

Bio393: Genetic Analysis

Developmental genetics I



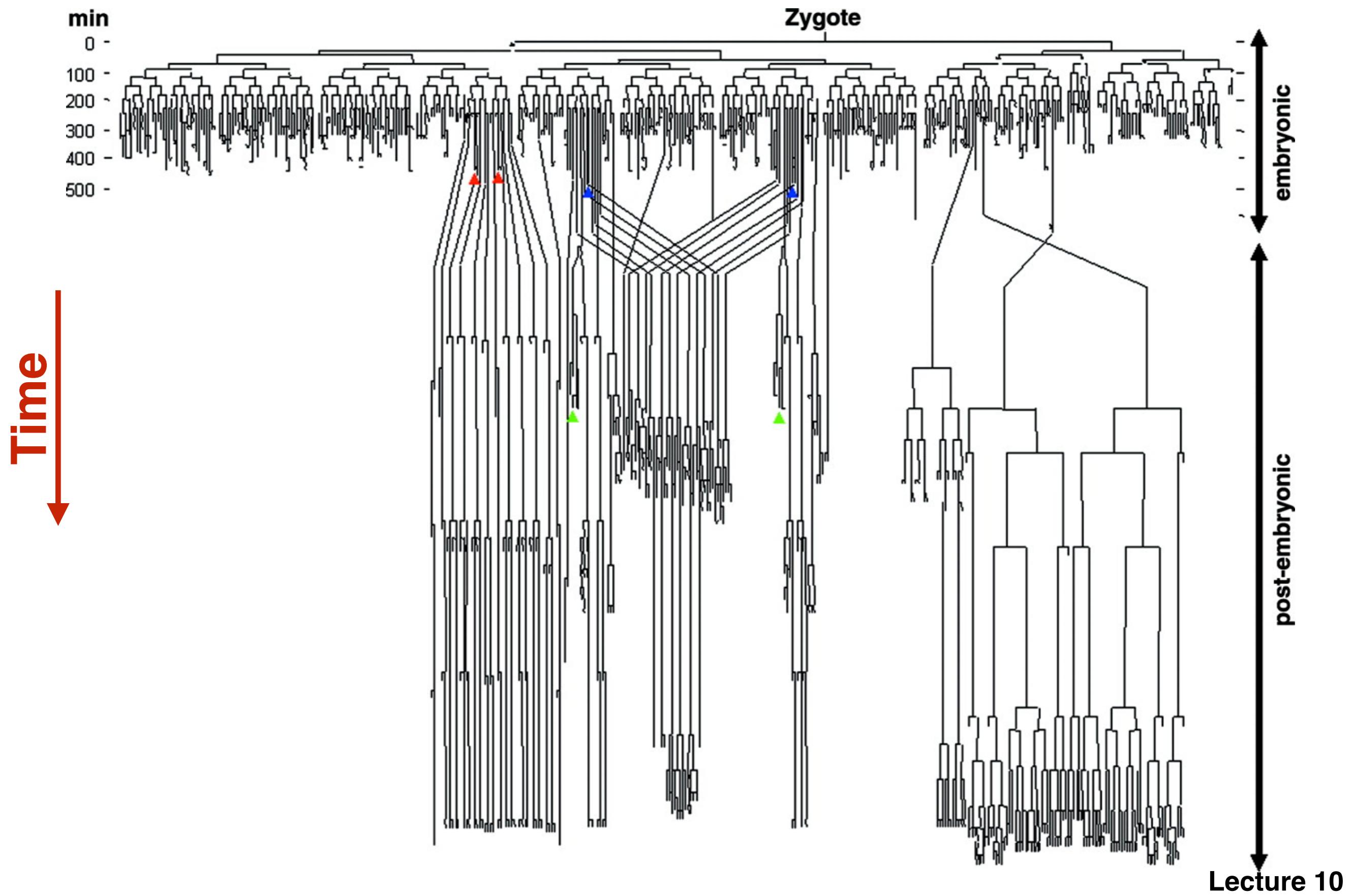
C. elegans

Developmental genetics is the study of how genes regulate the growth and development of an organism.



