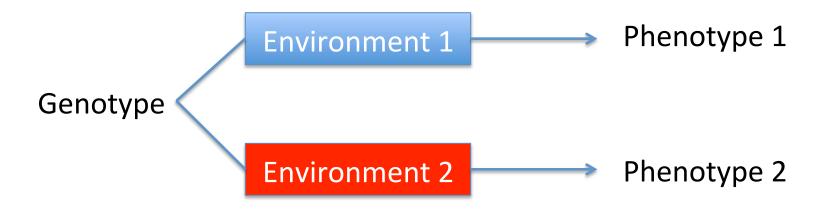
### Genetic bases of Phenotypic Plasticity

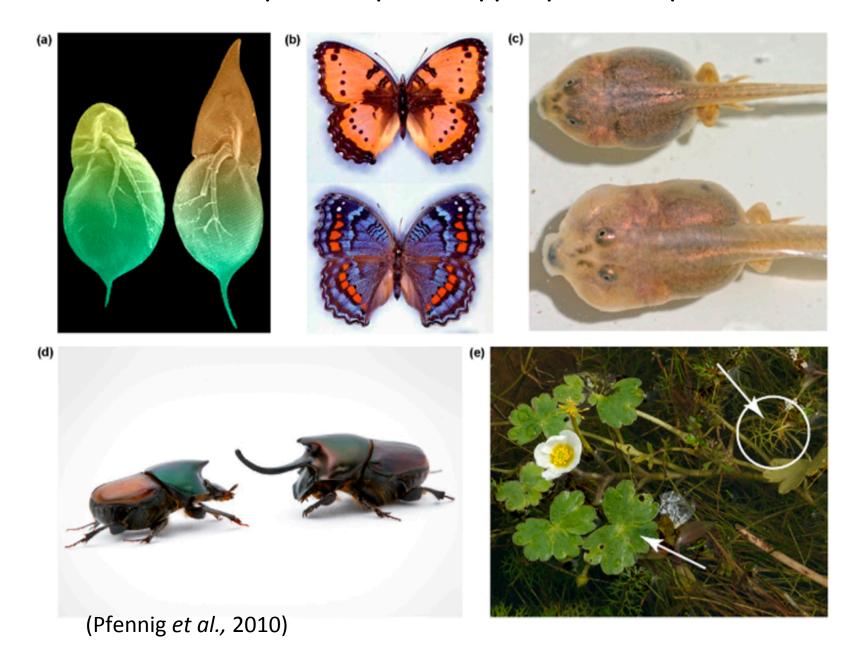
Jean-Michel Gibert
Developmental biology laboratory
CNRS-Sorbonne Université, IBPS
Paris

#### Phenotypic plasticity



« the property of a given genotype to produce different phenotypes in response to distinct environmental conditions » (Pigliucci, 2001)

### Examples of phenotypic plasticity



### Phenotypic plasticity and adaptation to environmental fluctuations



Summer Winter

Snow hare

#### Polyphenisms vs Polymorphisms

"In order to make the term 'polymorphism' more useful and precise, there is now a tendency to restrict it to genetic polymorphism. Since this would leave nongenetic variation of the phenotype without a designation, the term 'polyphenism' is here proposed for it. Polyphenism is discontinuous when definite castes are present (certain social insects) or definite stages in the life cycle (larvae vs. adults; sexual vs. parthenogenetic) or definite seasonal forms (dry vs. wet; spring vs. summer). Polyphenism may be continuous, as in the cyclomorphosis of fresh-water organisms and some other seasonal variation."

(Mayr, 1963)

#### Canalization

Canalization describes the ability of a organism to maintain the wild-type phenotype despite genetic and environmental variations (Waddington 1942; see also Schmalhausen 1949).

Environmental canalization can be seen as the opposite of phenotypic plasticity (Flatt, 2005).

However, environmental canalization and phenotypic plasticity are not mutually exclusive:

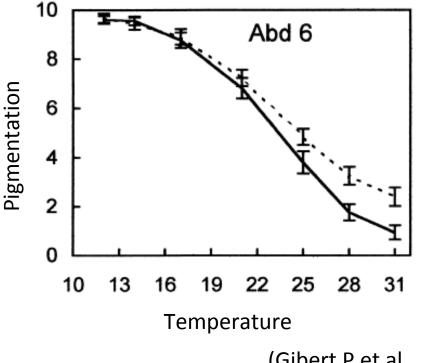
- Polyphenisms can be robust (canalized): no intermediate phenotype observed between alternative morphs.
- A plastic molecular response to environmental variation can be used to maintain the phenotype.

Role in evolution: release of accumulated cryptic genetic variation upon decanalization.

### The reaction norm: a major tool to represent phenotypic plasticity

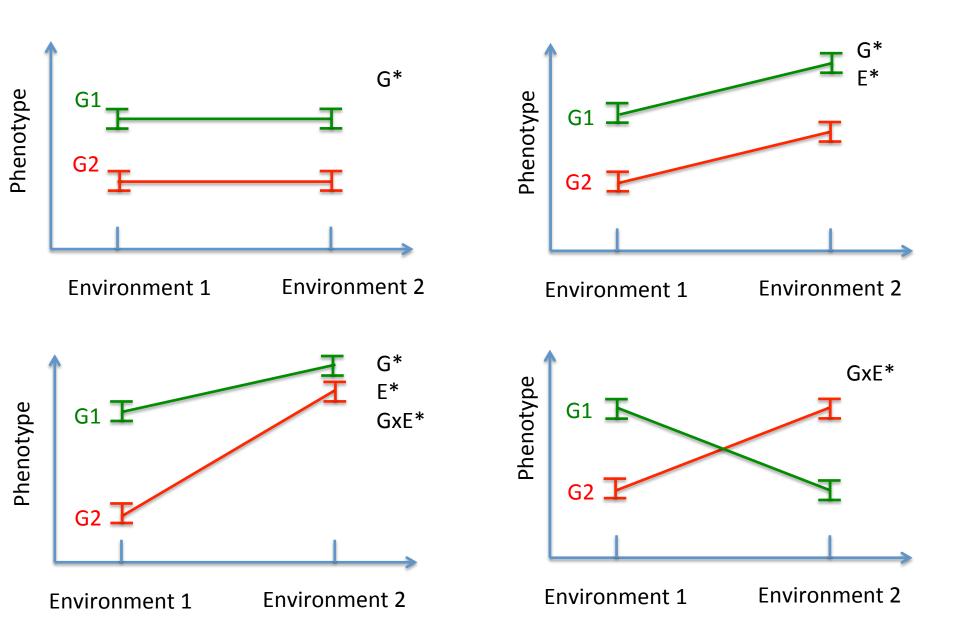
The reaction norm: graph representing the phenotype as a function of the environment

First drawn by Woltereck (1909) who however mis-interpreted them as the distinction between Genotype and Phenotype was made only in 1911 by Johannsen.



(Gibert P et al., 2000)

#### Analyses of reaction norms (using Analysis of Variance)



#### The genetics of phenotypic plasticity:

The idea that plasticity is under genetic control was initially developed by Bradshaw (1965)

Two genetic mechanisms were proposed (Via, 1995):

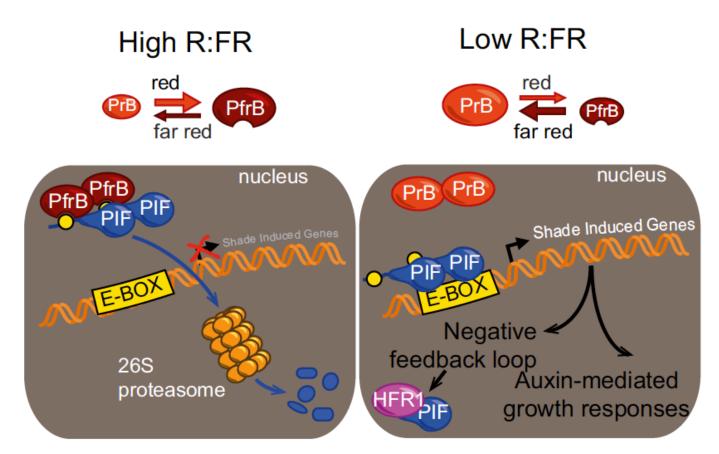
-allelic sensitivity

-gene regulation

But these to categories may blur.

### How is the environmental cue perceived and integrated in gene regulatory network?

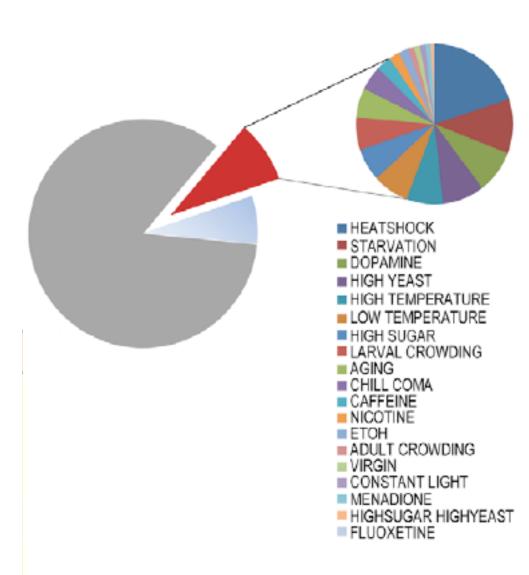
Shade avoidance in plants: role of phytochromes in plants: detection of Red/Far Red ratio



(Hersch et al., 2014)

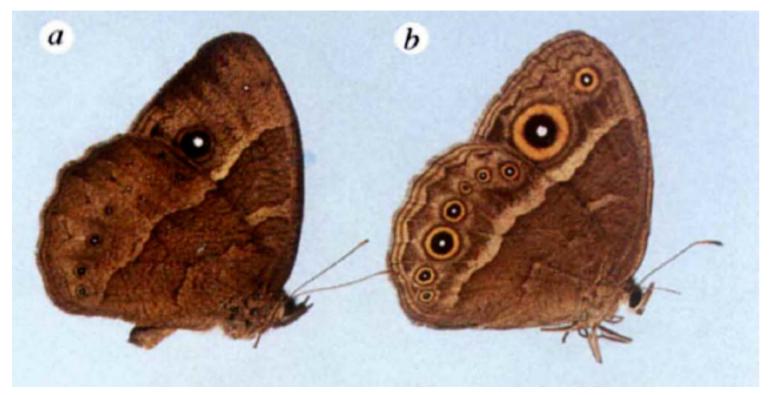
# Environmental conditions can strongly affect the transcriptome

Study of Drosophila adults transcriptome in 20 different environmental conditions: 15% of expressed genes show transcriptional plasticity (Zhou et al., 2012).



