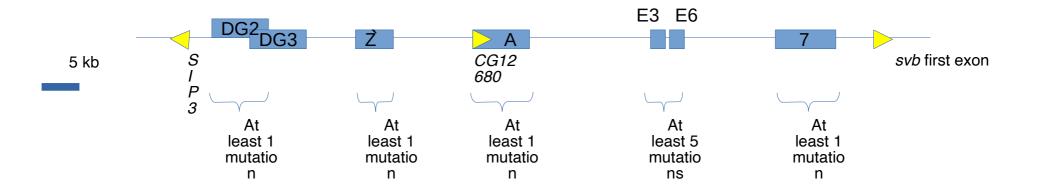
D. mauritiana

D. sechellia



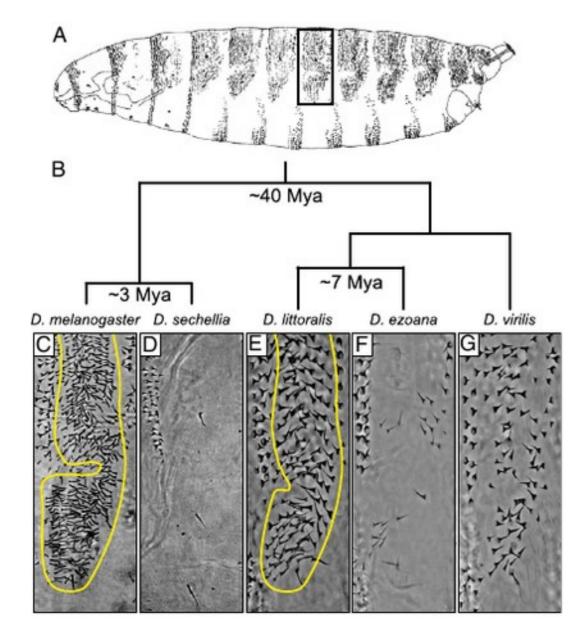






- McGregor, Orgogozo et al. 2007
 Nature
 - Frankel et al 2010 Nature
 - Frankel et al 2011 Nature

Convergent evolution in *D. ezoana*



Frankel, Wangl and Stern 2012
 PNAS

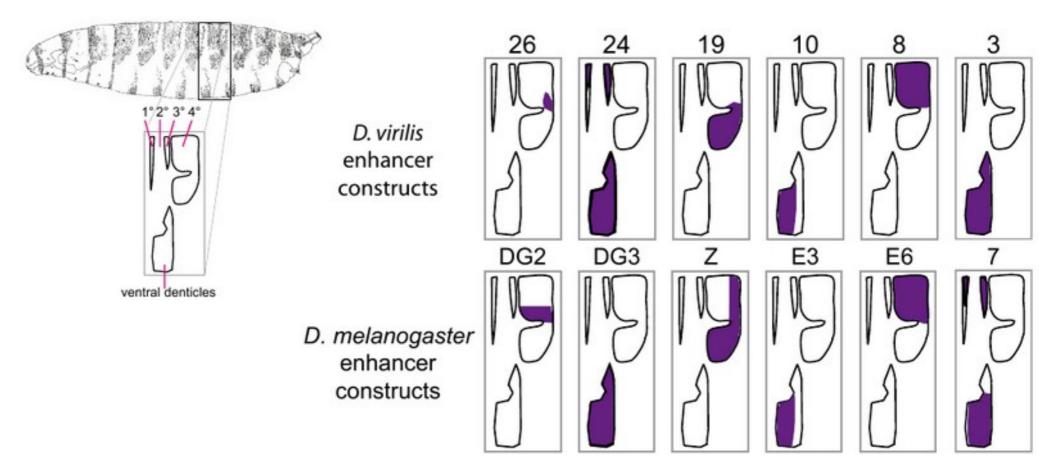
• D.



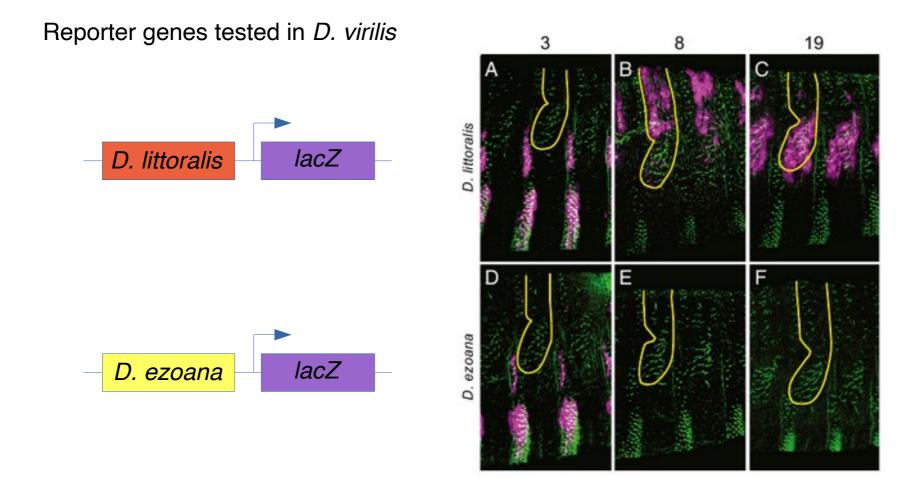
• D. melanogaster

Links = 30-bp-conserved sequences

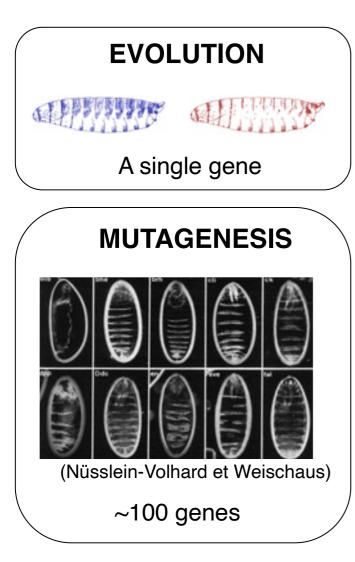
340

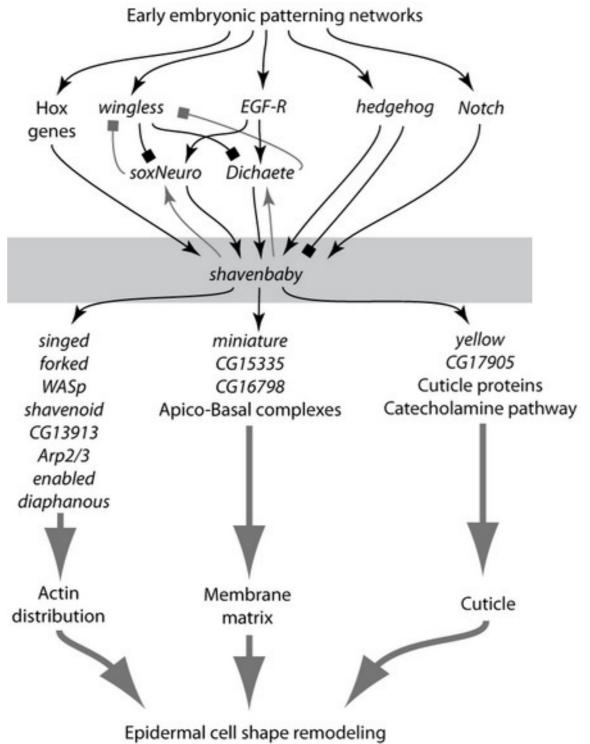


At least two cis-regulatory mutations



svb is a hotspot gene for evolution





(Stern et Orgogozo, 2008 Evolution – Stern et Orgogozo, 2009 Science)



BioAdhesio Expertise



The Organism and its Environment



The Organism and its Environment

Environment T, humidity, etc. chemicals food pathogens other species conspecifics

Ramona Bàdescu

omelo

découvre

Genes mediating interactions with E Immune defence Stress response Chemoreception Reproduction *Rapidly evolving*

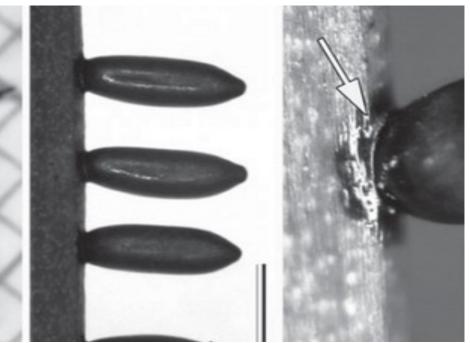
Bioadhesives



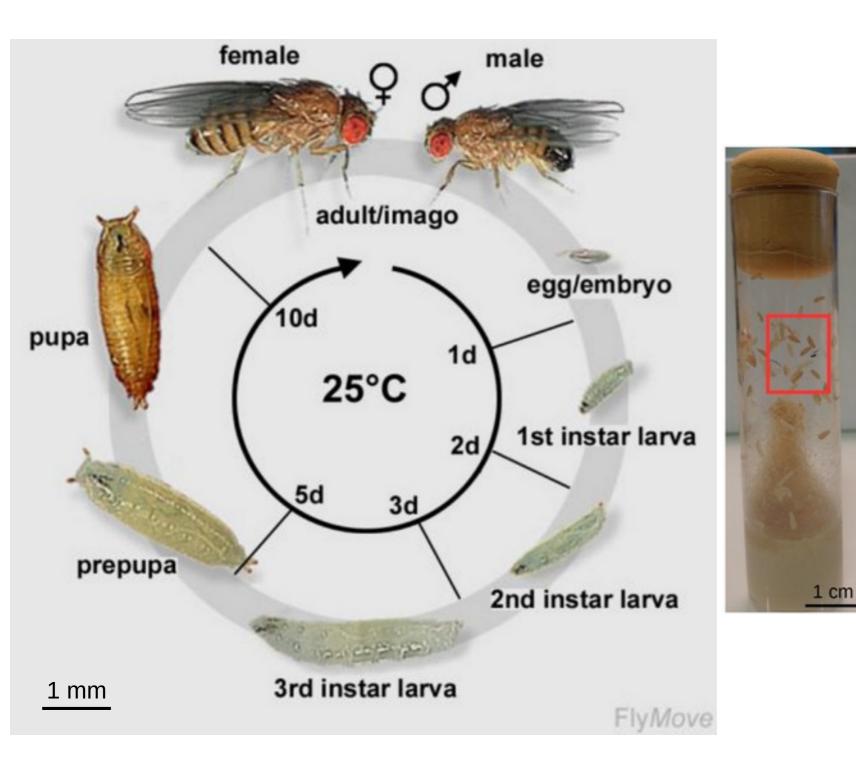
Mussle adhesion Burkett et al. 2009

In insects

- Locomotion
- Egg attachment
- Cocoon formation (silk)
- Pupa attachment



Asparagus beetle egg attachment Voigt and Gorb 2006





Drosophila glue

- A wide variety of environments
- Stick to a wide variety of substrates
- Difficult to observe pupae in nature