Cell location
Cell fate
Cell-cell communication
Maternal effects
Cell autonomy
Epistasis

The cell lineage of *C. elegans* is known and invariant

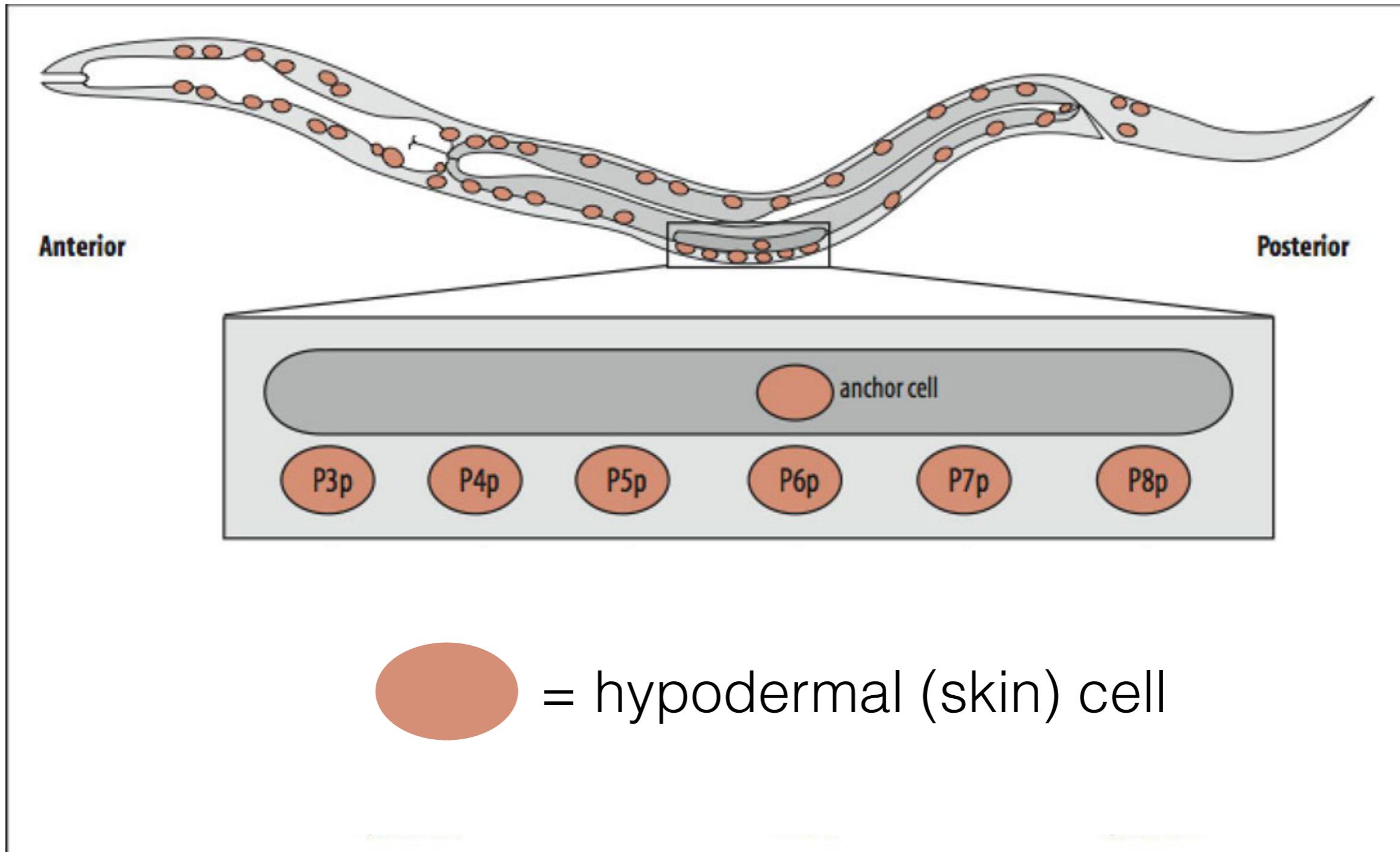


The cell lineage of *C. elegans* is known and invariant



John Sulston

C. elegans vulval development

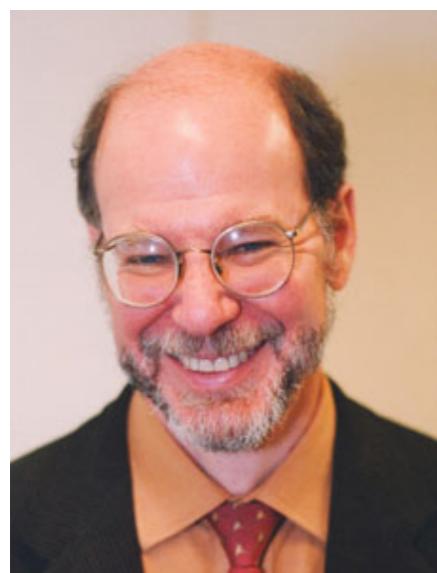
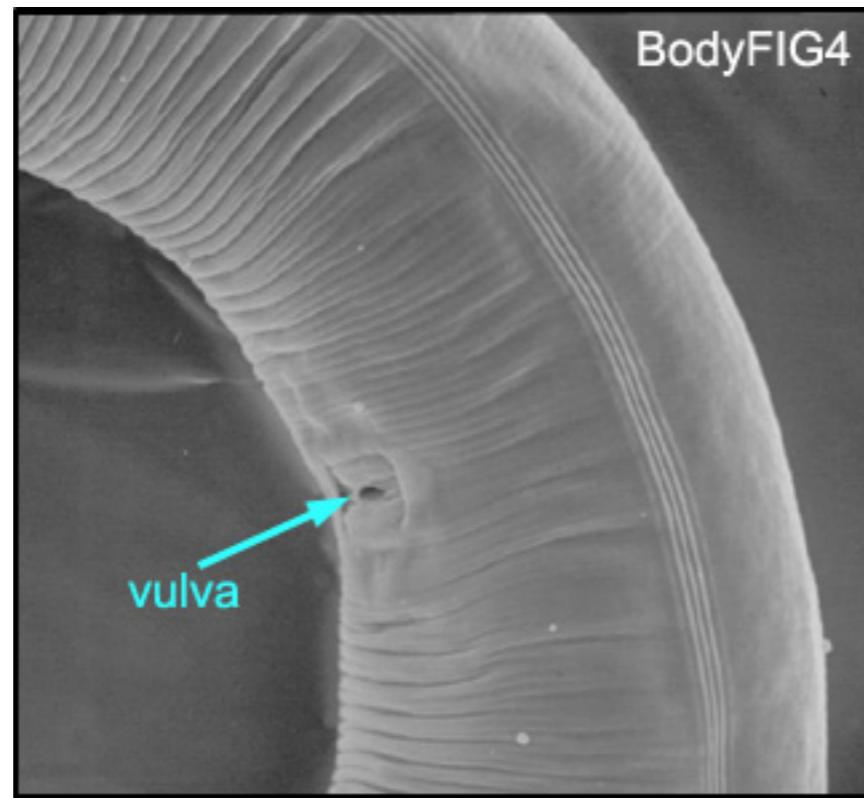