# Environmental conditions modulate the expression of developmental regulatory genes:

#### Bicyclus anynana polyphenism

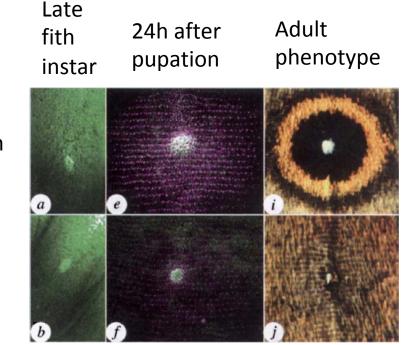


Dry season form 17°C

Wet season form 27°C

(Brakefield et al., 1996)

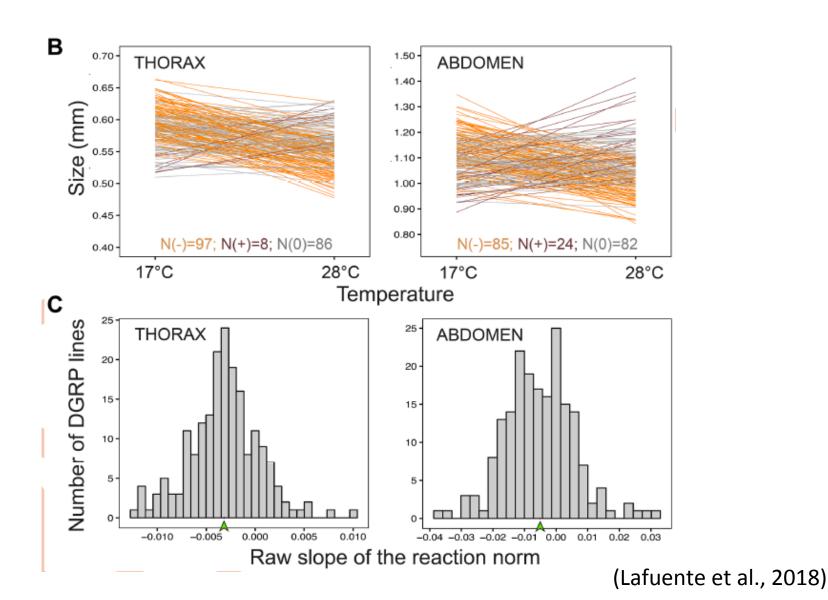
Distalless expression modulation in the butterfly Bicyclus anynana correlates with wing eyespot plasticity (Brakefield et al., 1996). Functional analyses show that Distalless is involved in eyespot formation (Monteiro et al., 2013).



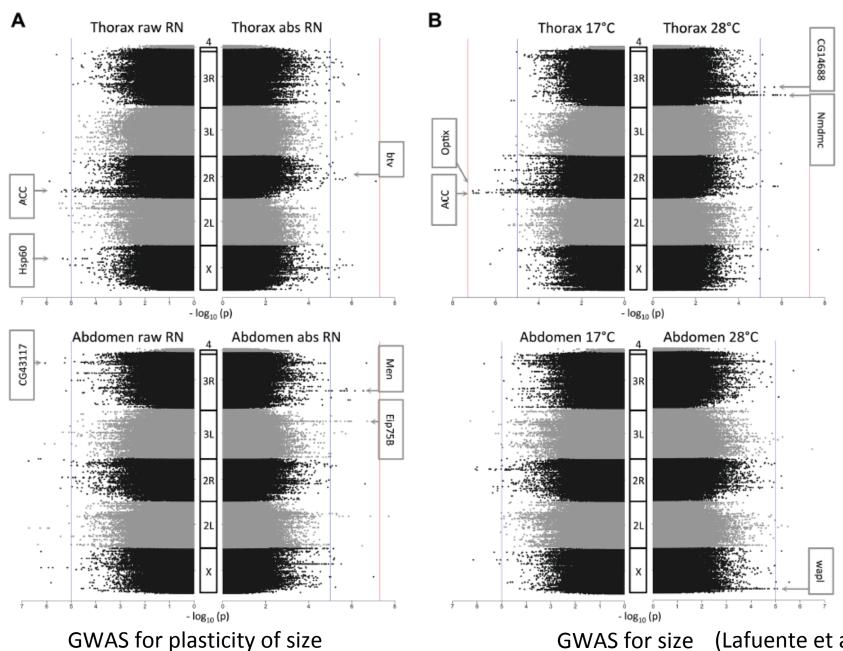
Wet-season form

Dry-season form

### Mapping genetic variation for plasticity: example of size thermal plasticity in *Drosophila*



#### Identification of SNPs affecting size thermal plasticity or size



GWAS for size (Lafuente et al., 2018)

#### Epigenetic bases of phenotypic plasticity

Epigenetics: The study of mitotically and/or meiotically heritable changes in gene function that cannot be explained by changes in DNA sequence" (Russo et al. 1996)

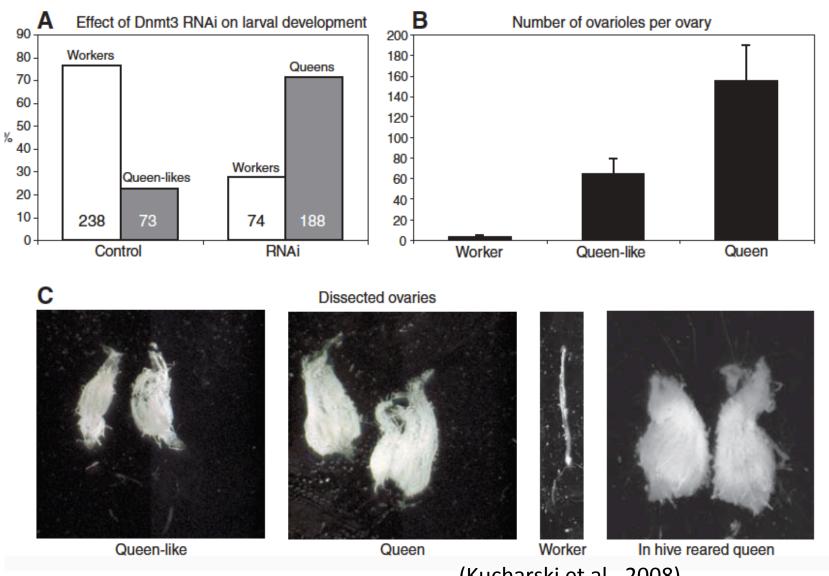


High fat diet induced chromatin remodeling in mouse liver (Leung et al., 2014).

Casts in the carpenter ant Camponotus floridanus and histone acetylation (Simola et al., 2012, 2016).

Honeybee casts and DNA methylation (Kucharski et al., 2008).

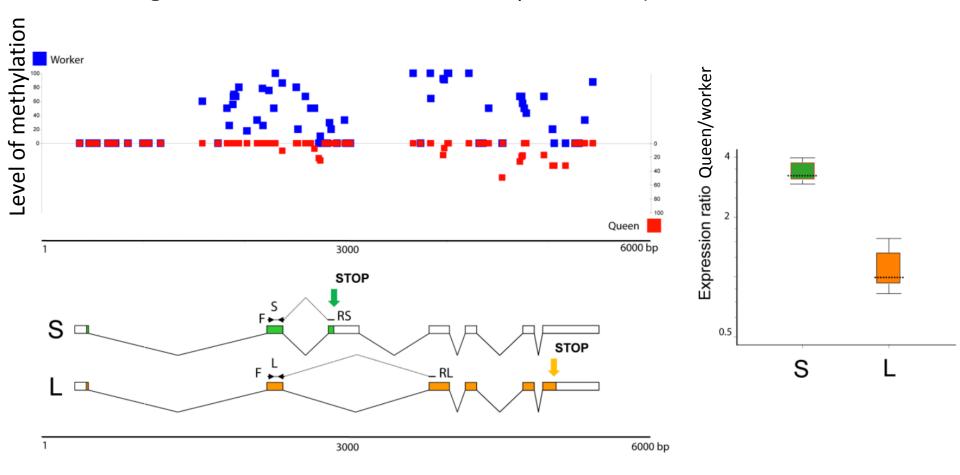
# Effect of inactivation of DNA methyl transferase Dnmt3 in the honeybee



(Kucharski et al., 2008)

#### Difference of DNA methylation between honeybee casts

Over 550 genes show differential methylation in queen and worker brains.



exemple of the gene GB18602 in queen and worker brains

(Lyko et al., 2010)

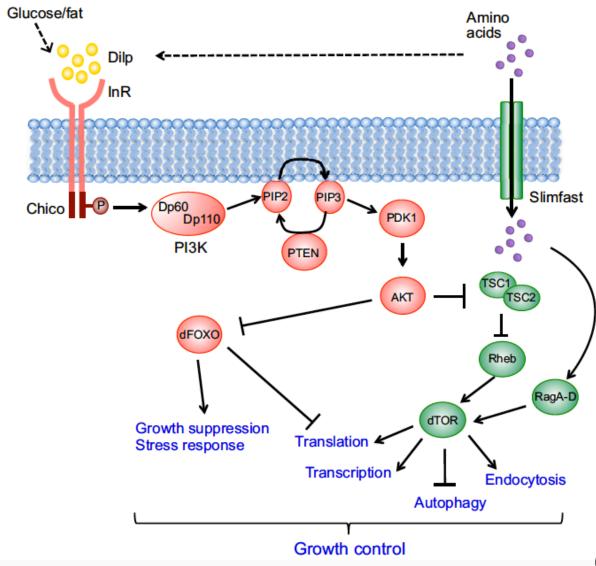
#### Hormonal bases of phenotypic plasticity

Corticotropin releasing hormone and reduction of developmental time in desiccating environment in amphibians (Denver, 1997).

Ecdysone and *Bicyclus anynana* eyespot plasticity (Monteiro et al., 2015).

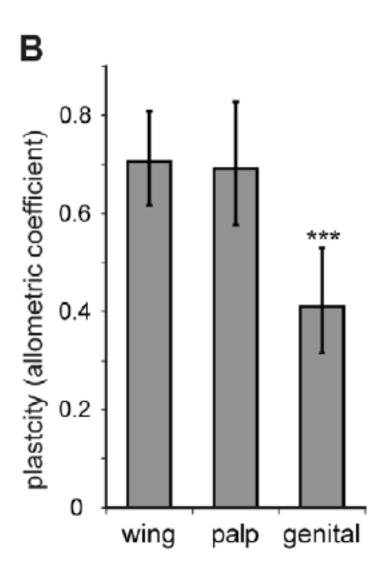
Insulin and nutritional plasticity in *Drosophila* (Tang et al., 2011).

### The Insulin pathway in Drosophila



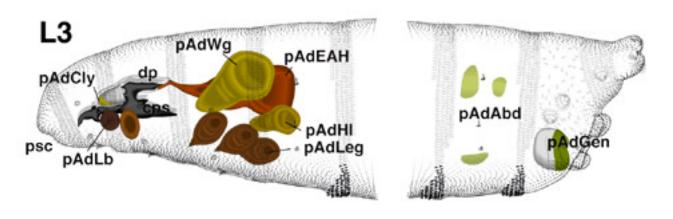
(Shim et al., 2013)

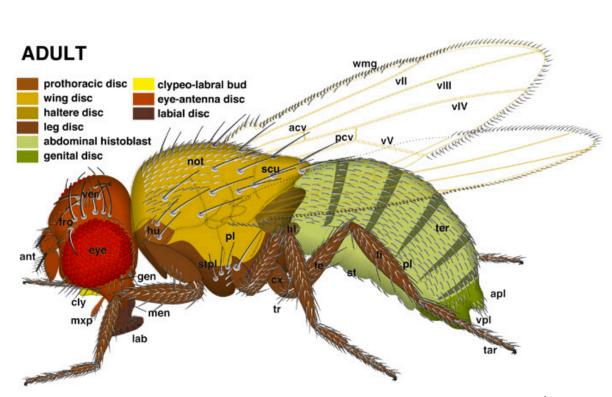
### Nutritional plasticity differs between appendages in *Drosophila*



(Tang et al, 2011)