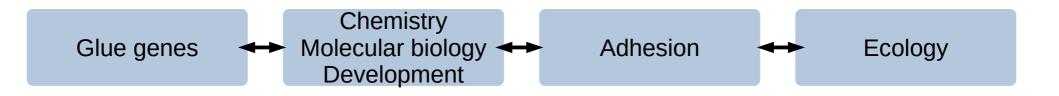


D. grimshawi

D. suzukii

- D. melanogaster D. repleta
- D. carcinophila

crab



Fly Glue Evolution

A Lalouette (genitalia) JN Lorenzi (SARS-CoV-2) R Vijendravarma (organ size)





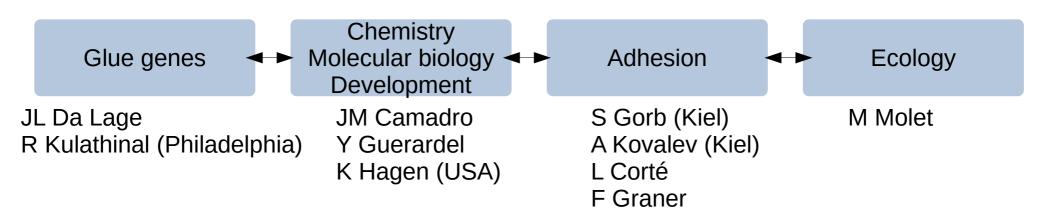


Flora Borne PhD (3 years)

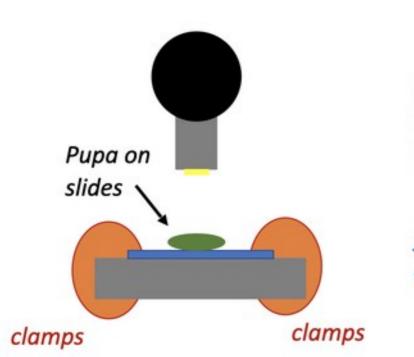
Manon Monier M2+PhD (3 years)

Isabelle Nuez Technician

Co-supervision: F. Graner & I

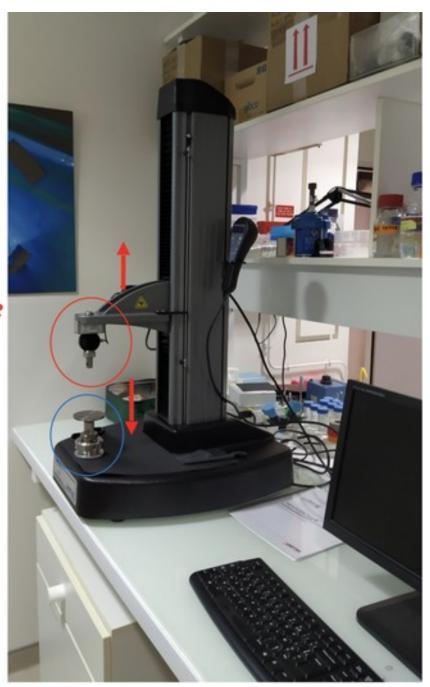


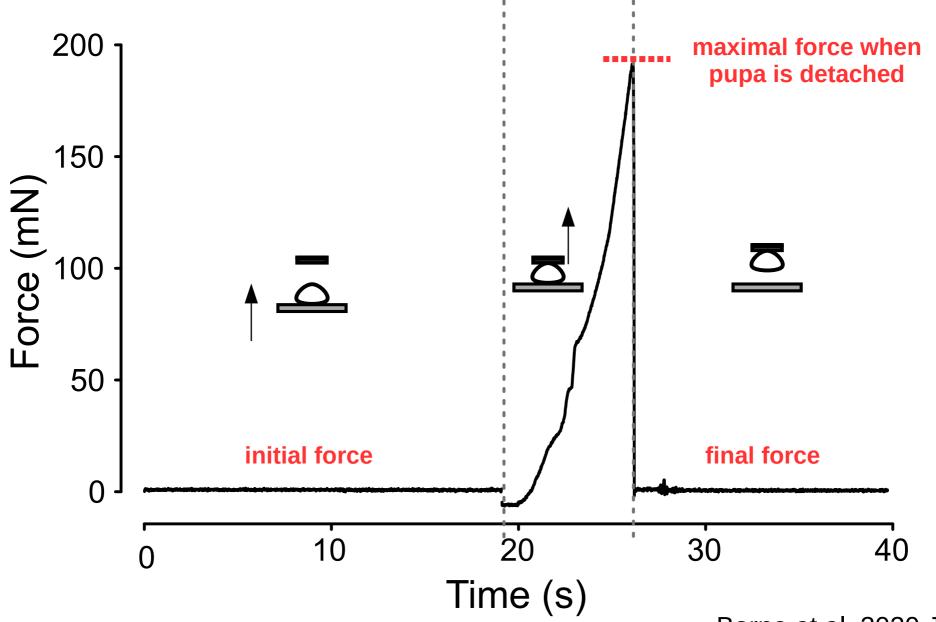
Experimental set up



Force sensor attached to the mobile part

Sample holder which is fixed





Borne et al. 2020 JEB

Drosophila glue is a universal glue

Same force on all tested substrates except Teflon

Force around 200 mN (like very adhesive tape) Strength around 200 kPa (~15 000 times the pupa weight)

moth eggs < Drosophila glue < mussel, barnacle < superglue</th>10-100 kPa100-300 kPa300-1000 kPa10 MPa

Borne et al. 2020 JEB

Function of fly glue Multiple hypotheses

Adhesion

Avoid predation (pupa hidden, cannot be removed) Resist to rain, wind, etc.

Help adult emerge from the pupal case

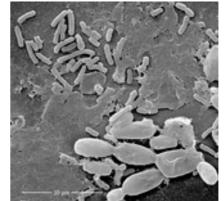
Repellent

Avoid predation (pupa are not eaten)

Preservative

Avoid fungi or bacteria infections on the pupal case

Beňová-Liszeková et al. 2019



Function of fly glue Multiple hypotheses

Adhesion

Avoid predation (pupa hidden, cannot be removed Resist to rain, wind, etc. Help adult emerge from the pupal case

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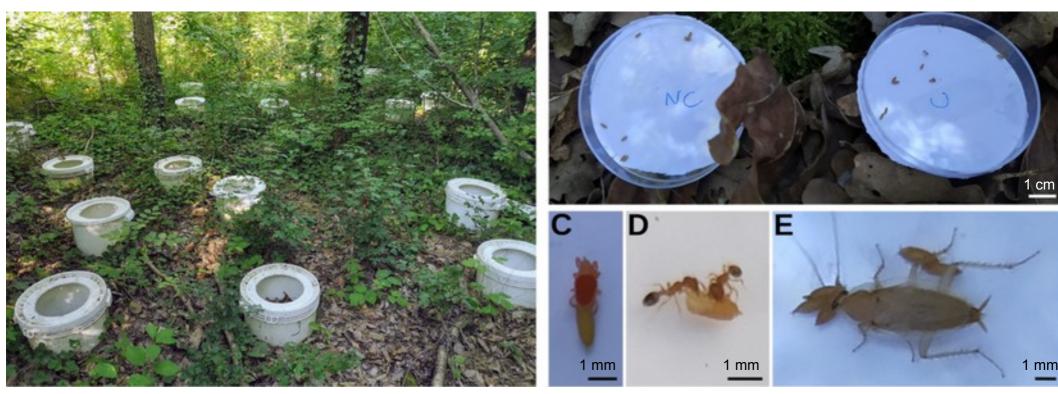
Preservative

Avoid fungi or bacteria infections on the pupal case

Beňová-Liszeková et al. 2019



Test in Bois de Vincennes

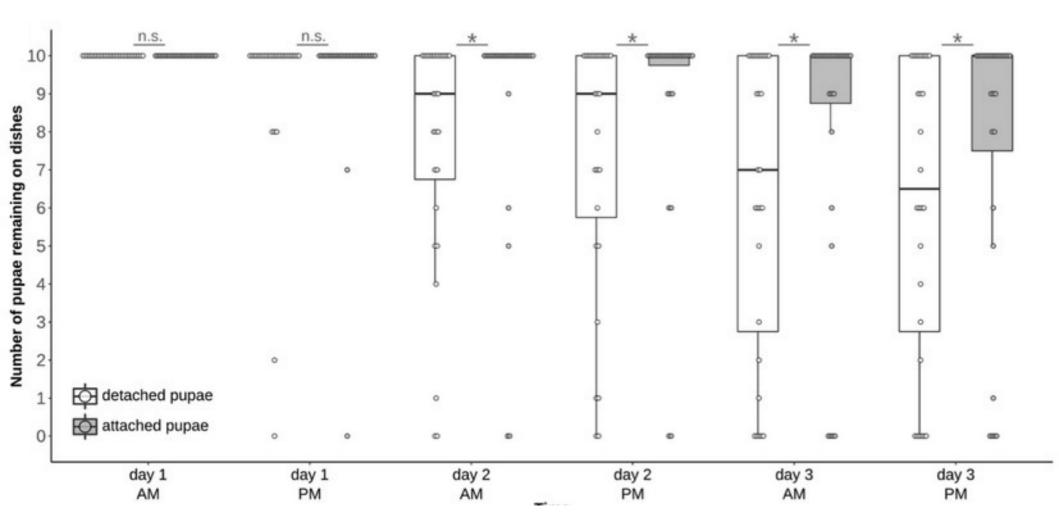


detached

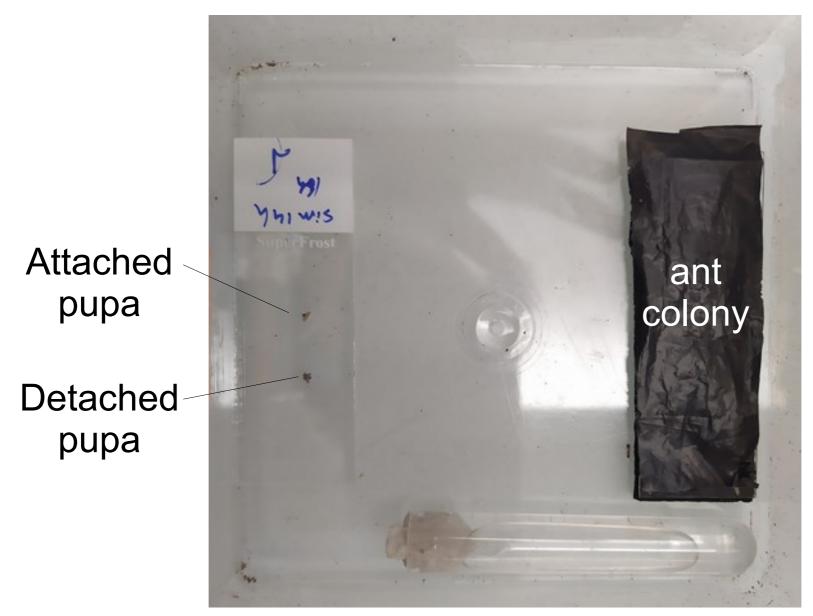
attached

In each bucket (n=28): one Petri dish w/ 10 detached pupae one Petri dish w/ 10 attached pupae Checked twice a day