One of the best genetic screens ever

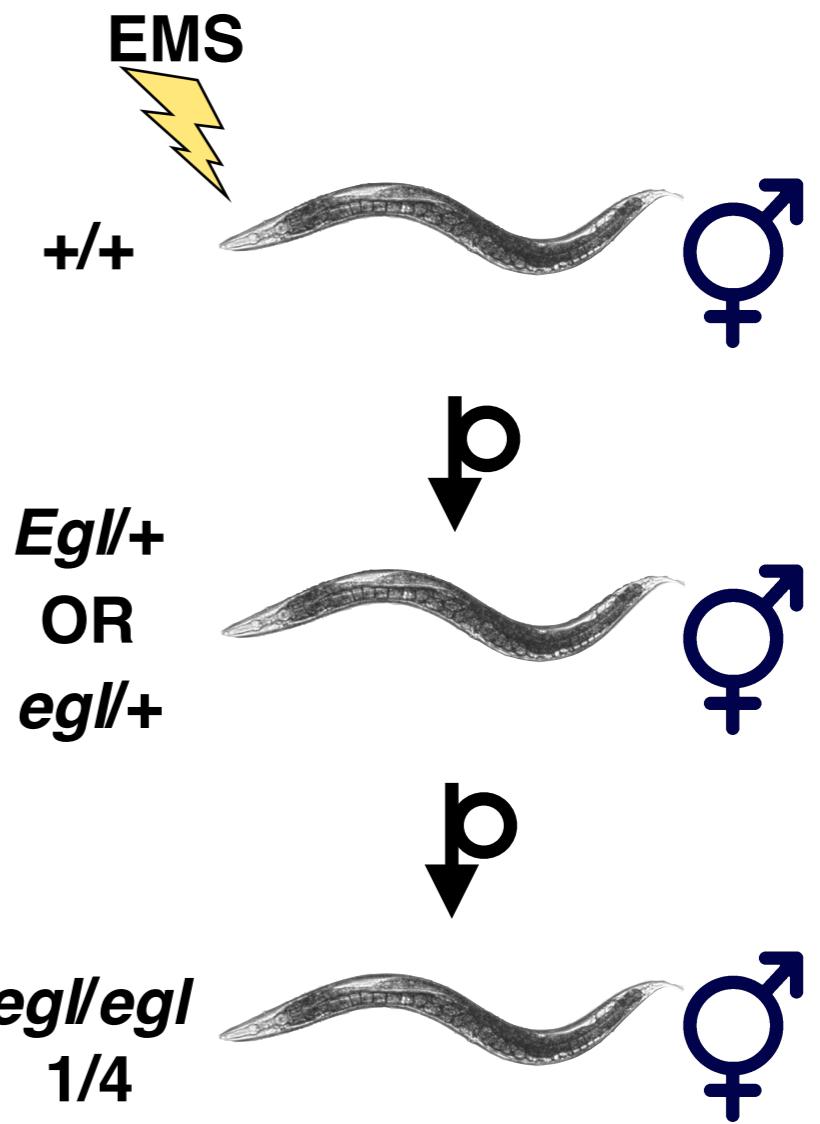


Let's say you screened for mutants that failed to lay eggs

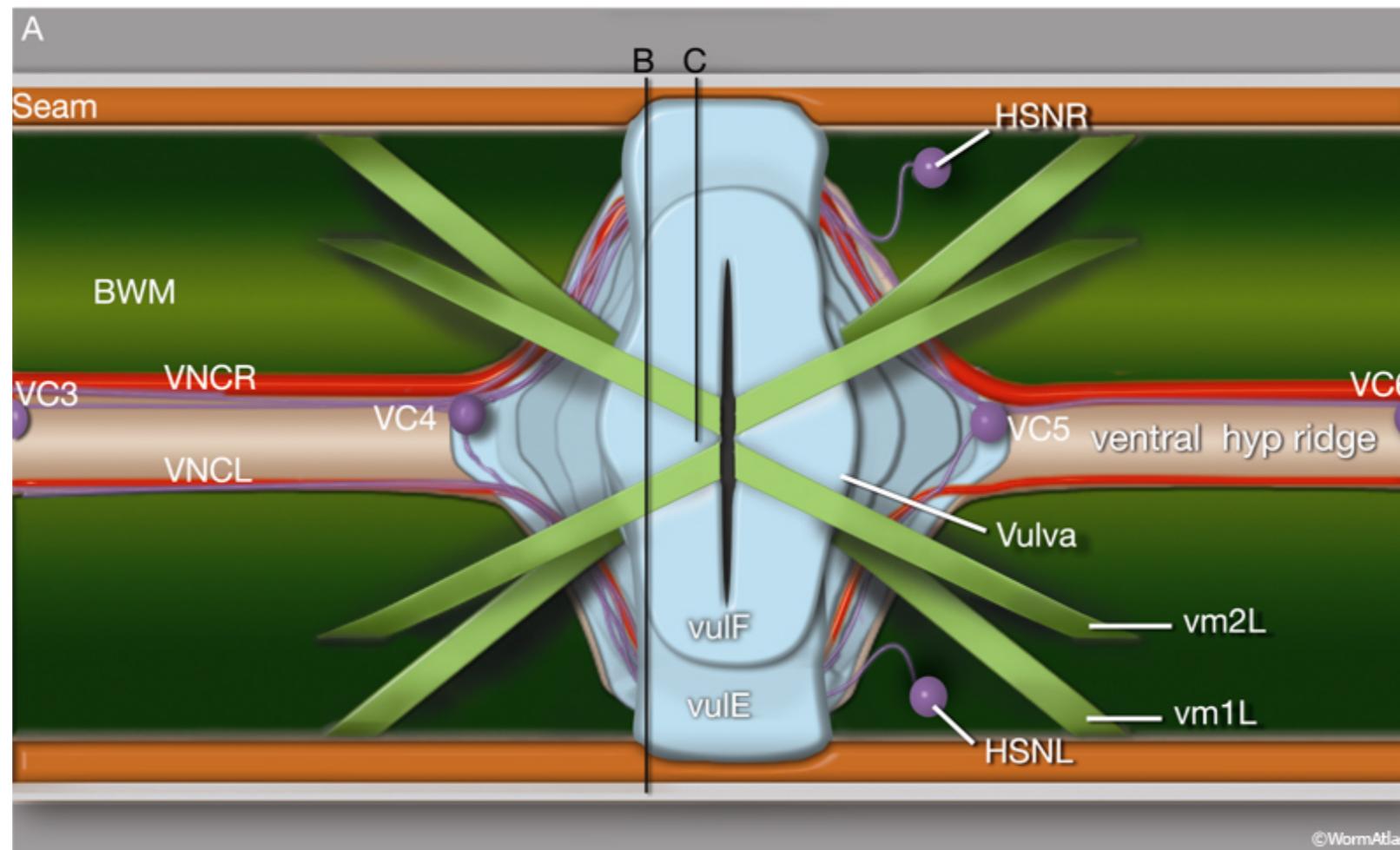
Called Egl for egg-laying defective