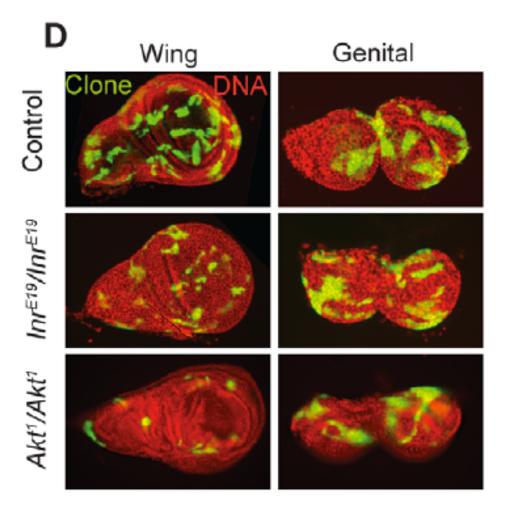
#### The imaginal discs of Drosophila



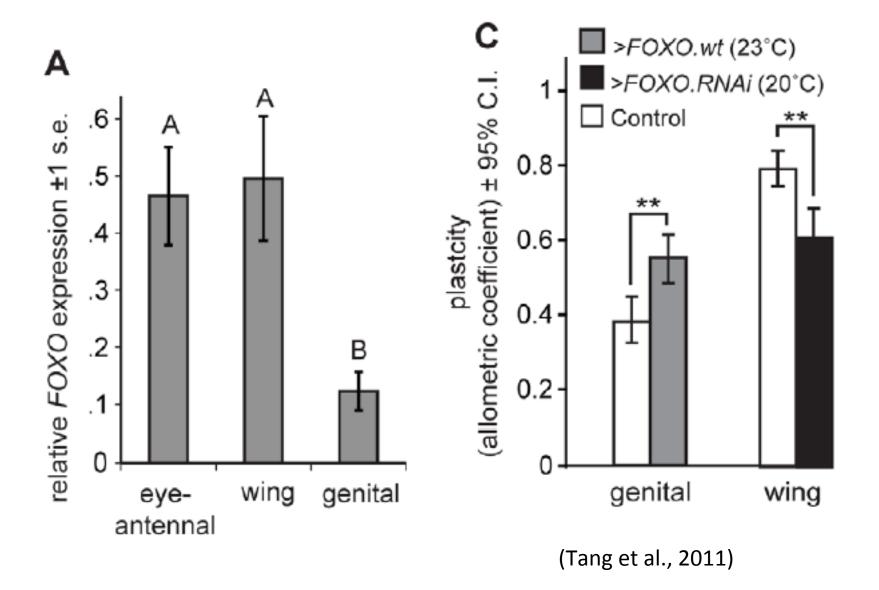


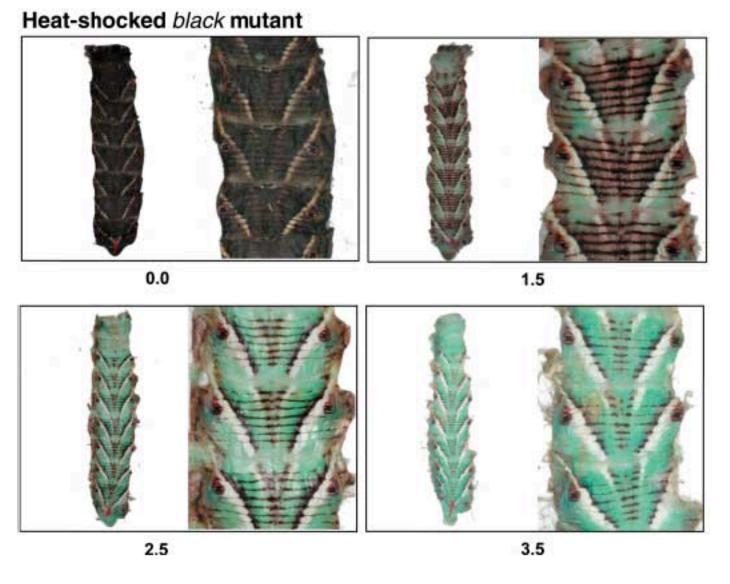
(Hartenstein, 1993)

Loss of activity of the insulin pathway has different effects depending on the appendage in *Drosophila* 

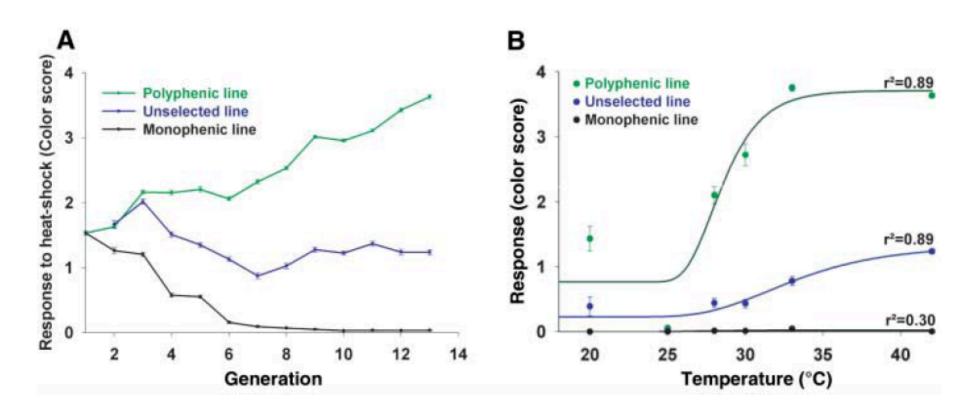


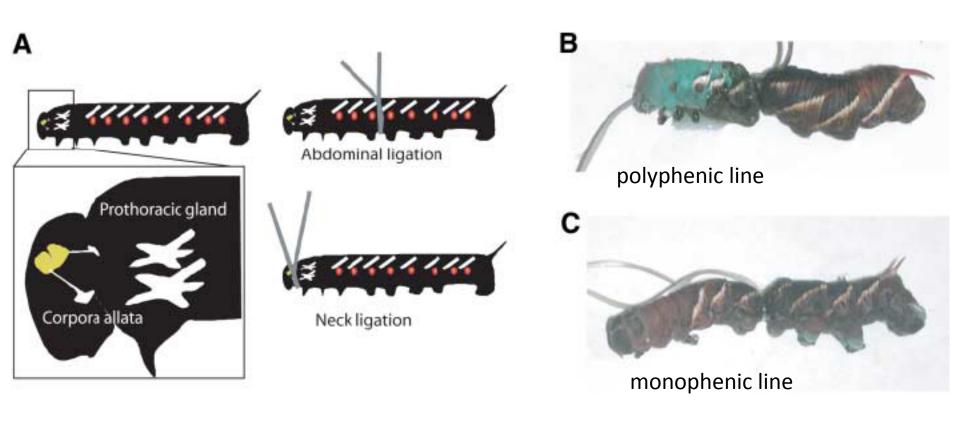
### Differential expression of *foxo* explains the difference of nutritional plasticity between the wing and the genitals

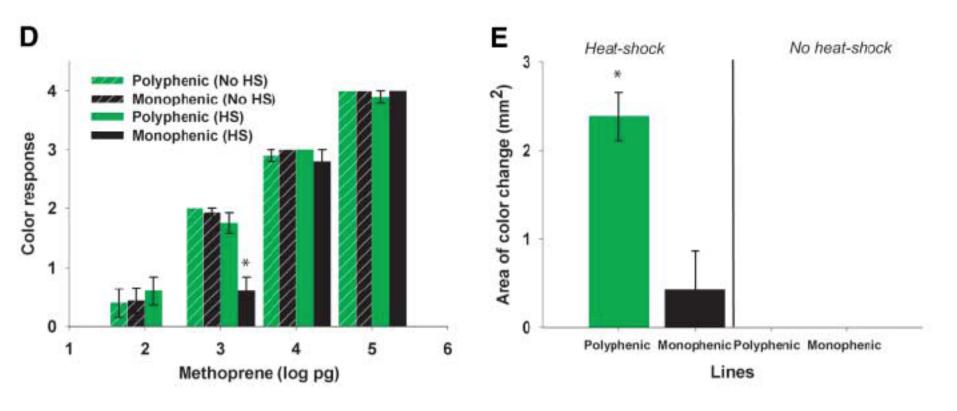




(Suzuki and Nijhout, 2006)



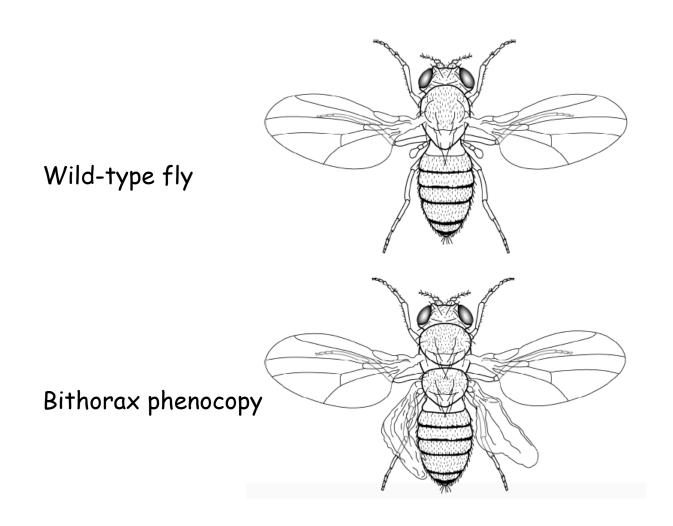




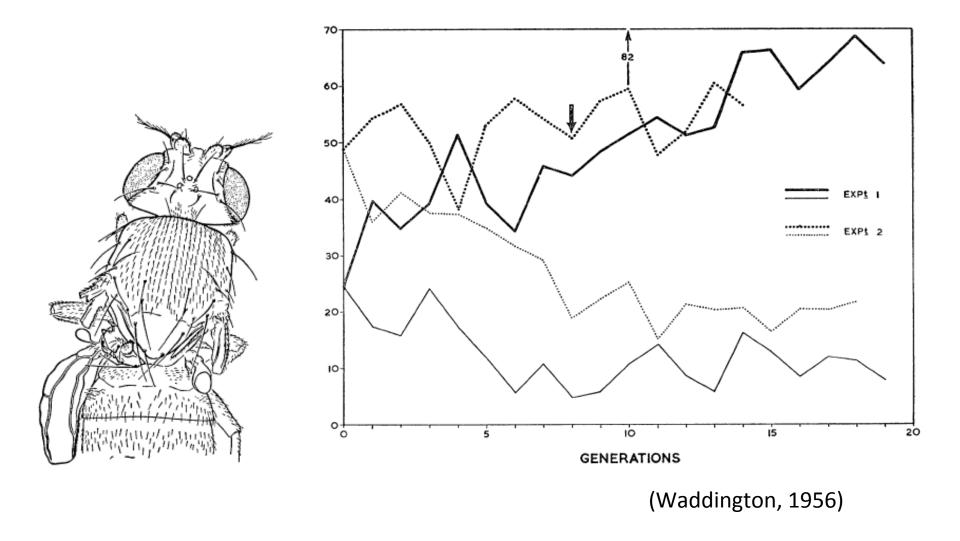
The monophenic line has a reduced JH sensitivity at high temperature.