Attached pupae are taken away less frequently than detached pupae

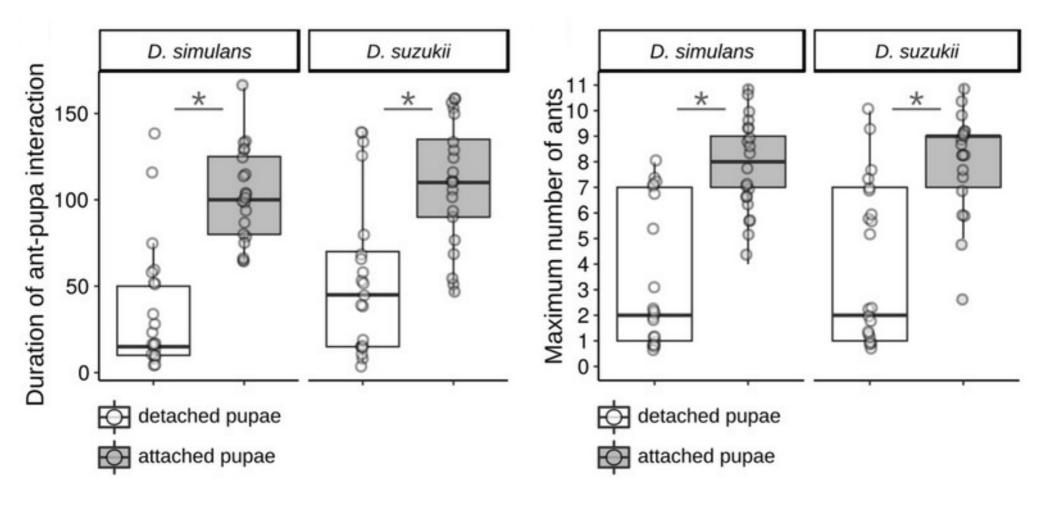


Test in the lab with Temnothorax ants



7 ant colonies, one test per day, 6 days either *D. simulans* or *D. suzukii*

Attached pupae require more time and more ants to go away



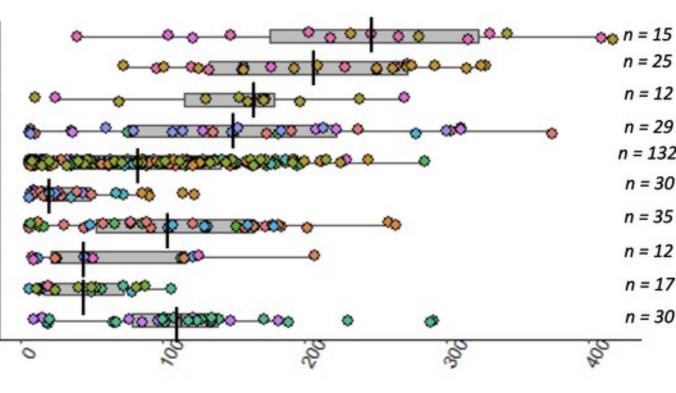
Ecology of fly glue: what is the function of the glue?

- Glue protects from predation
- Detached pupae are brought to the nest while attached pupae are eaten on site and require more ants and more time

Borne, Prigent, Molet and Courtier, Proc B, 2021

Weaker adhesion in *D. suzukii* and *D. biarmipes*

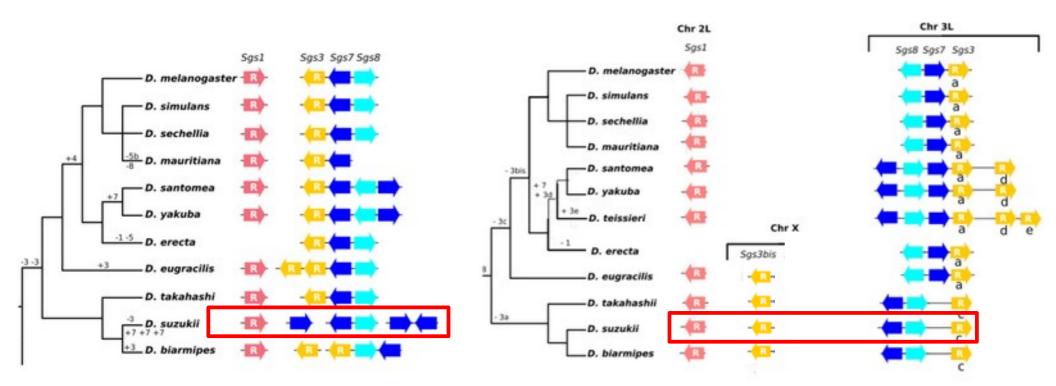
Drosophila melanogaster Drosophila simulans Drosophila mauritiana Drosophila eugracilis Drosophila suzukii Drosophila biarmipes Drosophila takahashii Drosophila prostipennis Drosophila rhopaloa Drosophila ananassae



Weaker adhesion in *D. suzukii* and *D. biarmipes*

Due to loss of Sgs3 in *D. suzukii* ?

No.

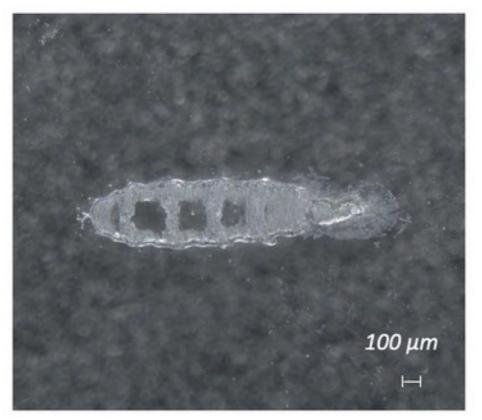


Da Lage 2019 BMC Evol Biol

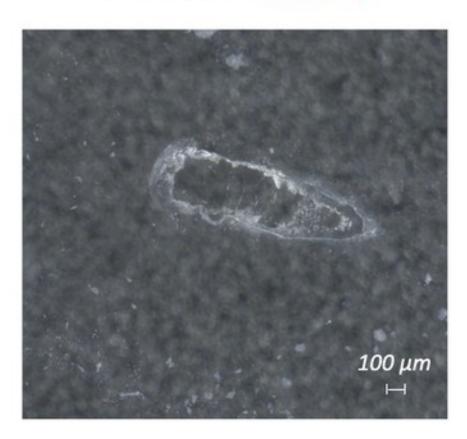
Our new study, unpublished

Little glue produced?

Drosophila simulans



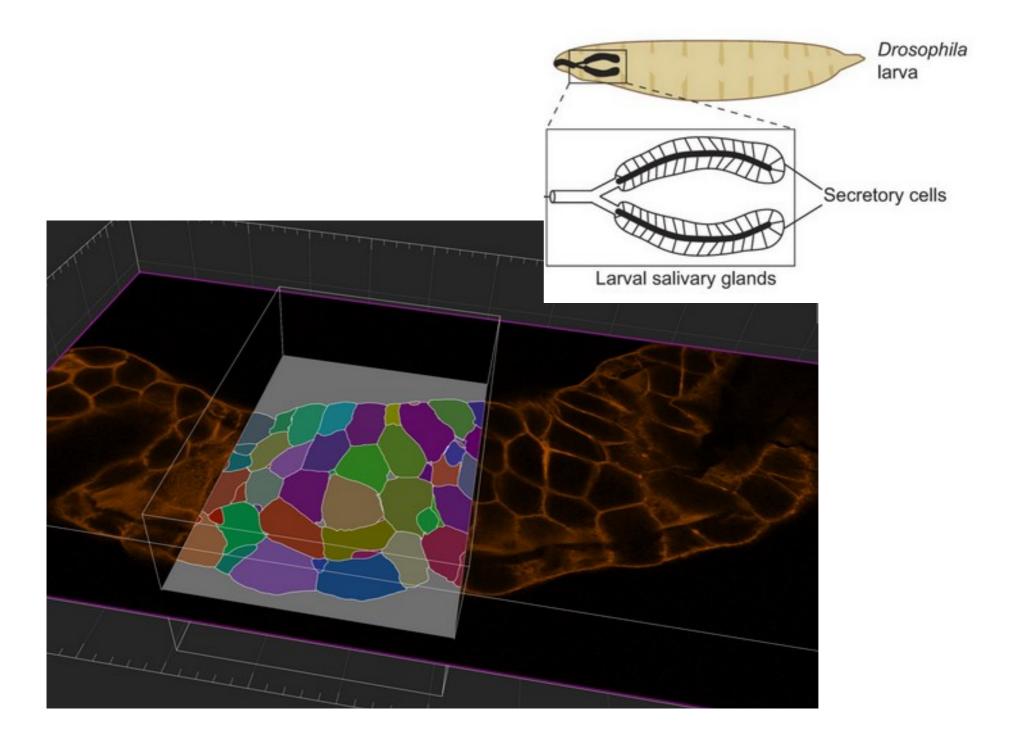
Drosophila suzukii

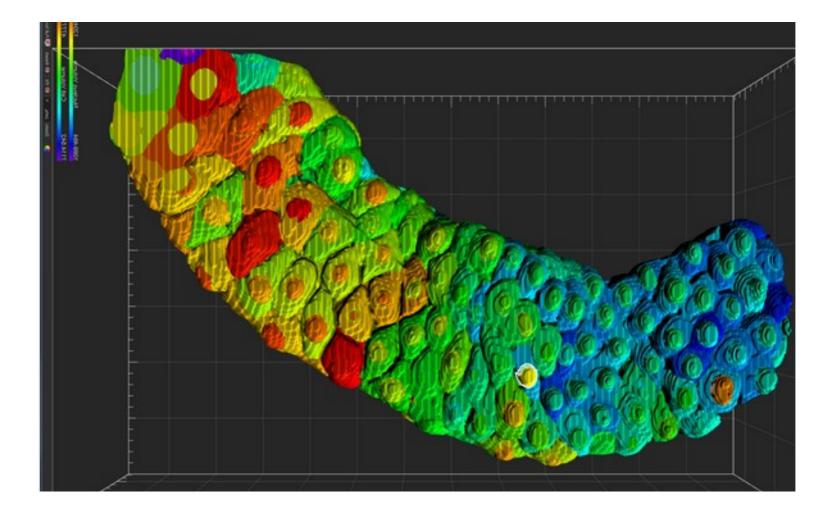




Ongoing: measure of glue prints and salivary gland size







The genotype-phenotype relationship

The distinction between genotype and phenotype is the basis of genetics

"The view of natural inheritance as realized by an act of transmission, viz., the transmission of the parent's (or ancestor's) personal qualities to the progeny, is the most naive and oldest conception of heredity."