Bob Horvitz



One of the best genetic screens ever

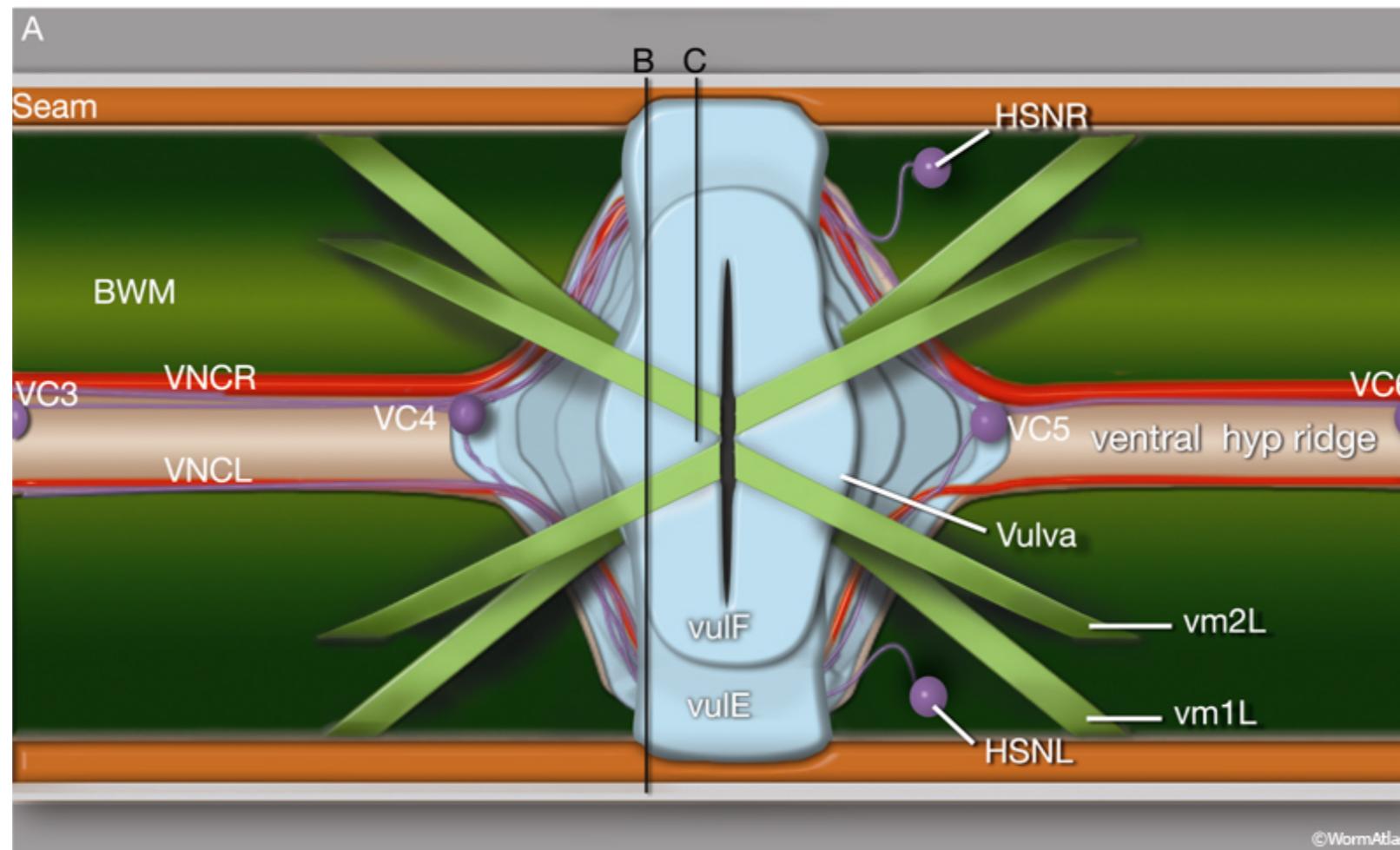


No neuron

egl-1 = inducer of programmed cell death

Hermaphrodite-Specific Neuron (HSN)
inappropriately dies