The polyphenic line has a higher juvenile hormone (JH) titer at high temperature;

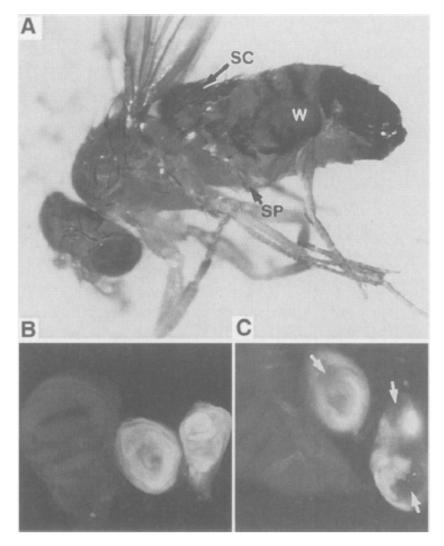
#### Genetic assimilation of ether induced Bithorax phenocopies



### Selection for increased and decreased proportions of ether induced Bithorax phenocopies

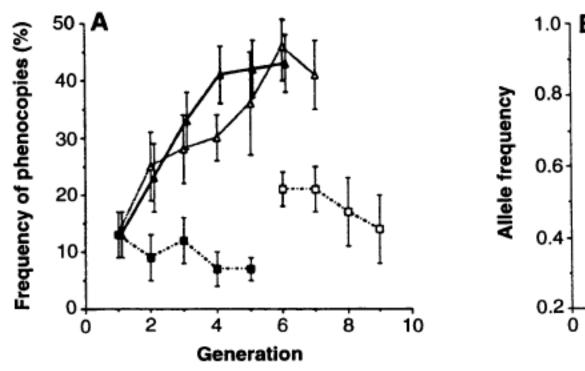


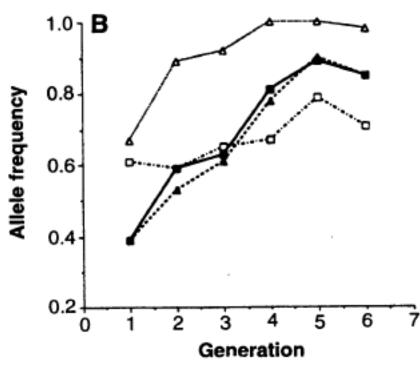
# Ether induced Bithorax phenocopies correspond to loss of expression of Ubx



(Gibson et Hogness, 1996)

### Selection of *Ubx* alleles during genetic assimilation of the Bithorax phenotype





(Gibson et Hogness, 1996)

# Genetic assimilation in the polyphenic butterfly Bicyclus anynana

17°C 27°C b C e

WT line

Low line

High line

#### Genetic assimilation

Defined by Waddington (1952, 1959):

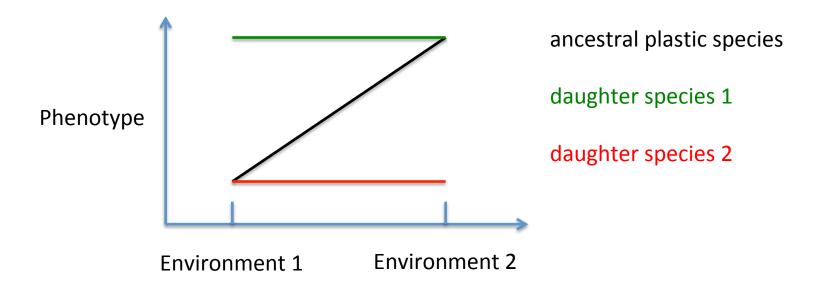
Environmental changes can reveal cryptic genetic variation and induce new phenotypes in some individuals.

This genetic variation can be selected allowing to fix a phenotype initially observed only in particular environmental conditions.

Some of Waddington's experiments were repeated with isogenic or outbred stocks and it was shown that genetic variation was necessary for genetic assimilation (Bateman, 1959).

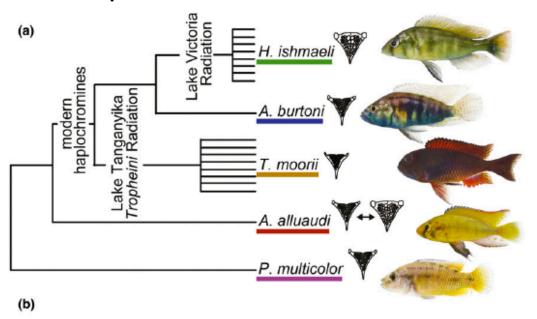
However, in a recent study, it was shown that de novo mutation induced by the environment (heatshock) can be involved in genetic assimilation (Fanti et al., 2017).

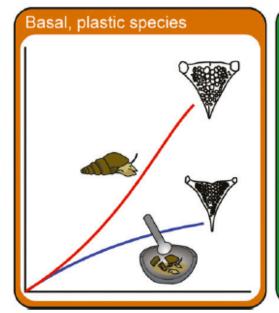
The idea that an ancestral plastic species can be at the origin of divergent species after fixation of the alternative morphs has been proposed by West-Eberhard as "the flexible stem hypothesis" (2003).

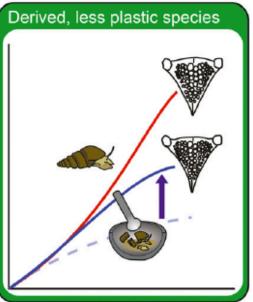


It is based on genetic assimilation discovered by Waddington. The "flexible stem hypothesis" is also called "plasticity first evolution".

#### Plasticity first evolution in cichlid fishes

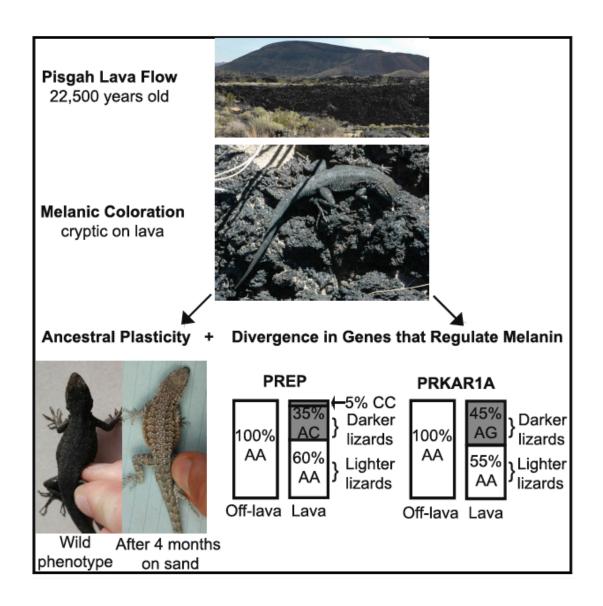


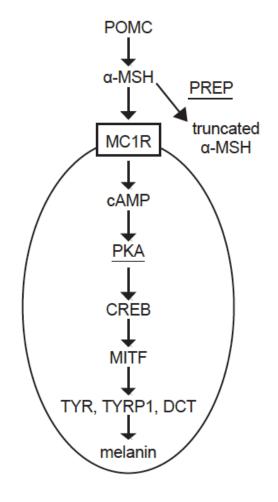




(Gunter et al., 2017)

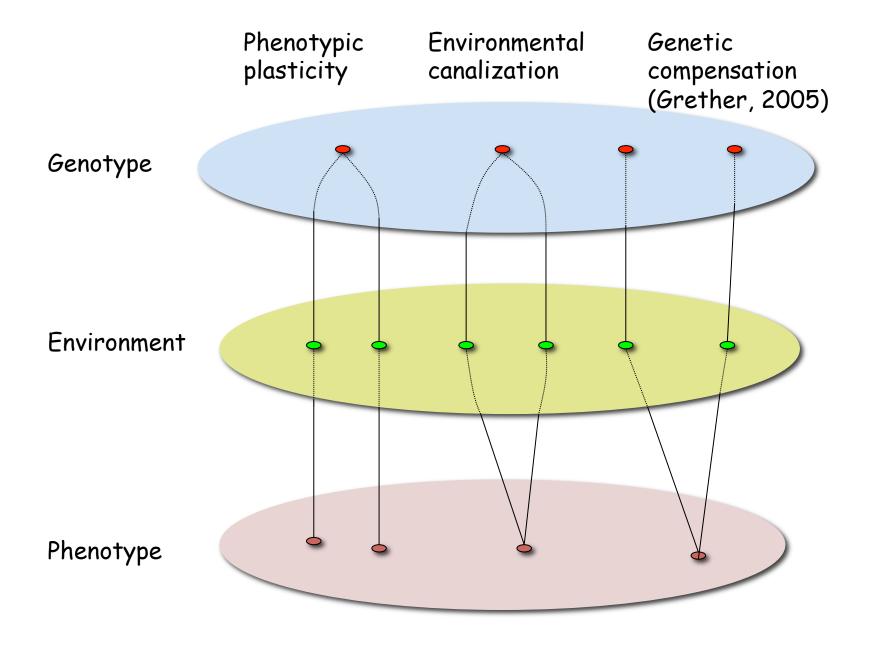
### Plasticity first evolution in the lizard Uta stansburiana





(Corl et al., 2018)

## The environment in the genotype-phenotype relation



# Phenotypic plasticity is not always linked to differential gene expression



Influence of carotenoids present in the diet on pigmentation

A study of the genetic bases of phenotypic plasticity: Pigmentation thermal plasticity in *Drosophila melanogaster* 

Jean-Michel Gibert\*, Emmanuèle Mouchel-Vielh\*, Sandra de Castro and Frédérique Peronnet

> Developmental biology laboratory CNRS-Sorbonne Université, IBPS Paris

### Drosophila and temperature

In Drosophila temperature affects many traits:

Developmental rate,

Size,

Ovariole number,

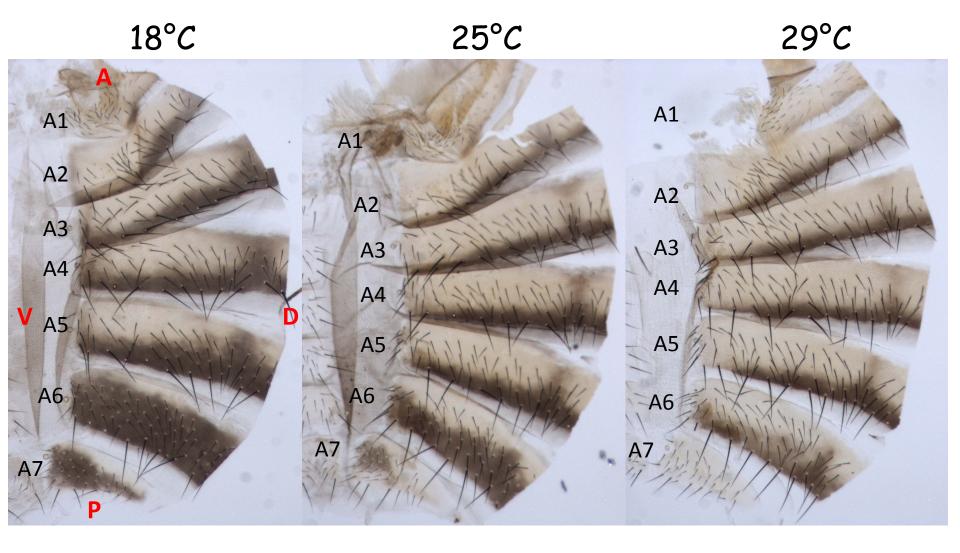
Bristle number,

Reproductive diapause,

Pigmentation

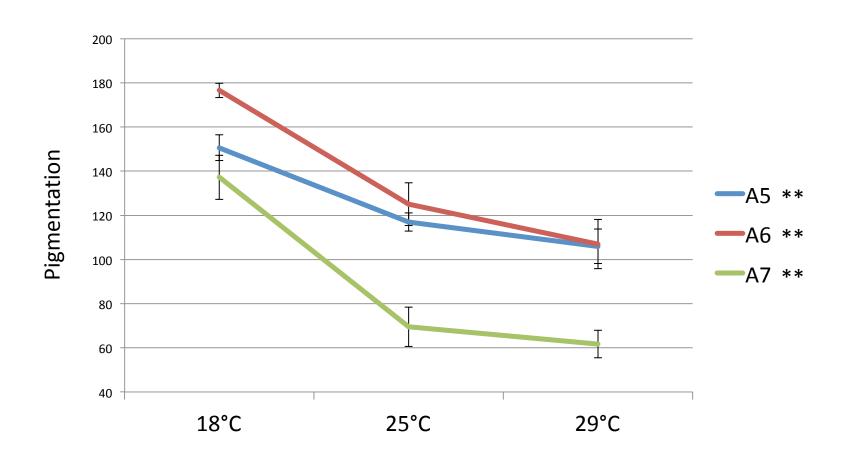
(Bouletreau-Merle et al., 2003; David et al., 2004; Schmidt et al., 2005; Trotta et al, 2006).