"All "types" of organisms, distinguishable by direct inspection or only by finer methods of measuring or description, may be characterized as "**phenotypes**."

" A "**genotype**" is the sum of all the "genes" in a gamete or in a zygote."



Johansen 1911

Phenotype = observable attributes of an individual

Genotype = inheritable genetic material = DNA or RNA

How do genotypes map onto phenotypes ?

A very brief history of genetics

Transmission + *Phenotypic* expression

- Mendel 1860s Controlled breeding, use of probabilities Dominant/recessive/intermediate expression of traits Random transmission of "factors"

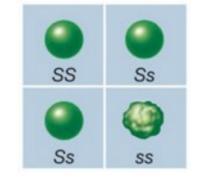
- Cytology from 1880-90s Flemming, Boveri, Sutton chromosomes as support of heredity, meiosis

- Classical genetics 1900-1950 distinction genotype-phenotype segregation of characters & genes genetic map, sex chromosomes penetrance - expressivity

"Rediscovery" of Mendel's law Extension to animals





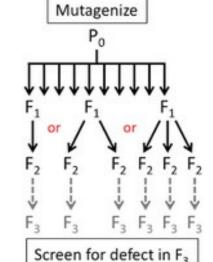


- Foundation of molecular biology 1940-1970

in part using bacteria and phage genetics DNA as the material support of heredity

- Deciphering the role of the genes: 1970-Cellular, developmental, behavioral genetics: Screens "High-throughput" versions 2000- (e.g. deletion libraries) **Reverse** genetics

- Association mapping on natural populations: 2000-





Divergences and syntheses

	Mendelian genetics "Mendelians"		Statistical genetics "Biometricians"		
1900					agriculture
			"Neodarwinian synthesis"		
1930	<u>Laboratory</u> genetics		Population genetics evolution		Quantitative genetics
	genetice		genotype		phenotype
1950	Molecular biolog	y			
1975	Molecular gene cell biology, development physiology, etc.	etics	Molecular evolution	Molecular markers	
1990	Functional genomics	<u>Evo</u> - Devo	Evolutionary genomics	/	Quantitative genetics molecular basis

How do genotypes map onto phenotypes ?

DEVELOPMENTAL BIOLOGY

EVO-DEVO

Both are direct descendants of Morgan's school. Emphasis on genes.

How does an organism form from a single cell?

What makes one organism different from another one?

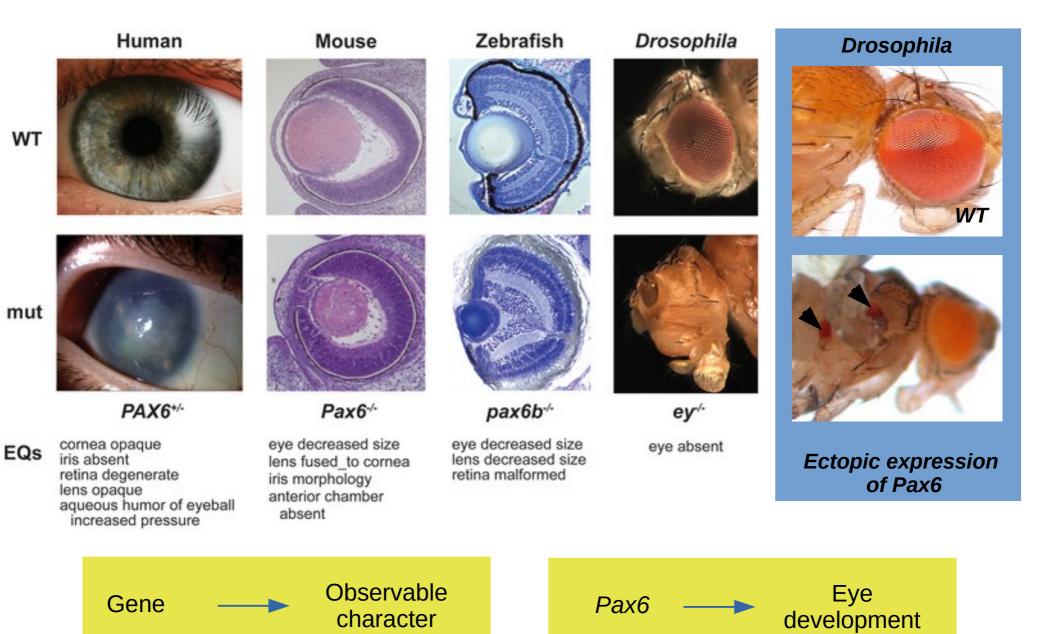
One of the central problems of biology is that of differentiation - how does an egg develop into a complex many-celled organism? That is, of course, the traditional major problem of embryology; but it also appears in genetics in the form of the question, **"How do genes produce their effects?**

Sturtevant, 1932

How do genes produce observable traits?

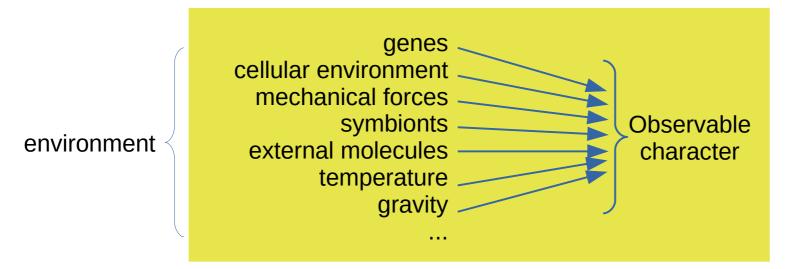


Pax6 : an eye gene ?



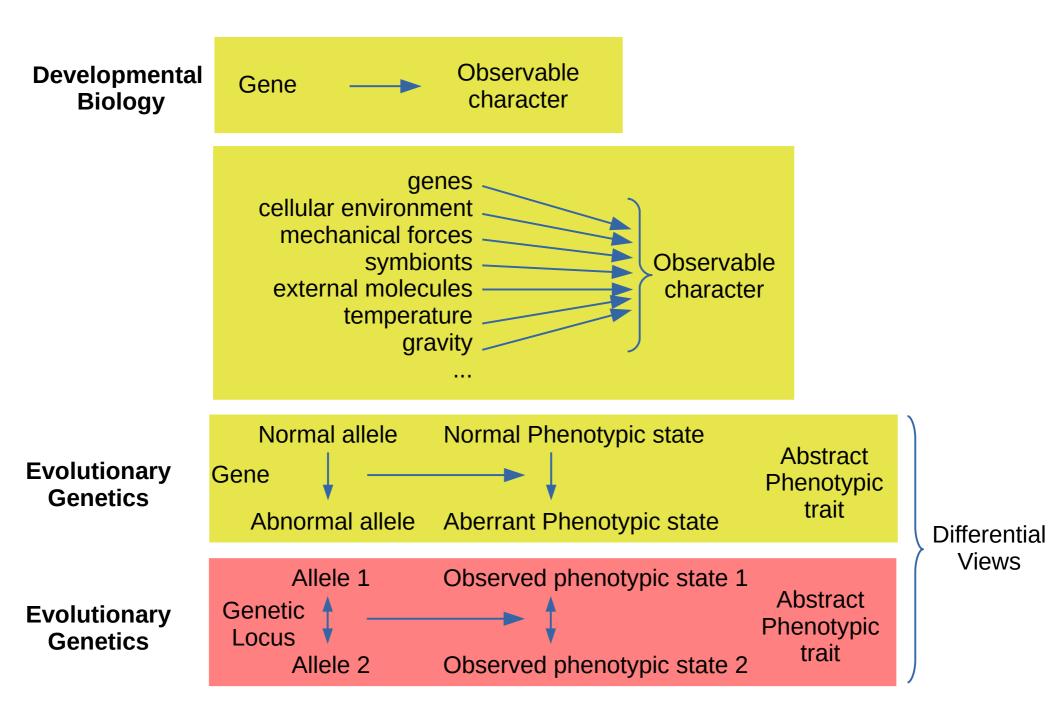






Better, but difficult to disentangle the effects

A Cause/Mechanism relates to a Difference



Different kinds of phenotypes

Morphology

Color

Size and shape

Presence/



absence Aristote, Historia animalium, book I, 2, 300BC Position Physiology

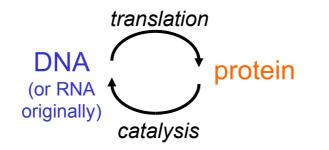
Behavior

Genotype & Phenotype = what engenders = what is apparent

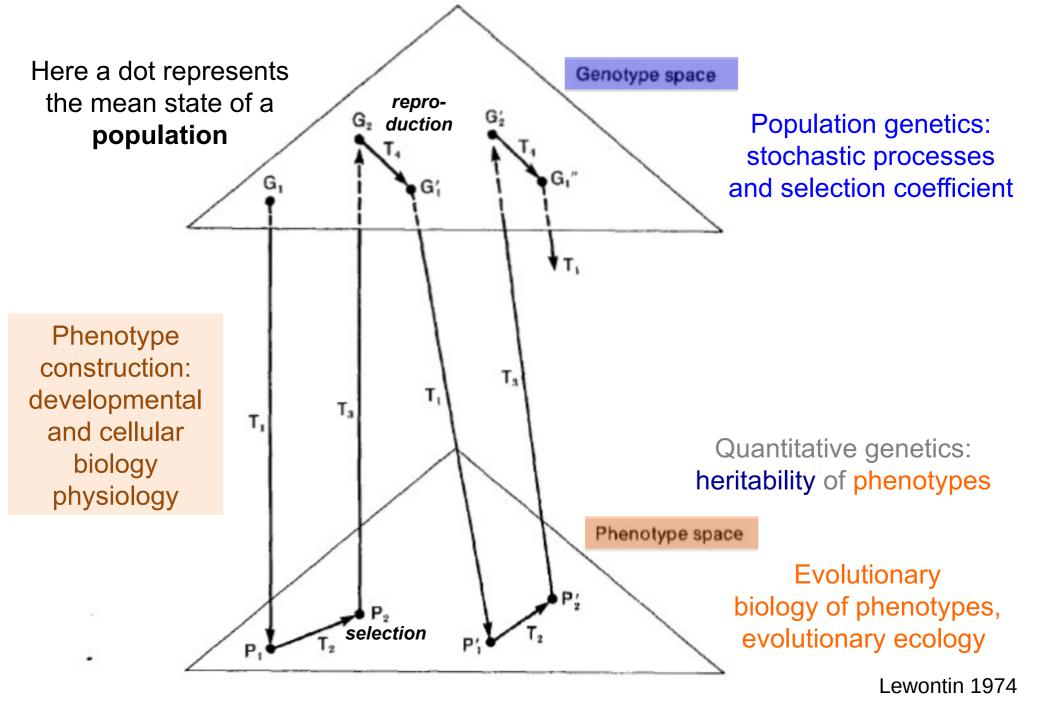
DNA/RNA • Regulation of gene expression

- Biochemical reactions
- Subcellular architecture
- Assembly of cells
- Organism morphology and behavior