One of the best genetic screens ever

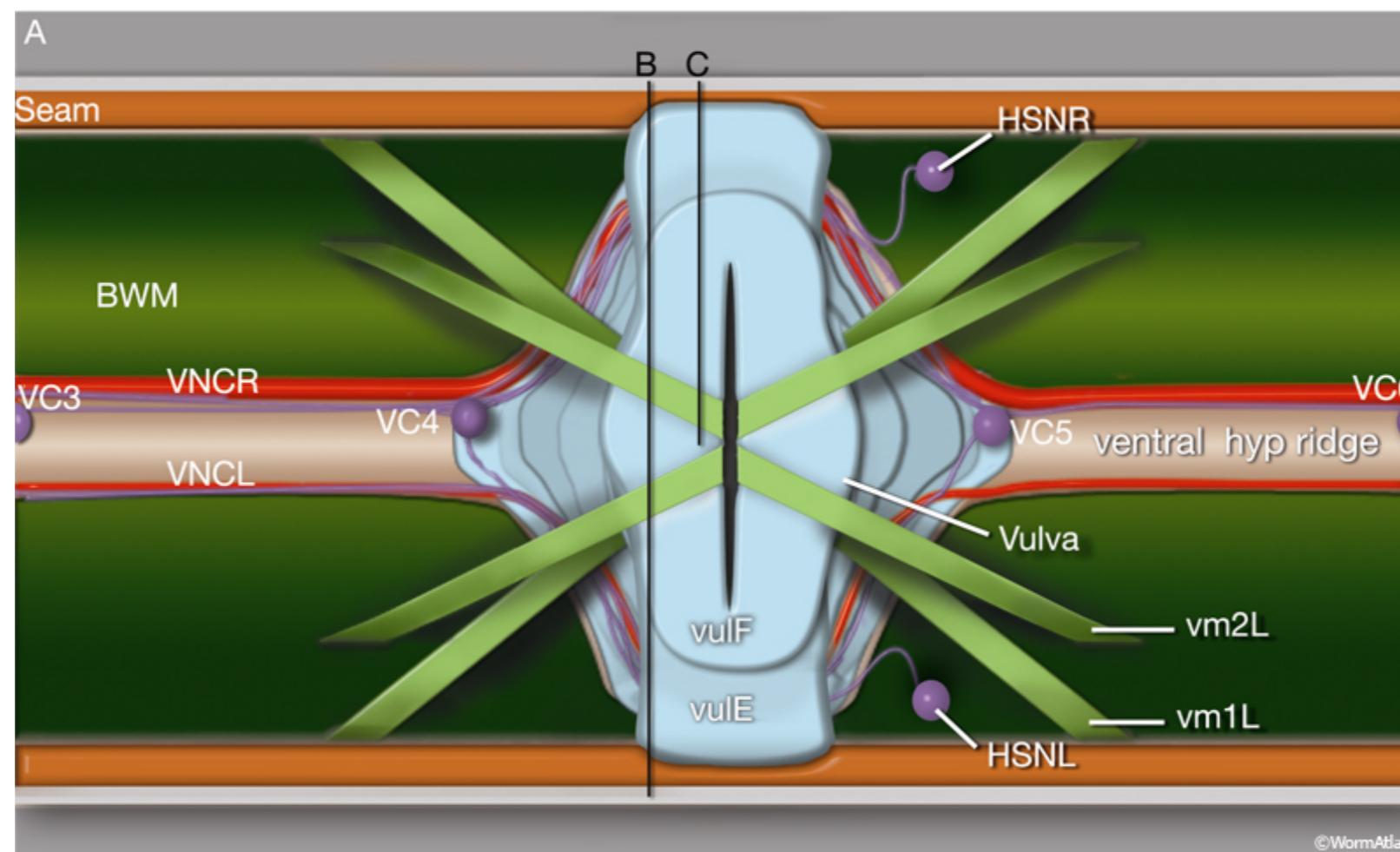


No neuron

tra-1 = inducer of sex determination

Mutants are partially male so HSNs die

One of the best genetic screens ever