# Temperature sensitivity of female abdominal pigmentation



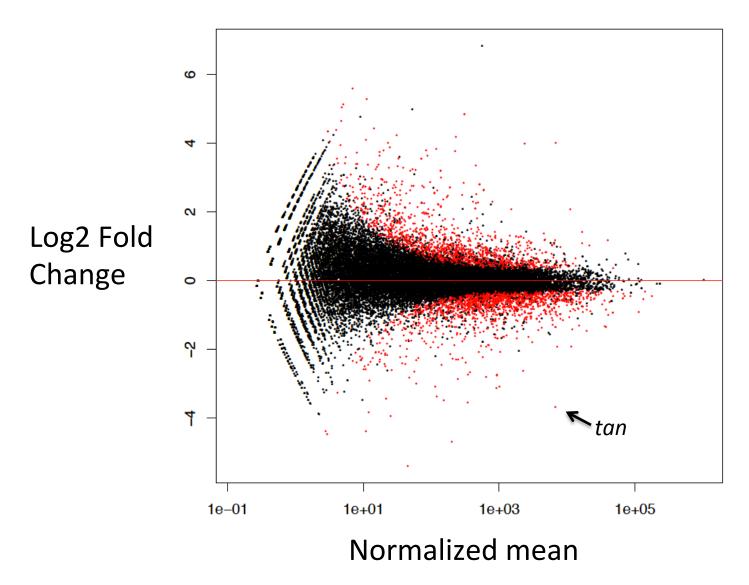
Drosophila melanogaster, isogenic line w<sup>1118</sup>

### Reaction norms of female abdominal pigmentation



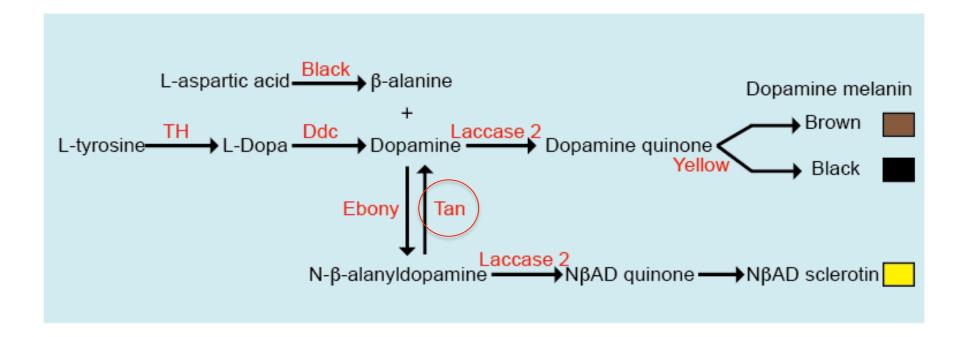
Quantification of pigmentation from mounted cuticles (ImageJ).

# Transcriptome analysis at 18°C and 29°C in young adult female posterior abdominal epidermis

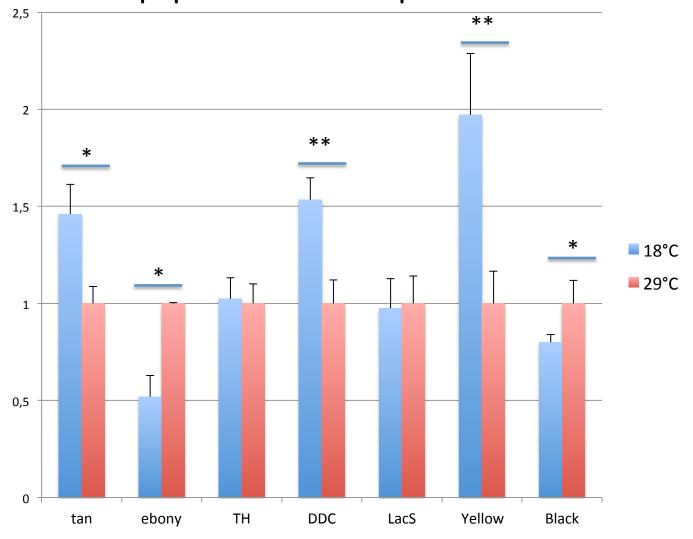


3000 transcripts=2097 genes (p<0.05), 200 transcripts (p<1E-10)

### Cuticular pigment synthesis pathway

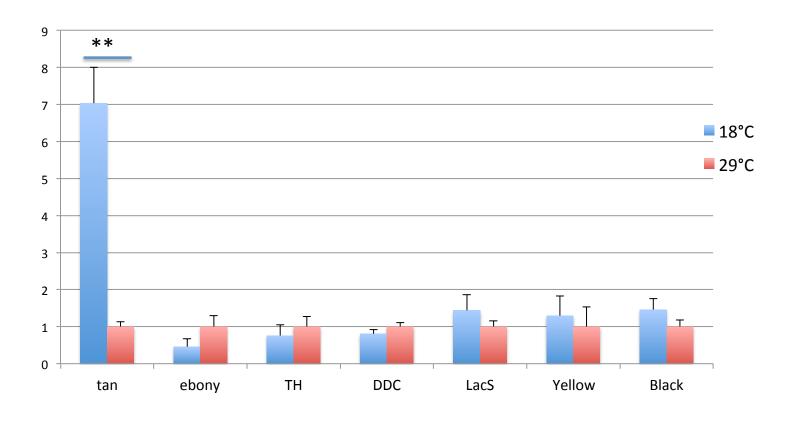


The expression of several pigmentation enzyme genes is modulated in pupal abdominal epidermis



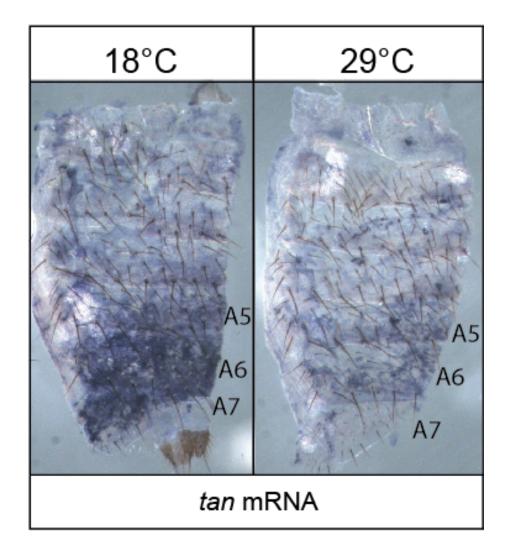
RT-qPCR on 3 biological replicates, normalized with Act5c and RP49

The expression of expression of tan is dramatically modulated by temperature in the abdominal epidermis of freshly hatched females



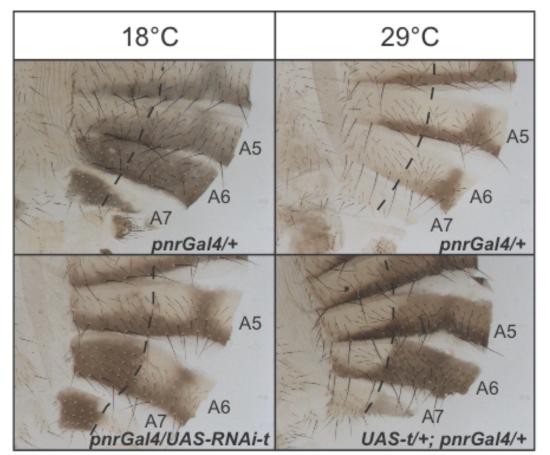
RT-qPCR on 3 biological replicates, normalized with Act5c and RP49

# tan expression is modulated by temperature



# Modulation of tan expression by temperature is essential for female abdominal pigmentation plasticity

control at 18°C



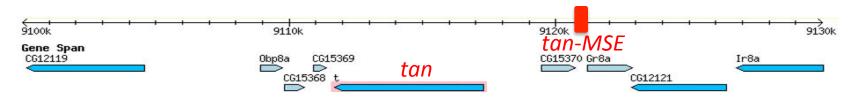
control at 29°C

tan LOF at 18°C

tan GOF at 29°C

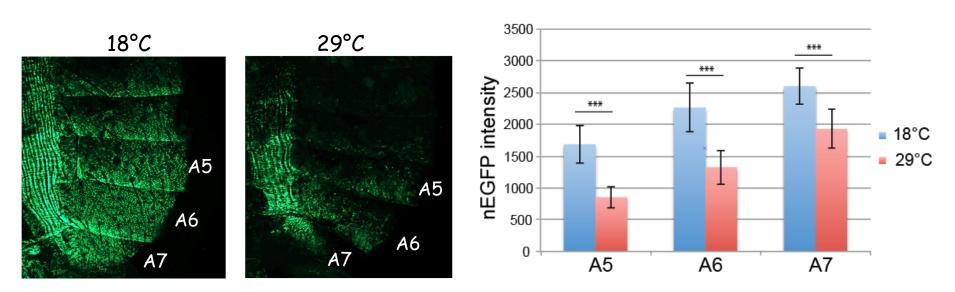
# The effect of temperature on tan expression is mediated by the tan-MSE enhancer

#### Structure of tan genomic region

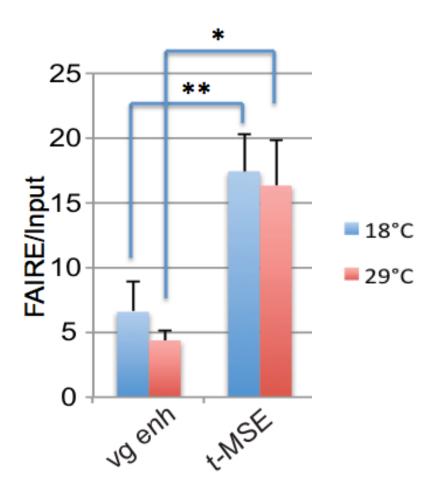


after Jeong et al., 2008

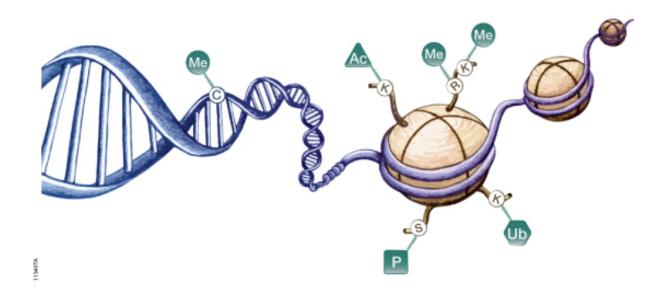
#### GFP expression in a tan-MSE GFP line is sensitive to temperature



FAIRE (Formaldehyde Assisted Isolation of Regulatory Element)-qPCR shows that  $t\_MSE$  is less compacted than vg enhancer