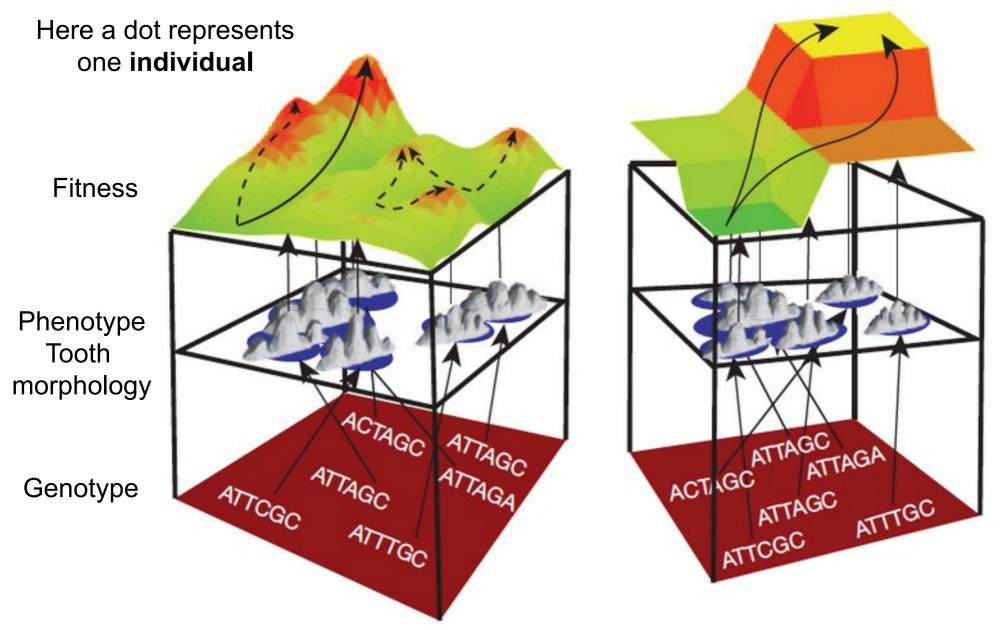
distinction appeared at the origin of life: etc.



The first genotype-phenotype map

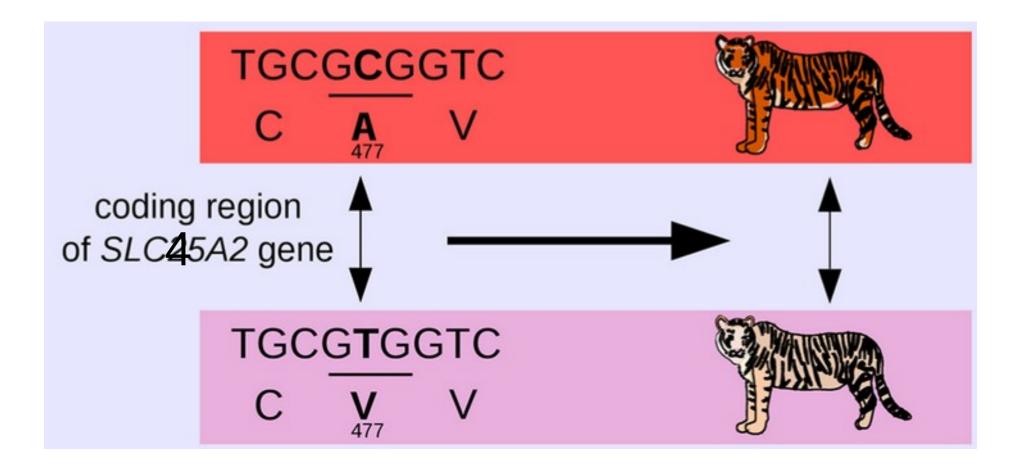


The genotype-phenotype-fitness map



Salazar-Cuidad & Martin-Riera 2013

The genotype-phenotype connection



Xu et al 2013 Current Biology Orgogozo et al 2015 Frontiers Genetics Genotype = "the genetic makeup of an organism that determines a specific phenotype (trait), from one generation to the next, and potentially throughout the population".

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





The genotype-phenotype connection is about differences

The genes contributing to evolution

www.Gephebase.org >2000 entries



Suggest an Article





The Database of Genotype-Phenotype Relationships Search Gephebase for genes, phenotypes, taxa, mutations, and

Gephebase compiles genotype-phenotype relationships, i.e. associations between a mutation and a phenotypic variation. Gephebase consolidates data from the scientific literature about the genes and the mutations responsible for phenotypic variation in Eukaryotes (mostly animals, yeasts and plants). We plan to include non Eukaryote species in the future. For now, genes responsible for human disease and for aberrant mutant phenotypes in laboratory model organisms are excluded and can be found in other databases (OMIM, OMIA, FlyBase, etc.). QTL mapping studies that did not identify single genes are not included in Gephebase.

Q

If you use Gephebase for your publication, please cite: Martin, A., & Orgogozo, V. (2013). The loci of repeated evolution: a catalog of genetic hotspots of phenotypic variation. Evolution, 67(5), 1235-1250.

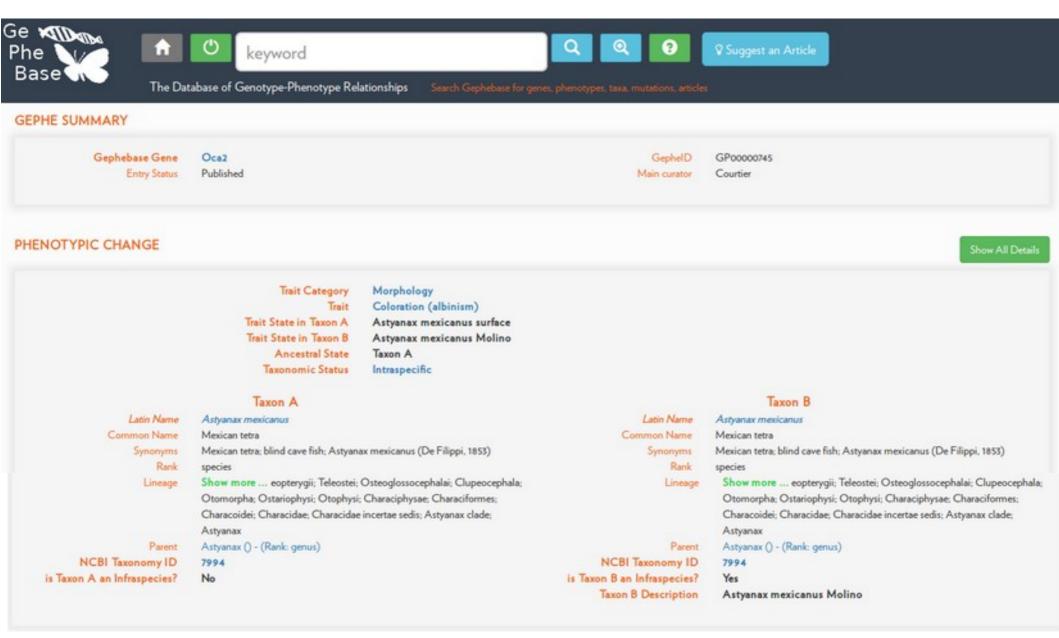
Conference on Gephebase and the loci of evolution (Paris, 2016)

You can retrieve data via HTTP requests through APIs. Below is the list of available APIs. By default, the response data is sent in xml format. For each field, it is possible to only enter a subset of a keyword and still be able to successfully retrieve the desired data. (example: "Bir" for "Birds" will display all data that have the characters "Bir").

Orgogozo et al Nucleic Acid Research 2019

oca2 in cavefish





Trait Category Trait Trait State in Taxon A Trait State in Taxon B Ancestral State Taxonomic Status

Morphology Coloration (albinism) Astyanax mexicanus surface Astyanax mexicanus Molino Taxon A Intraspecific

Taxon A

Latin Name	Astyanax mexicanus		
Common Name	Mexican tetra		
Synonyms	Mexican tetra; blind cave fish; Astyanax mexicanus (De Filippi, 1853)		
Rank	species		
Lineage	Show more eopterygii; Teleostei; Osteoglossocephalai; Clupeocephala;		
	Otomorpha: Ostariophysi: Otophysi: Characiphysae: Characiformes:		
	Characoidei; Characidae; Characidae incertae sedis; Astyanax clade;		
	Astyanax		
Parent	Astyanax () - (Rank: genus)		
BI Taxonomy ID	7994		
an Infraspecies?	No	is Tax	