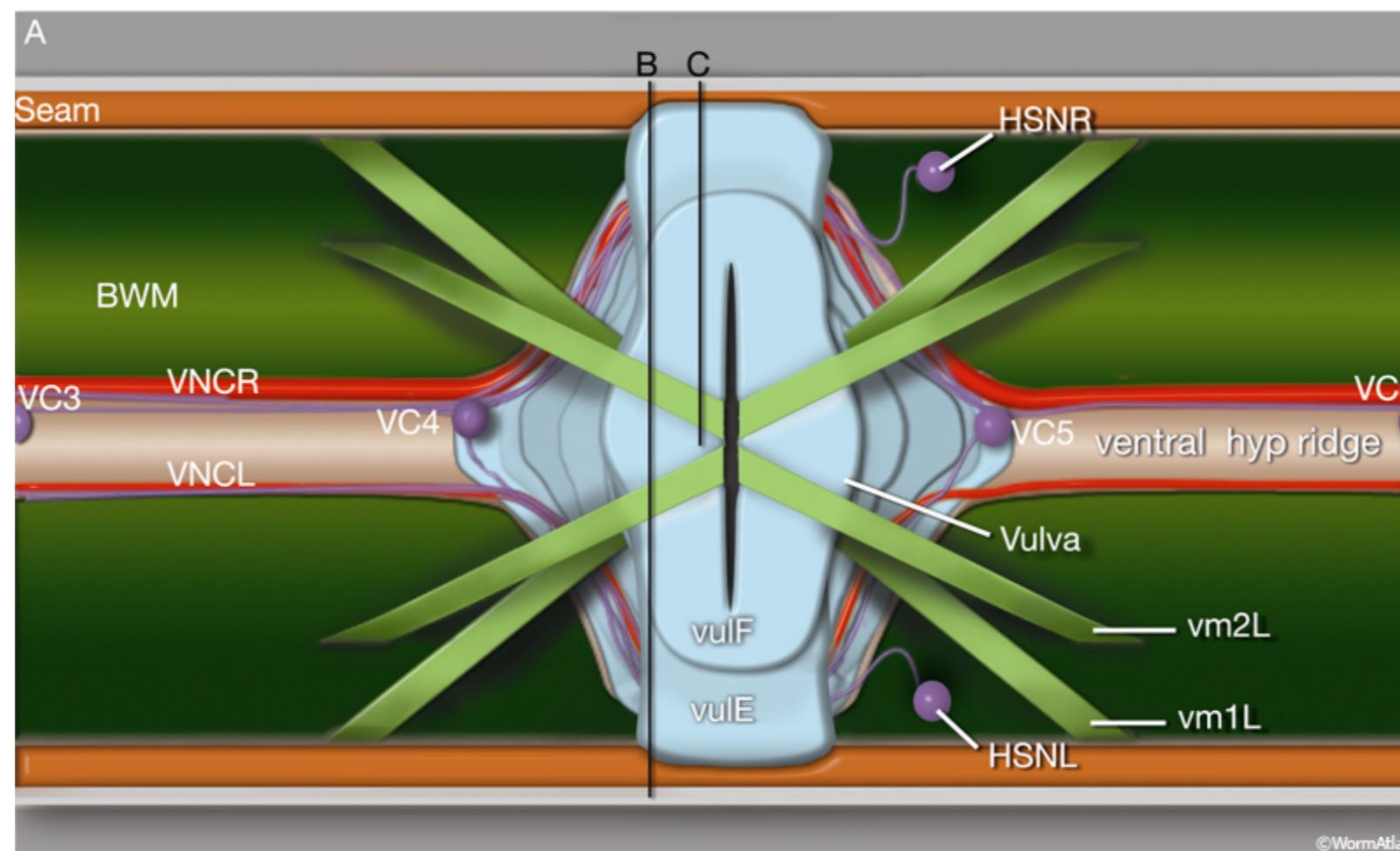


No neuron

No neuronal
signaling

egl-6 = serotonin signaling from HSN is defective

One of the best genetic screens ever



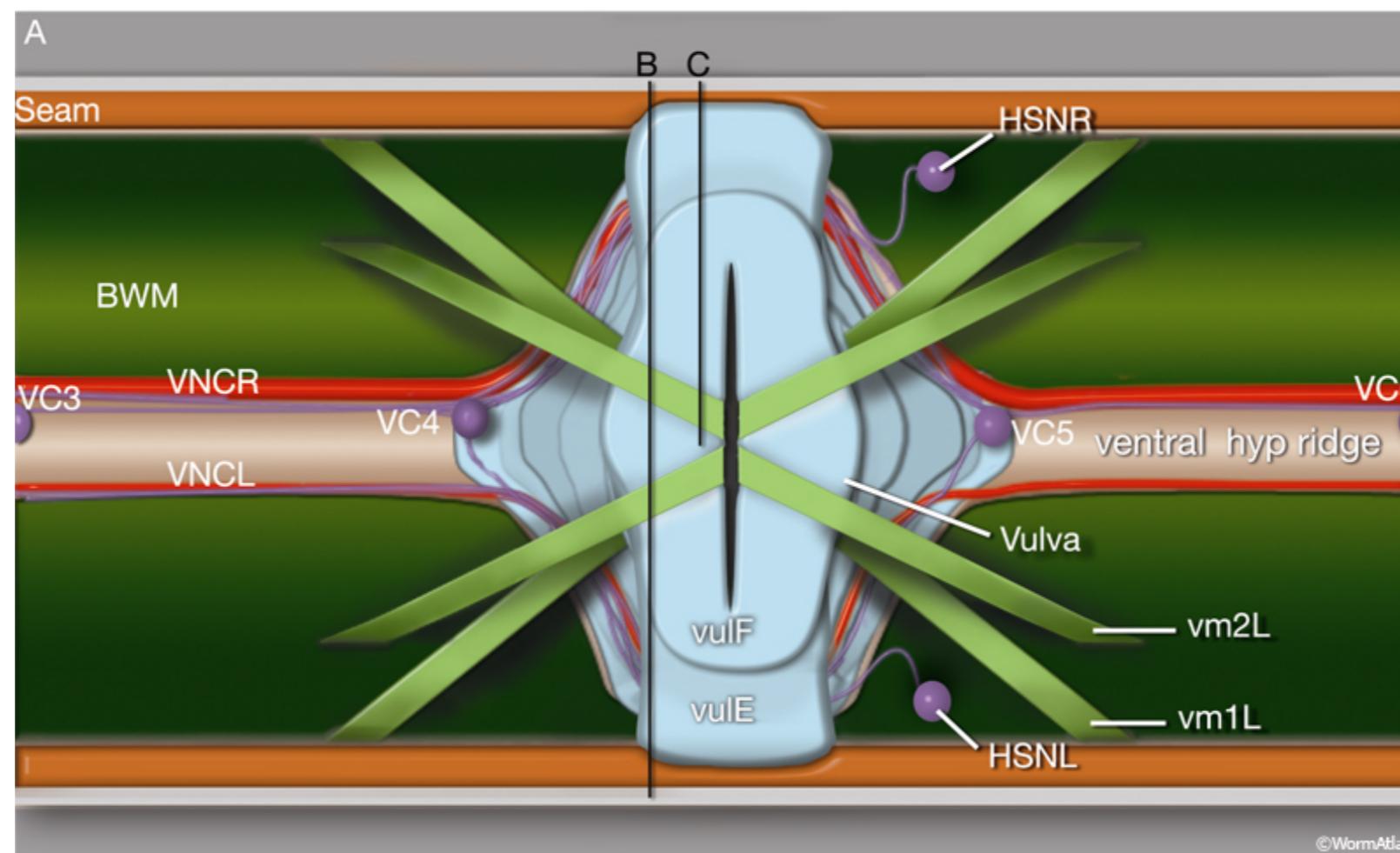
No neuron
No neuronal
signaling