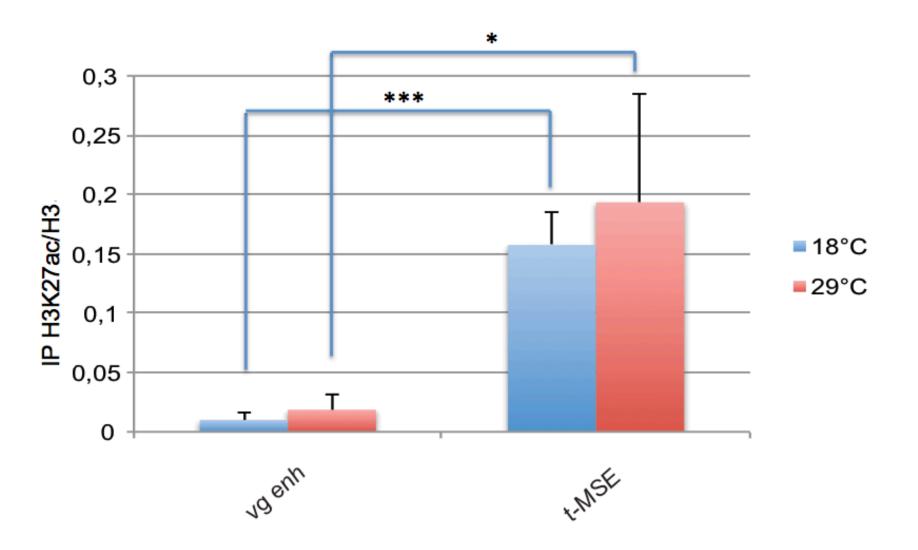


# Epigenetic marks analysed in tan region

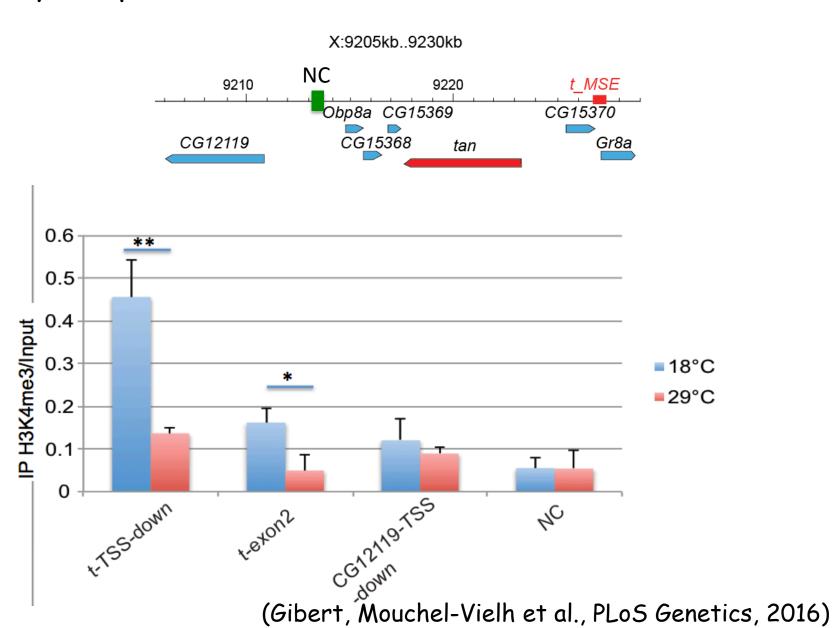


Mark	location	Indication
H3K4me3	Promoter	Active genes
H3K27ac	Enhancer	Active enhancer

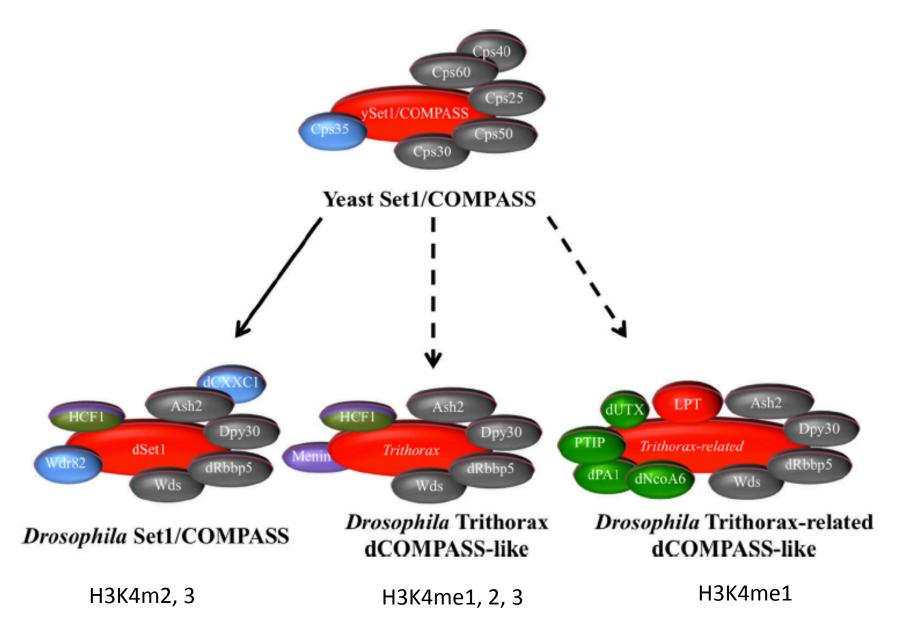
t\_MSE is enriched in H3K27ac, but this mark is not modulated by temperature



# H3K4me3 on tan promoter is strongly modulated by temperature

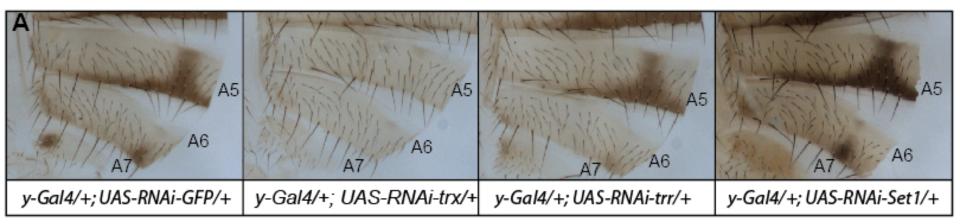


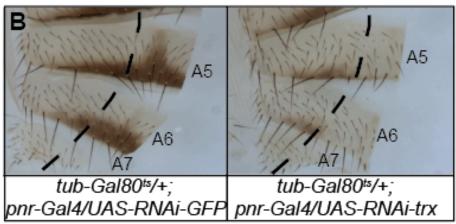
## Complexes involved in H3K4 methylation



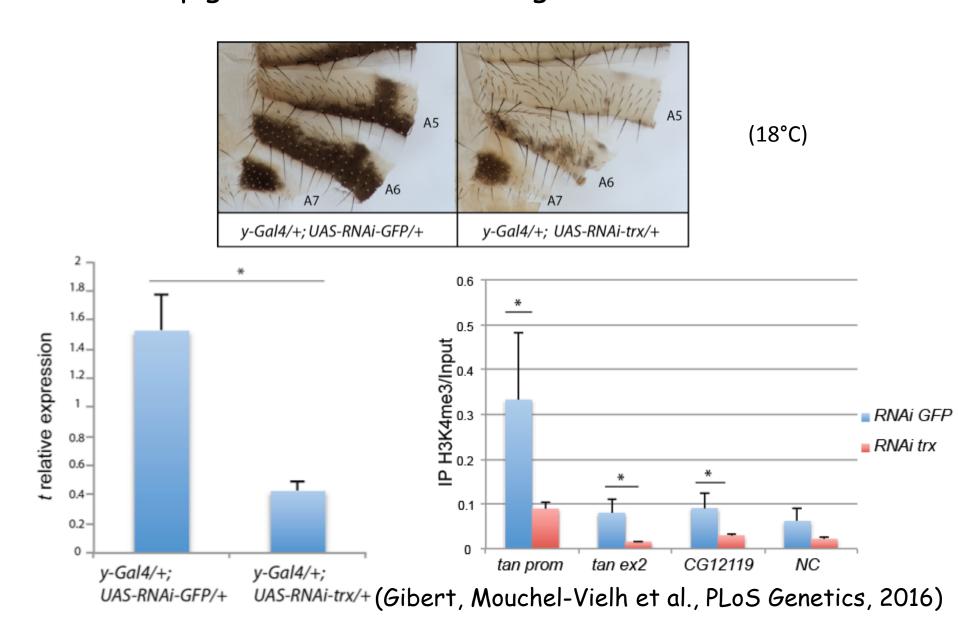
(Mohan et al., 2011; Herz, et al., 2012; Hallson et al., 2012; Tie et al;, 2014; Smith et al., 2004)

#### Female pigmentation phenotypes of H3K4 methyl-transferase LOF





# The H3K4 methyl-transferase Trithorax is involved in female abdominal pigmentation and tan regulation



#### Conclusions

tan temperature sensitive expression plays a major role in female abdominal pigmentation plasticity. Modulation of *yellow* expression by temperature is also involved (Gibert et al., Scientific Reports, 2017).

The effect of temperature is mediated at least partly by  $t\_MSE$ . However we did not detect modification of chromatin structure on  $t\_MSE$ .

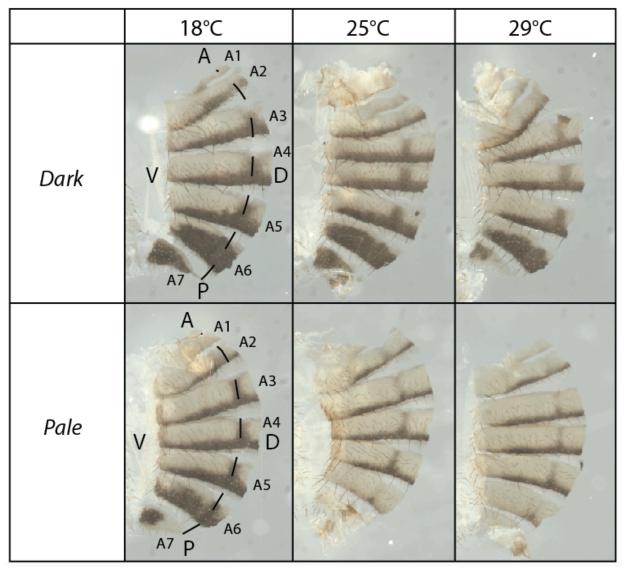
In contrast H3K4me3 level is strongly modulated by temperature on tan promoter.

The H3K4me3 methyl-transferase involved is likely Trithorax as it regulates female abdominal pigmentation, tan expression and H3K4me3 level on tan promoter.

25°C
Pale line Dark line

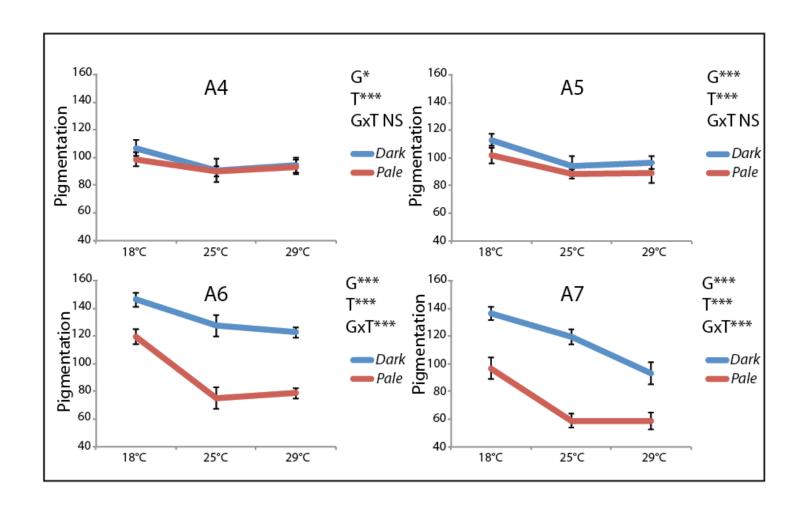


#### Phenotypes of the Dark and Pale lines at different temperatures

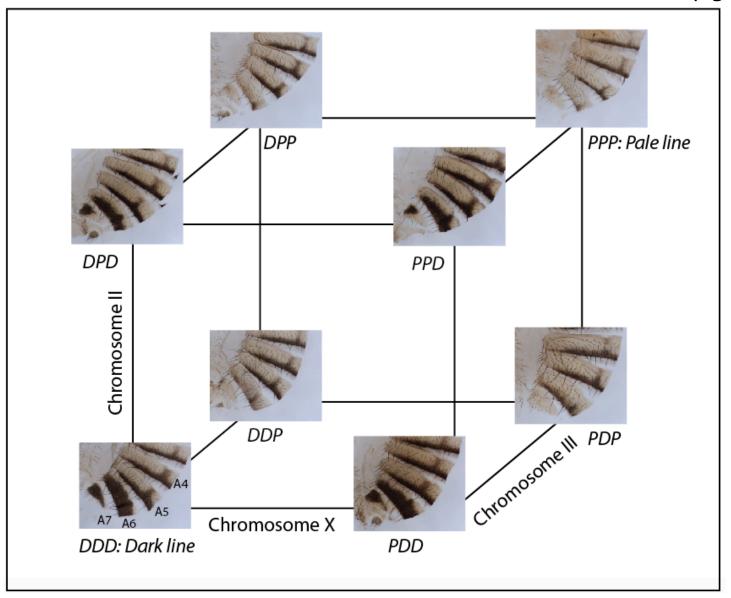


(De Castro et al., PLOS Genetics, 2018)