Taxon B

Astyanax mexicanus Mexican tetra Mexican tetra; blind cave fish; Astyanax mexicanus (De Filippi, 1853) species Show more ... eopterygii; Teleostei; Osteoglossocephalai; Clupeocephala; Otomorpha; Ostariophysi; Otophysi; Characiphysae; Characiformes; Characoidei; Characidae; Characidae incertae sedis; Astyanax clade; Astyanax Astyanax () - (Rank: genus) 7994 Yes

xon B an Infraspecies? Taxon B Description

NCBI Taxonomy ID

Latin Name

Synonyms Rank

Lineage

Parent

Common Name

Astyanax mexicanus Molino

- NCBI Tax
- is Taxon A an Inf

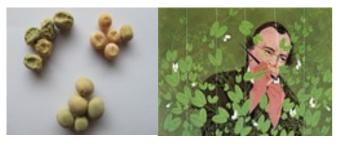
GENOTYPIC CHANGE

Generic Gene Name	Oca2	UniProtKB	Q62052		
Synonyms	p: D7Nic1; p <cas>; D7H15S12; D7Icr28RN; P</cas>	Mus musculus			
String	10090.ENSMUSP00000032633	GenebankID or UniProtKB	ABB29299		
Sequence Similarities	Belongs to the CitM (TC 2.A.11) transporter family.				
GO - Molecular Function	-				
GO - Biological Process	GO:0055085 : transmembrane transport show more				
GO - Cellular Component	GO:0016021 : integral component of membrane show more				
	Presumptive Null	Yes			
	Molecular Type	Coding			
	Aberration Type	Deletion			
	Deletion Size	-			
	Molecular Details of the Mutation	Deletion of exon 21			
	Experimental Evidence	Linkage Mapping			
	Main Reference	Genetic analysis of cavefish reveals molecular conv	ergence in the evolution of albinism. (2006)		
	Authors	Protas ME; Hersey C; Kochanek D; Zhou Y; Wilker	ns H: Jeffery WR: Zon LI: Borowsky R: et al show more		
	Abstract	The genetic basis of vertebrate morphological evolution has traditionally been very difficult to examine in naturally occurring			
		populations. Here we describe the generation of a g	enome-wide linkage map to allow guantitative trait analysis of evolutionarily		
			pecies that has, in a series of independent caves, repeatedly evolved specialized		

characteristics adapted to a unique and well-studied ecological environment. We focused on the trait of albinism and discovered

		Additional References	, ,	· · · · · · · · · · · · · · · · · · ·	
RELATED GEPHE					
	Related Genes 1 (MC1R)			Related Haplotypes 1	
COMMENTS					
YOUR FEEDBACK is welcom	el				
Feedback			Your E-mail	Optional and remains confidential (not displayed	online). Only used to contact you if

Wrinkled seed: TE insertion (Bhattacharyya 1990)



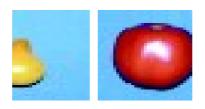
myostatin coding region (Grobet 1997)



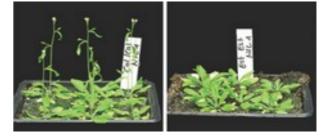
Mc1r coding region (Eizirik 2003)



OVATE coding region (Liu 2002)



FRI coding region (Johanson 2000)



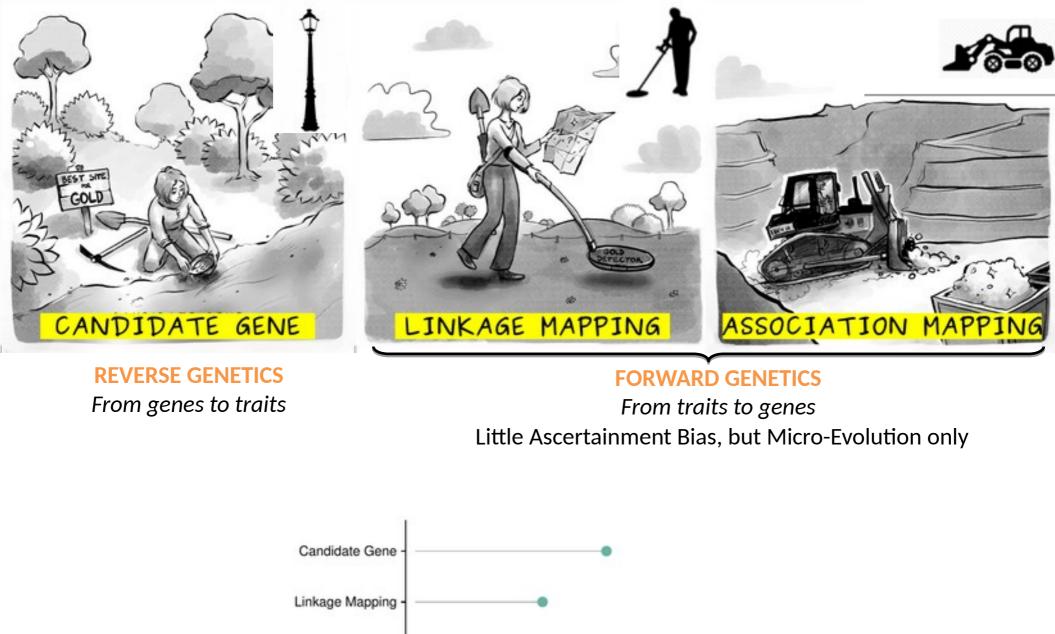
luciferase coding region (Stolz 2003)

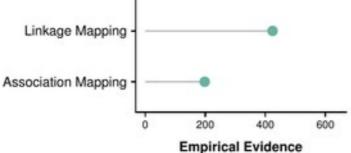


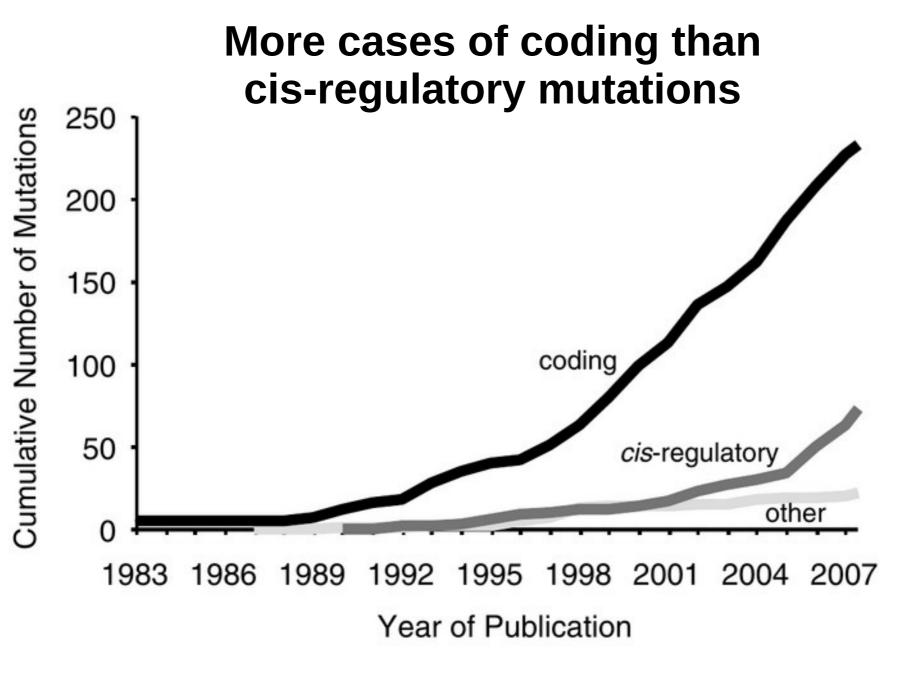
anthocyanin-2 coding region (Quattrocchio 1999)



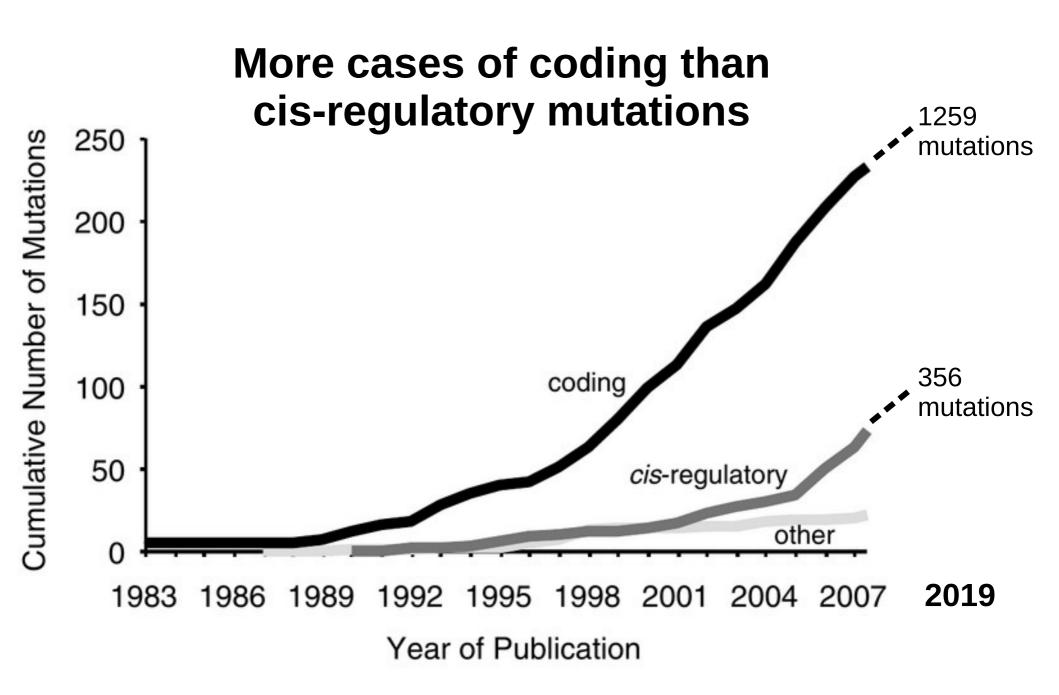
THREE APPROACHES to FIND the GOLDEN LOCI of EVOLUTION







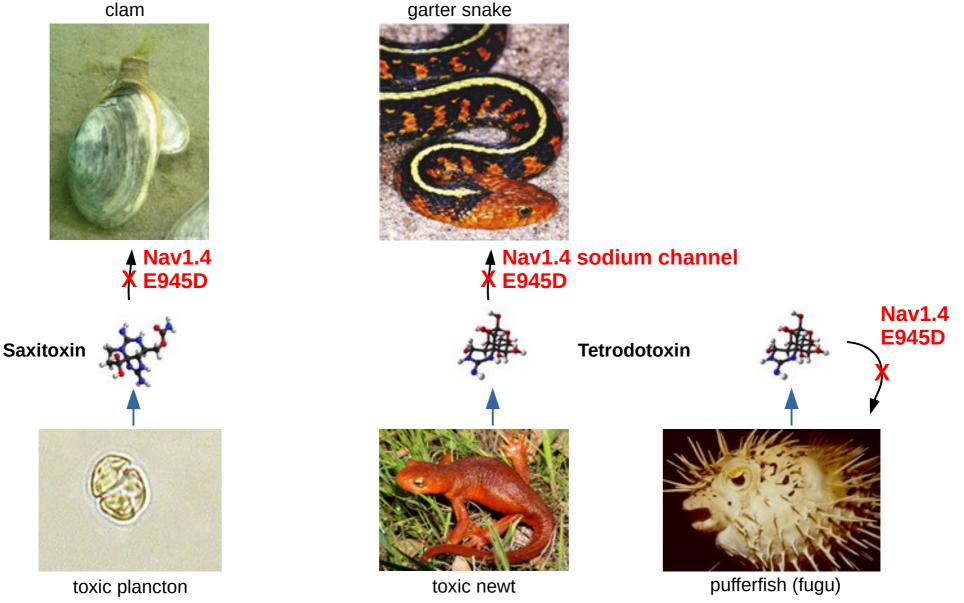
survey of ~300 articles Stern and Orgogozo 2008 Evolution



Evolution repeats itself

Repeated evolution via the same amino acid change

clam



Bricelj 2005 Geffeney 2005 Venkatesh 2005

Repeats in..