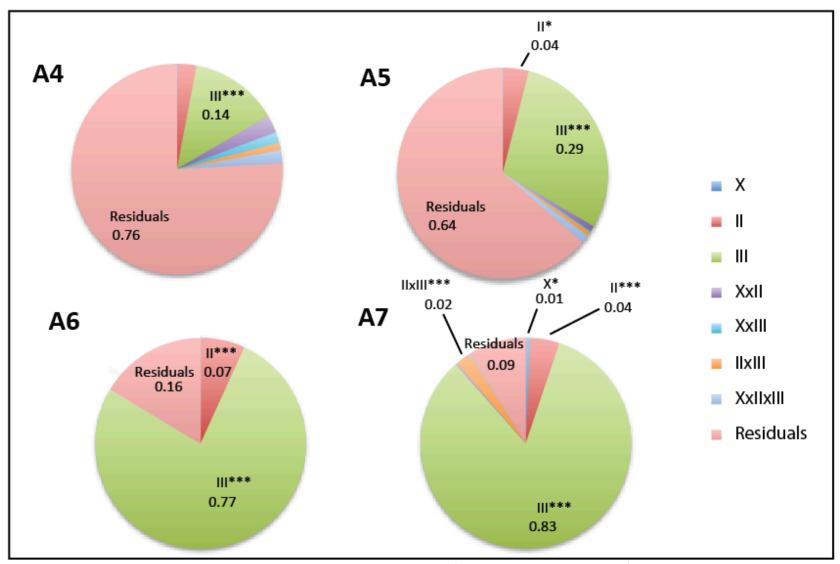
#### Reaction norms of the Dark and Pale lines



Effects of the different chromosomes of the Dark and Pale lines on pigmentation



The third chromosome plays a major role in the difference of pigmentation between the *Dark* and *Pale* lines



### bab, a major QTL for female abdominal pigmentation

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#### Quantitative Trait Loci Responsible for Variation in Sexually Dimorphic Traits in *Drosophila melanogaster*

Artyom Kopp,\* Rita M. Graze,† Shizhong Xu,‡ Sean B. Carroll\* and Sergey V. Nuzhdin<sup>†,1</sup>

2003

\*Howard Hughes Medical Institute and Laboratory of Molecular Biology, University of Wisconsin, Madison, Wisconsin 53706,

†Section of Evolution and Ecology, University of California, Davis, California 95616 and †Department of Botany and

Plant Sciences, University of California, Riverside, California 92521

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PLOS GENETICS

# Composite Effects of Polymorphisms near Multiple Regulatory Elements Create a Major-Effect QTL

2011

Ryan D. Bickel<sup>1,2\*</sup>, Artyom Kopp<sup>3</sup>, Sergey V. Nuzhdin<sup>2</sup>

1 School of Biological Science, University of Nebraska – Lincoln, Lincoln, Nebraska, United States of America, 2 Program in Molecular and Computational Biology, Department of Biological Sciences, University of Southern California, Los Angeles, California, United States of America, 3 Department of Evolution and Ecology, University of California Davis, Davis, California, United States of America

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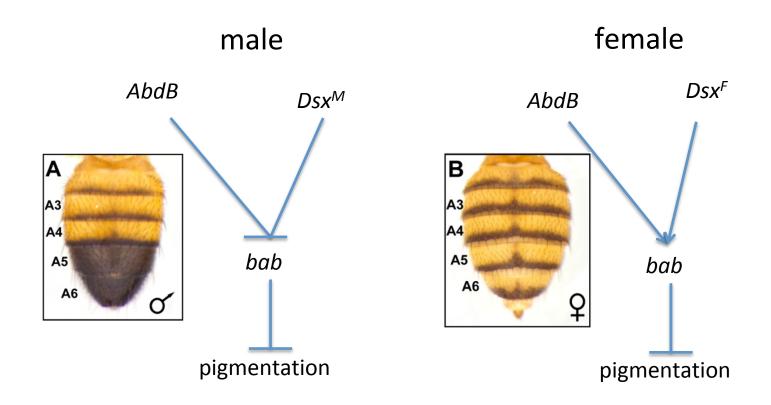
# Recurrent Modification of a Conserved *Cis*-Regulatory Element Underlies Fruit Fly Pigmentation Diversity

William A. Rogers<sup>1</sup>, Joseph R. Salomone<sup>1</sup>, David J. Tacy<sup>1</sup>, Eric M. Camino<sup>1</sup>, Kristen A. Davis<sup>1</sup>, Mark Rebeiz<sup>2</sup>, Thomas M. Williams<sup>1,3</sup>\*

2013

1 Department of Biology, University of Dayton, Dayton, Ohio, United States of America, 2 Department of Biological Sciences, University of Pittsburgh, Pittsburgh, Pennsylvania, United States of America, 3 Center for Tissue Regeneration and Engineering at Dayton, University of Dayton, Dayton, Ohio, United States of America

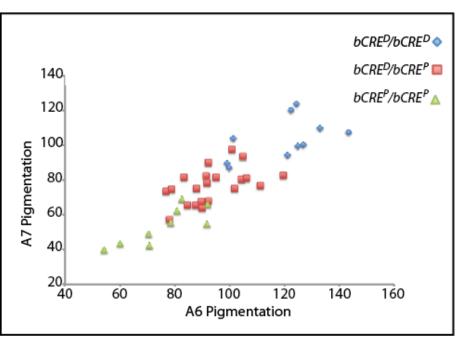
### bab and sex-specific pigmentation

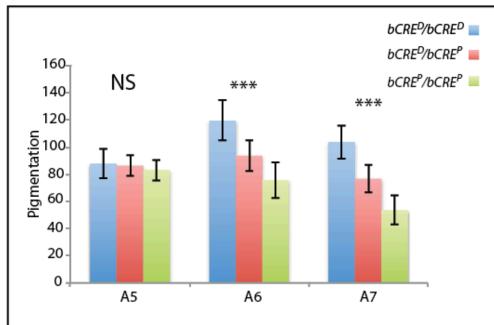


#### bab dimorphic CRE in $w^{1118}$ , Dark and Pale

Canton/w Dark Pale	TTTTAAGACCATAAATTCAGCTCACTCTCTCTCTCTCTCT	480
Canton/w Dark Pale	AbdB1 AbdB2 AbdB3 CTTTTATTACTCTTAATATAAAAAAGCTGGCTAGATGCGGGCCAGCTGTAAAAATGCACG CTTTTATTACTCTTAATATAAAAAAAGCTGGCTAGATGCGGGCCAGCTGTAAAAATGCACG CTTTTATTACTCTTAATATAAAAAAAGCTGGCTAGATGCGGGCCAGCTGTAAAAATGCACG	540 520 540
Canton/w Dark Pale	AbdB4 CGGTCATAAAAAGTTGCAGGAGGCATGTTGCCAGTTGCCTGCAACCGGCAACATTCGCAG CGGTCATAAAAAAGTTGCAGGAGGCATGTTGCCAGTTGCCTGCAACCGGCAACATTCGCAG CGGTCATAAAAAAGTTGCAGGAGGCATGTTGCCAGTTGCCTGCAACCGGCAACATTCGCAG	544
Canton/w Dark Pale	AbdB5 D Dsx1  AACAGCAGCAACATCGTAAAATAACTTCTTCCTCTGCGGTCTGAGTTTGGCCGCAACAAT  AACAGCAGCAACATCGTAAAATAACTTCTTCCTCTGCGGTCTGAGTTTGGCCGCAACAAT  AACAGCAGCAACATCGTAAAATAACTTCTTCCTCTGCGGTCTGAGTTTGGCCGCAACAAT	660 604 660
Canton/w Dark Pale	AbdB6  GTTGCTGCATTTATTCGTATTATTACATTTTAATGAATAATTCTAATTATATGCAAC  GTTGCTGCATTTATTCGTATTATTATTACATTTTAATGAATAATTCTAATTATATGCAAC  GTTGCTGCATTTATTCGTATTATTATTACATTTTAATGAATAATTCTAATTATATGCAAC	720 664 720
Canton/w Dark Pale	AbdB7 TTGAATAAGCCCGCCGATGCCAATAAAAAGCGGCGTGGCAAAGTGGAGTGGACTGGGTTT TTGAATAAGCCCGCCGATGCCAATAAAAAGCGGCGTGGCAAAGTGGAGTGGACTGGGTTT TTGAATAAGCCCGCCGATGCCAATAAAAAAGCGGCGTGGCAAAGTGGAGTGGACTGGGTTT	780 724 780
Canton/w Dark Pale	Abdb8  GTGTGGCGCCCCTGCTAGTGGCACATAAAAATTGGCGCAAGTTAATTGTGGTAGTTATTT  GTGTGGCGCCCCTGCTAGTGGCACATAAAAATTGGCGCAAGTTAATTGTGGTAGTTATTT  GTGTGGCGCCCCTGCTAGTGGCACATAAAAATTGGCGCAAGTTAATTGTGGTAGTTATTT	840 784 840
Canton/w Dark Pale	AbdB9 AbdB10  GCTGTTTTGCCATTTGGTCATTTTACAATTTTACCATTTCAGCCACAACTTTTCGCACTG GCTGTTTTGCCATTTGGTCATTTTACAATTTTACCATTTCAGCCACAACTTTTCGCACTG GCTGTTTTGCCATTTGGTCATTTTACAATTTTACCATTTCAGCCACAACTTTTCGCACTG	844

# Genotyping of the F2 of a *Dark x Pale* cross shows that the *bab* locus is linked to the pigmentation phenotype



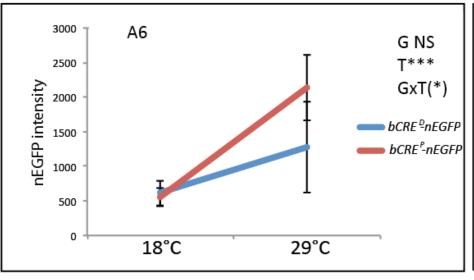


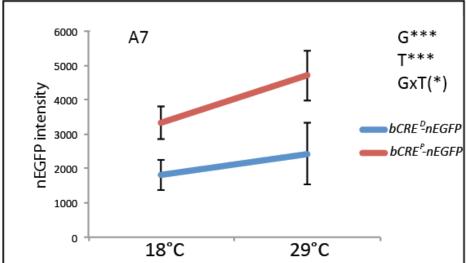
Comparison of the activities of bab dimorphic enhancers from the Dark and Pale lines