.. the genes responsible for natural evolution

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

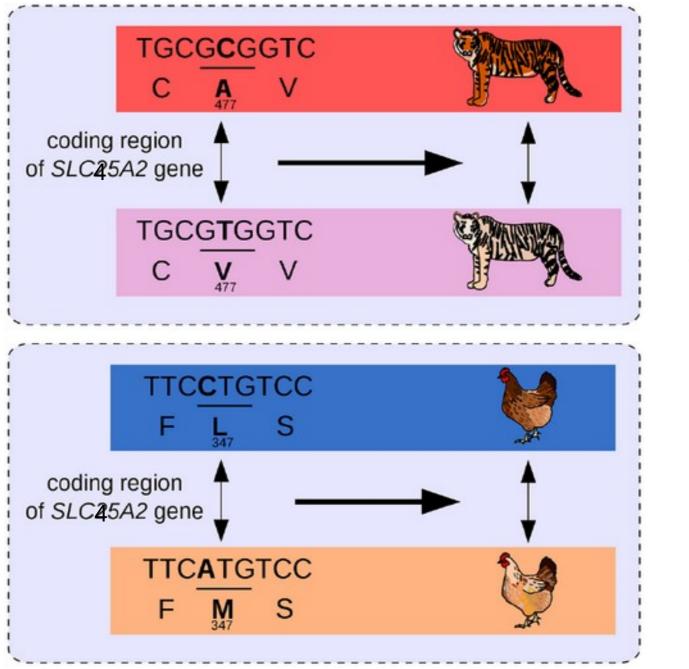


.. the genes responsible for experimental evolution

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

Orgogozo 2014 Interface Focus

Repeated evolution



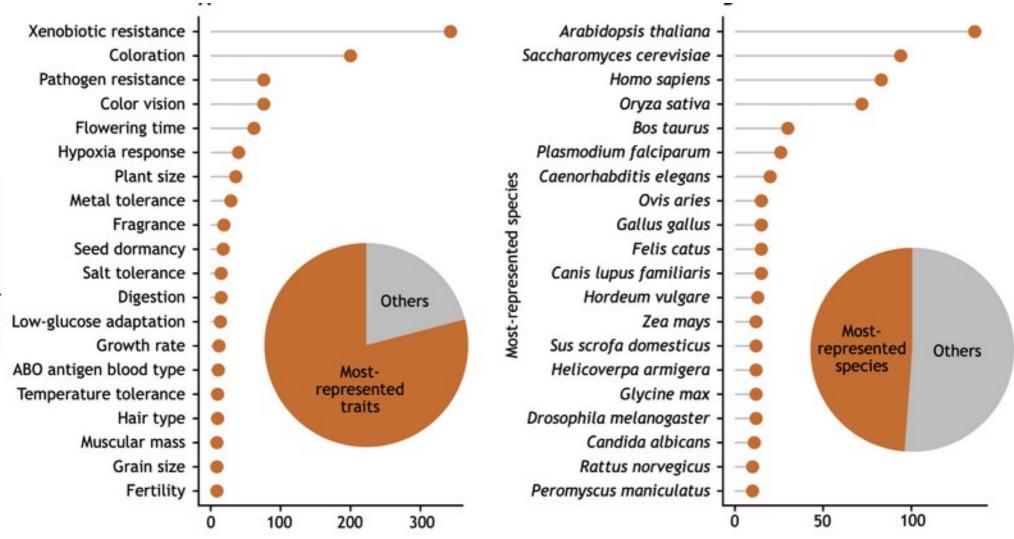
Also in: Humans Horses Quails Chickens Mice Pigeons

Orgogozo et al 2015 Frontiers Genetics

TOP HOTSPOT GENES 2022

MC1R (93) para (kdr) (54) Kit (type III receptor protein-tyrosine kinase) (46) Na/K-ATPase alpha-subunit (45) beta-tubulin (33) beta-tubulin (ben-1) (27) SCN4A (Nav1.4) (26) Agouti (ASIP) (24) tyrosinase (TYR) (24) tyrosinase-related protein 1 (TYRP1) (24) AtGA20ox1 (=GA5=Sd1) (24) Agouti (24) alcohol dehydrogenase (Adh) (23) opsin - (SWS1) (20)

Bias towards certain traits and species



Most-represented traits

Specialized loci in the genome

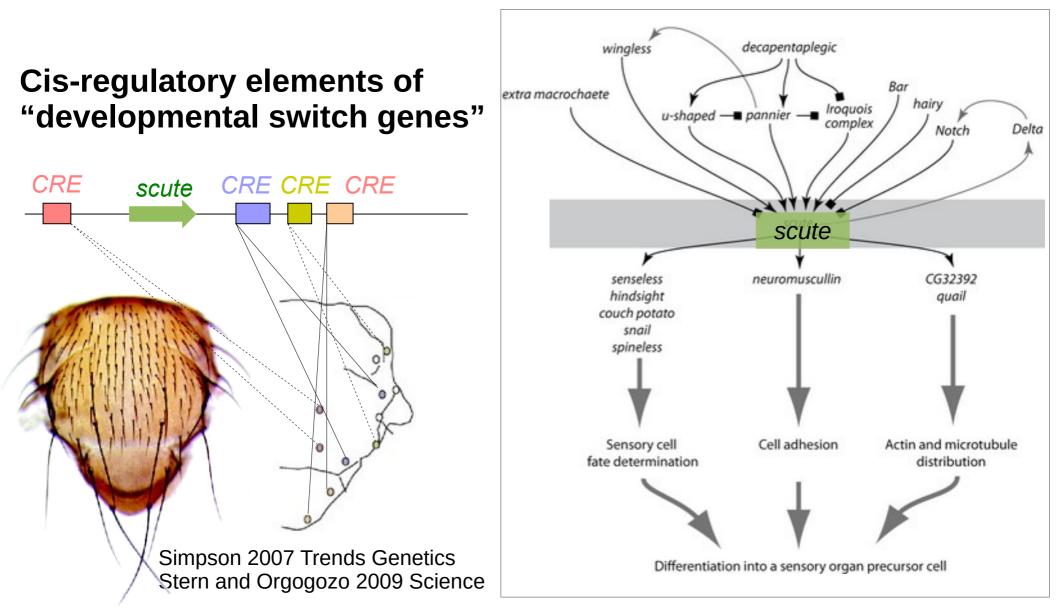
Proteins that interact with external molecules

oxygen, photons, insecticide, cholesterol...

Specialized loci in the genome

Proteins that interact with external molecules

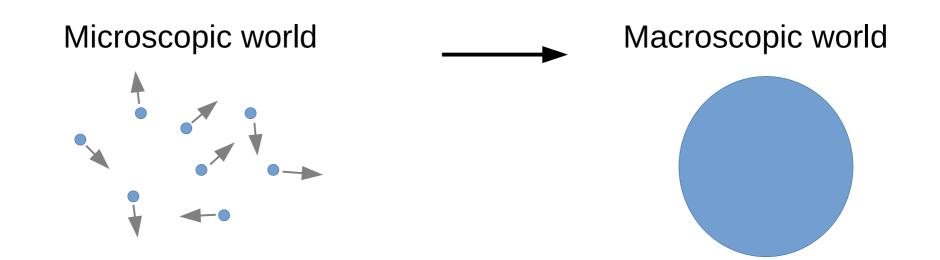
oxygen, photons, insecticide, cholesterol...



From random processes can emerge predictability

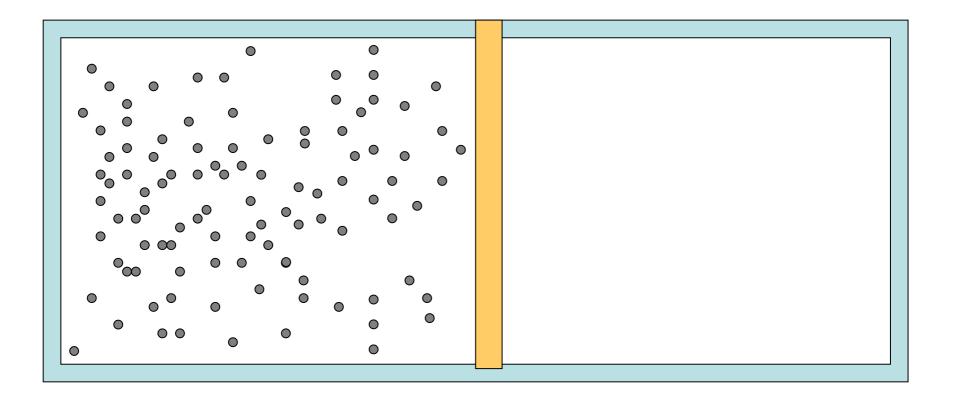
Many unpredictable processes at a low level Mutations in DNA Chromosome segregation during meiosis Assortative mating Gamete competition during fecondation Life history traits Genetic linkage Environmental changes (meteorite, etc.)

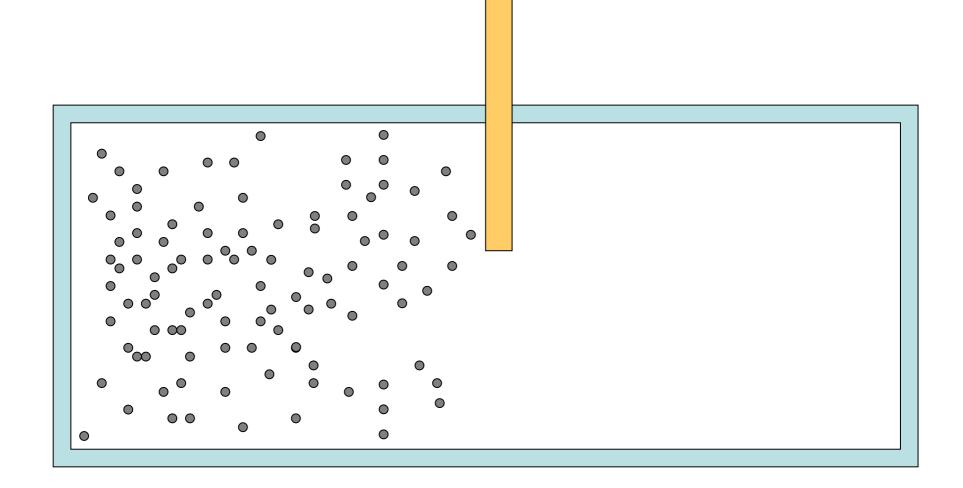
From random processes can emerge predictability

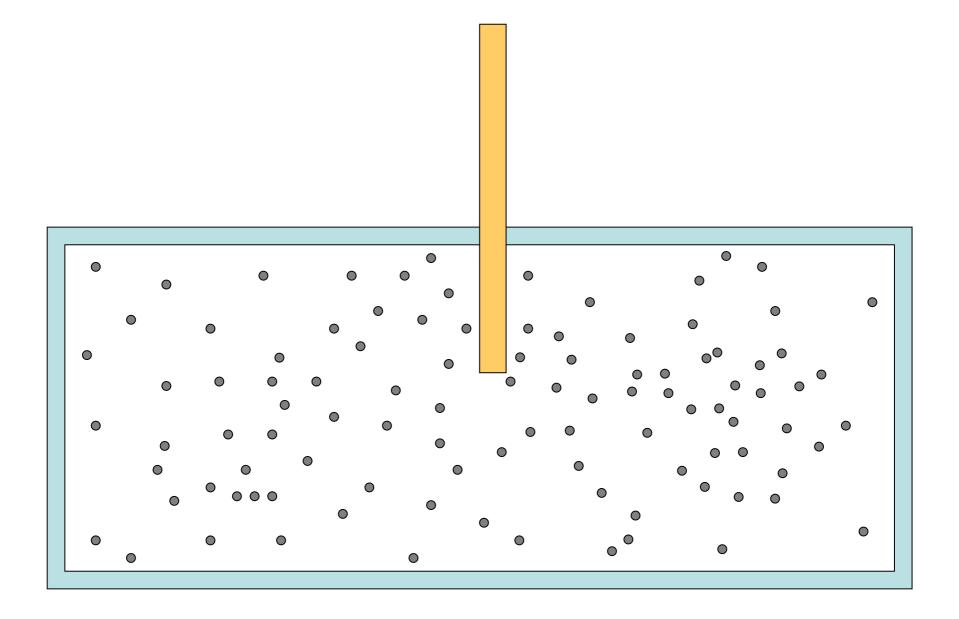


Position, mass, velocity of each particle

Pressure, Volume, Temperature, Number of moles

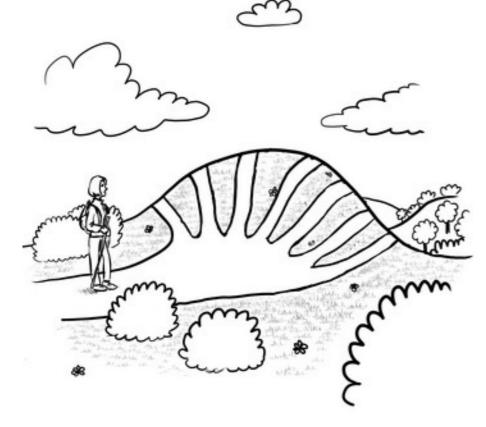


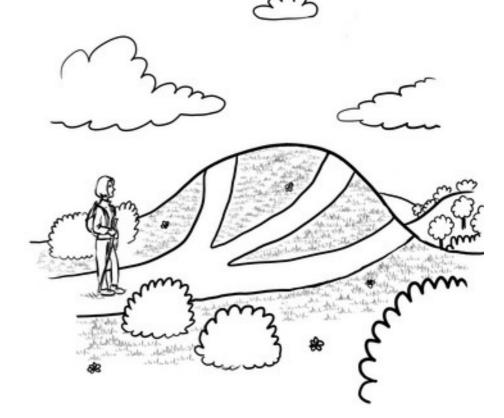




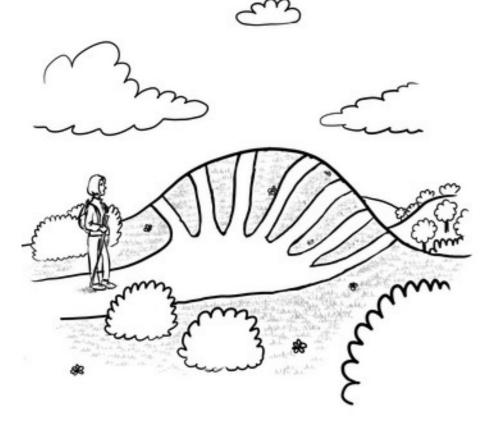
After a few seconds

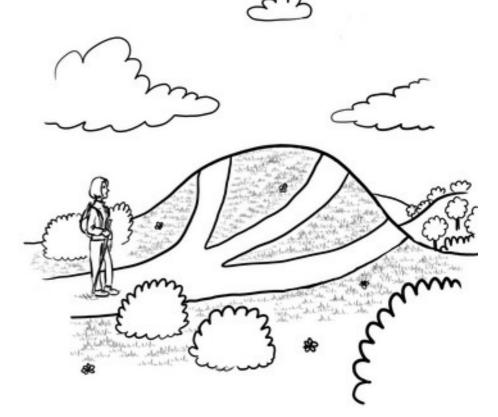
A small number of genetic solutions for a given phenotypic change





A small number of genetic solutions for a given phenotypic change





We sometimes seem to have forgotten that the original question in genetics was not what makes a protein but rather 'what makes a dog a dog, a man a man'. (D. Noble – The Music of Life)

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.

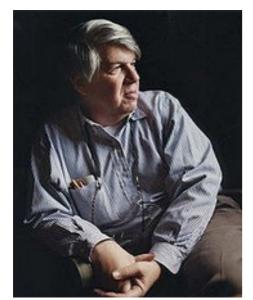
It is hard to realize that the living world as we know it is just one among many possibilities; that its actual structure results from the history of the earth.

1977

Evolution and Tinkering

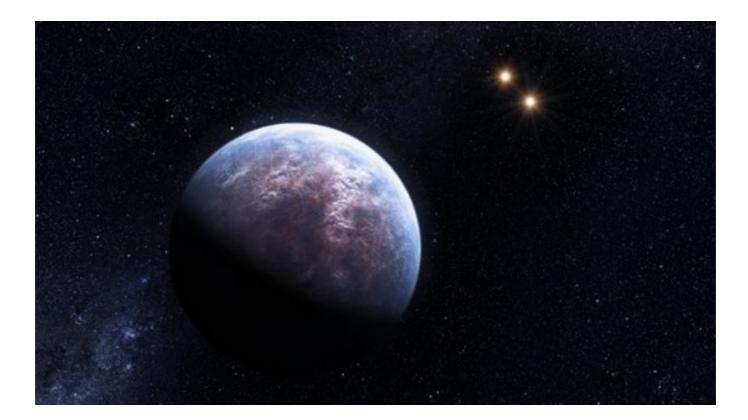
François Jacob

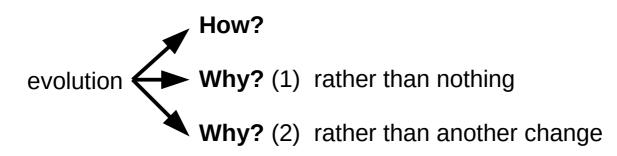
Stephen Jay Gould, 2002





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





Lab

Olga Nagy (past member) Isabelle Nuez (IE)

Michael Lang (CR) Alexis Lalouette (MCU)

Bénédicte Lefèvre (PhD) Flora Borne (PhD) Manon Monier (PhD)

Main collaborators

Fly anatomy Jean David (Gif-sur-Yvette)

> *Morphometrics* Vincent Debat (Paris) François Graner (Paris)

Genotyping David L. Stern (Janelia Farm, USA)

> Fly crosses Daniel Matute (UNC, USA)

Gephebase Arnaud Martin (Berkeley, USA)









The differential view of genotype-phenotype relationships



Virginie Orgogozo1*, Baptiste Morizot2 and Arnaud Martin3

The "Mendelian Gene" and the "Molecular Gene": Two Relevant Concepts of Genetic Units

V. Orgogozo*1, A.E. Peluffo*, B. Morizot*

*Institut Jacques Monod, UMR 7592, CNRS-Université Paris Diderot, Sorbonne Paris Cité, Paris, France ¹Université Aix-Manseille, CNRS UMR 7304, Aix-en-Provence, France ¹Corresponding author: e-mail address: virginie.orgogoro@normalesup.org



THE LOCI OF REPEATED EVOLUTION: A CATALOG OF GENETIC HOTSPOTS OF PHENOTYPIC VARIATION

Arnaud Martin^{1,2} and Virginie Orgogozo³

Nucleic Acids Research, 2019 1 doi: 10.1093/nar/gkz796

Gephebase, a database of genotype–phenotype relationships for natural and domesticated variation in Eukaryotes

Virginie Courtier-Orgogozo^{91,*}, Laurent Arnoult¹, Stéphane R. Prigent¹, Séverine Wiltgen² and Arnaud Martin^{3,*}

INTERFACE FOCUS

Replaying the tape of life in the twenty-first century

rsfs.royalsocietypublishing.org

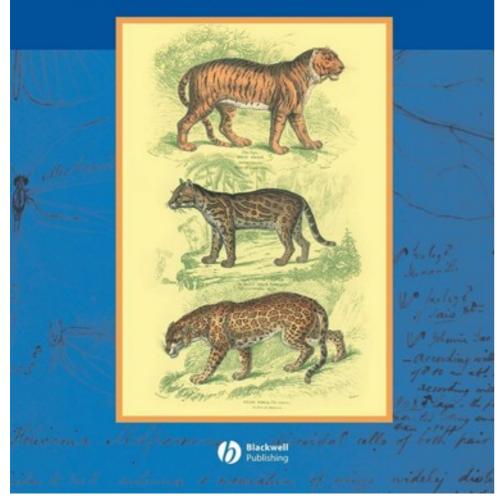
Virginie Orgogozo

The MUSIC of LIFE

Biology Beyond Genes

Denis Noble

SEAN B. CARROLL JENNIFER K. GRENIER SCOTT D. WEATHERBEE FROM DNA TO DIVERSITY MOLECULAR GENETICS AND THE EVOLUTION OF ANIMAL DESIGN SECOND EDITION



'Compelling . . . masterful . . . outstandingly good." Richard Dawkins, TLS WHY **EVOLUTION** IS E JERRY A. COYNE

https://www.youtube.com/watch?v=Ov30-7rXZCY Une brève histoire de l'évo-dévo (Denis Duboule)