29°C 18°C Α6 bCRE<sup>D</sup>nEGFP A5 1 A5 bCRE <sup>P</sup>-nEGFP

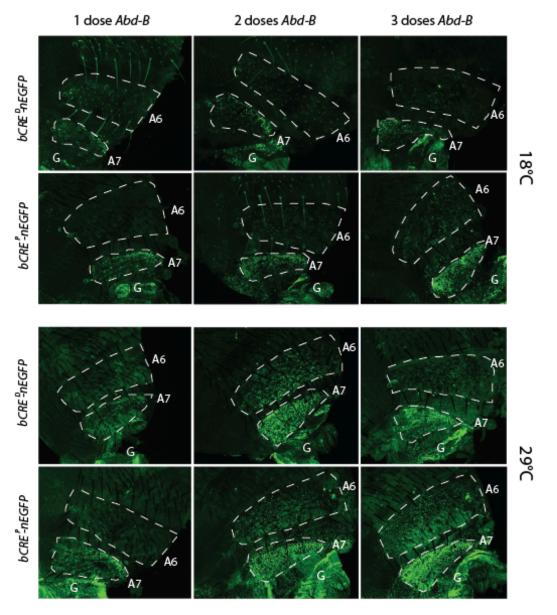
(De Castro et al., PLOS Genetics, 2018)

Comparison of the activities of bab dimorphic enhancers from the Dark and Pale lines



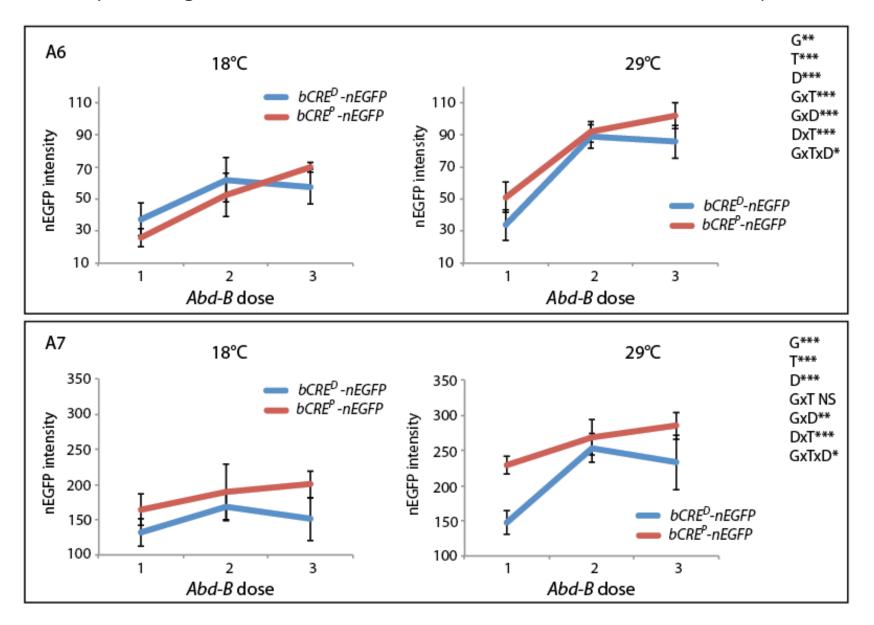


#### Impact of genetic variation in the enhancer on its activation by AbdB



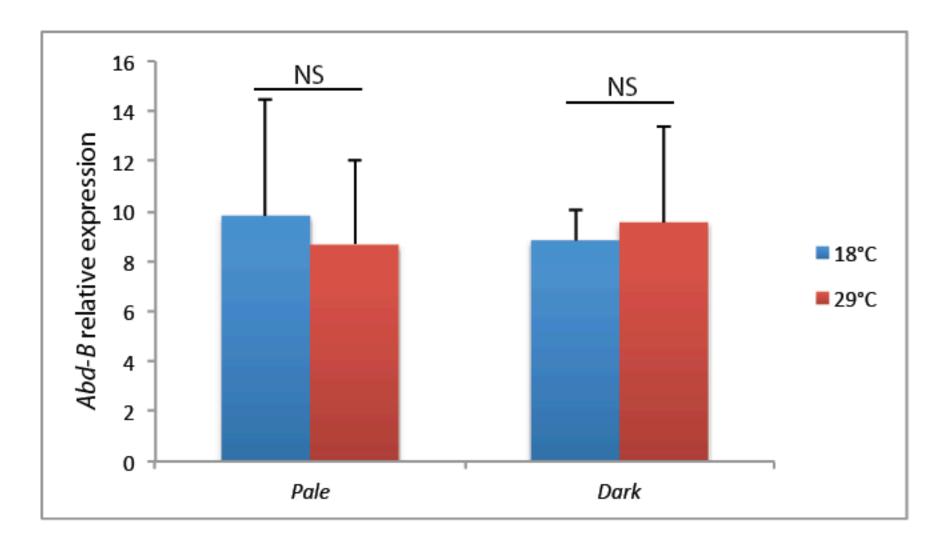
(De Castro et al., PLOS Genetics, 2018)

#### Impact of genetic variation in the enhancer on its activation by AbdB

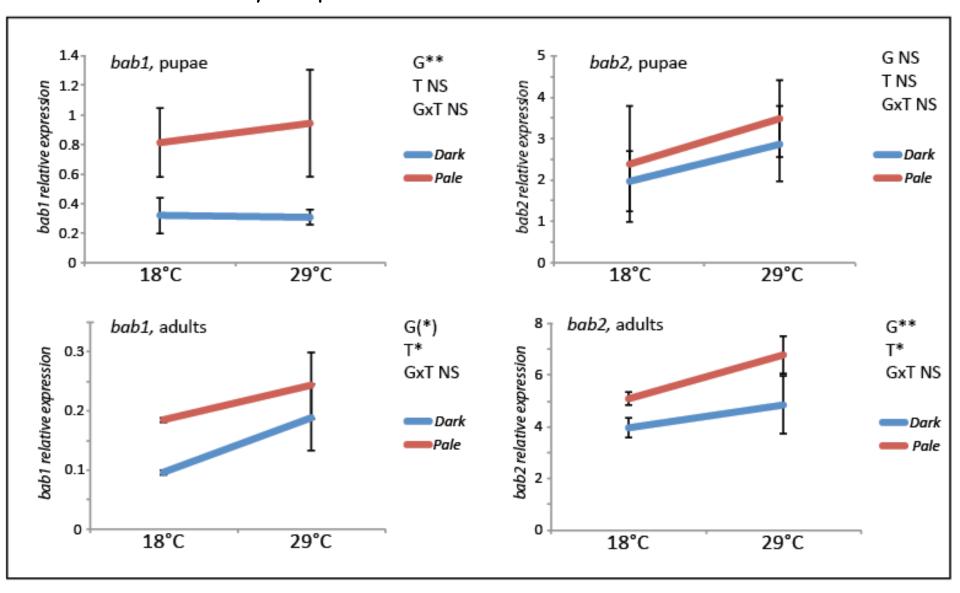


(De Castro et al., PLOS Genetics, 2018)

#### AbdB expression is not modulated by temperature

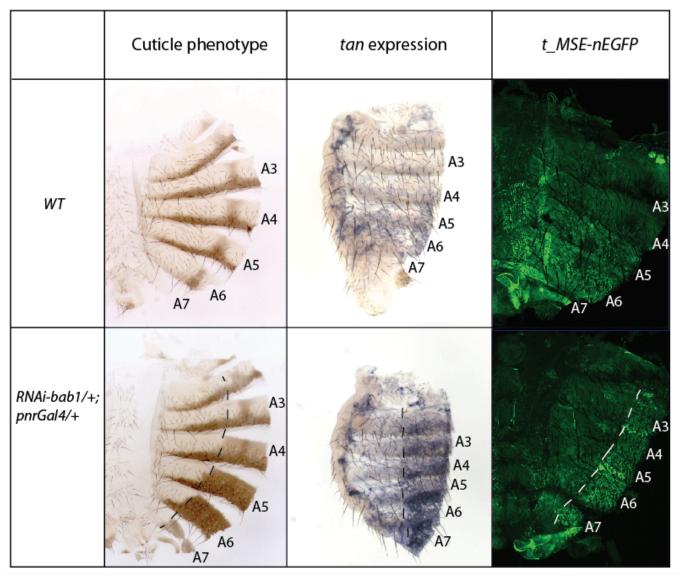


bab1 and bab2 expressions are different between the Dark and Pale lines and modulated by temperature



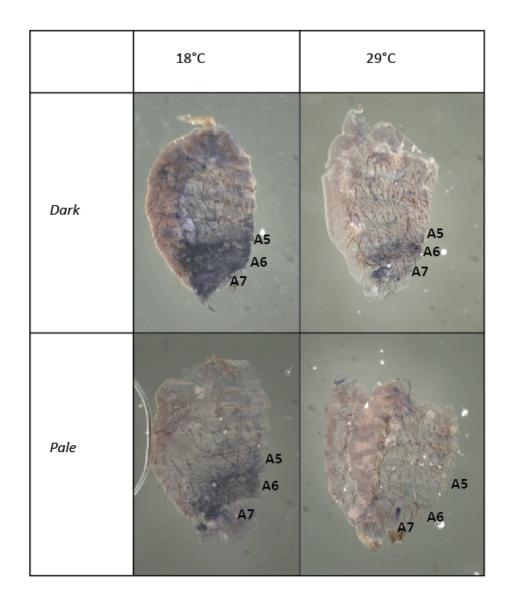
(De Castro et al., PLOS Genetics, 2018)

#### bab represses tan via the t\_MSE

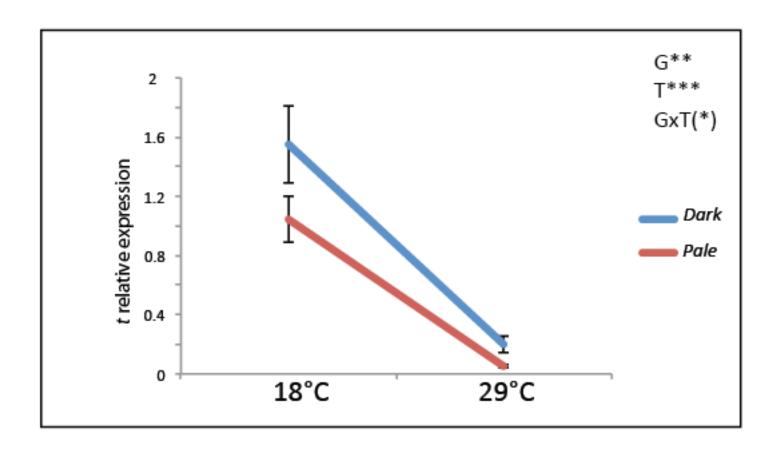


(De Castro et al., PLOS Genetics, 2018)

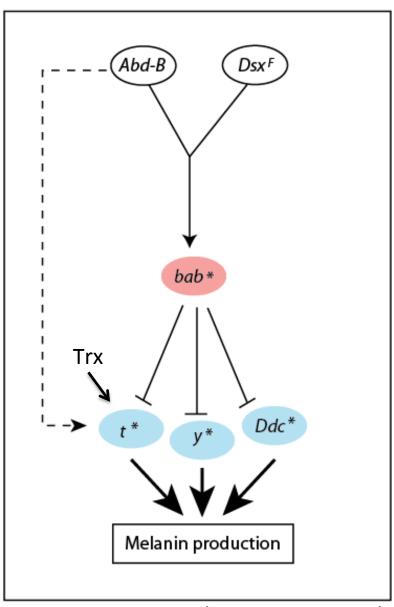
#### Analysis of tan expression in the Dark and Pale lines



#### Analysis of tan expression in the Dark and Pale lines



#### Model



Interestingly, genetic variation in tan t-MSE and bab dimorphic element is involved in within and between *Drosophila* species pigmentation variation (Bastide et al., 2013; Yassin et al., 2016; Jeong et al., 2008; Rogers et al., 2013).

This suggests that the temperature sensitivity of these regulatory sequences turns them into evolutionary hotspots by facilitating the selection of the genetic variation they carry.

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