No vulva

lin-3 = lineage defective gene 3

Vulval cells are not specified

One of the best genetic screens ever



No neuron

No neuronal
signaling

No vulva

Vulval muscle fails to function

Vulval opening fails to form

Uterus doesn't connect to vulva

One of the best genetic screens ever



C. elegans



Sydney Brenner



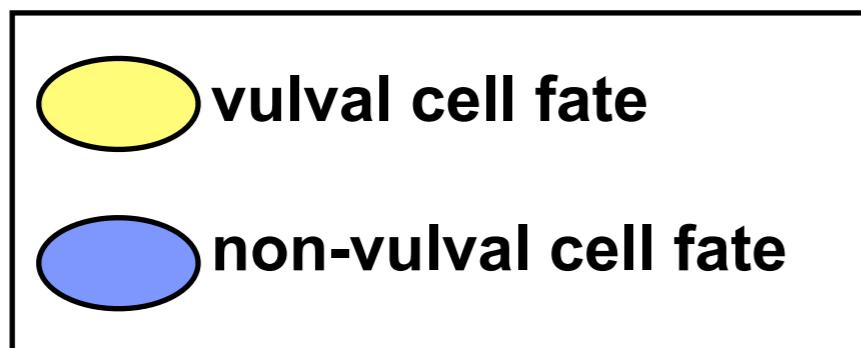
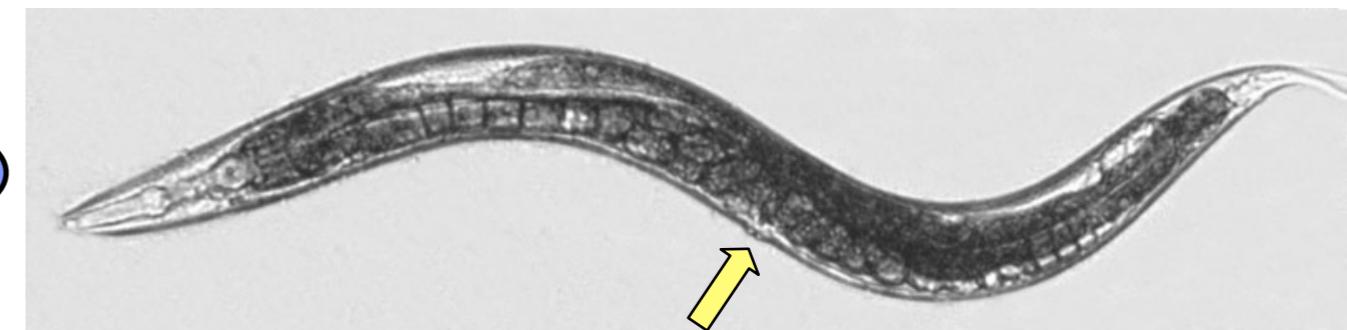
H. Robert Horvitz



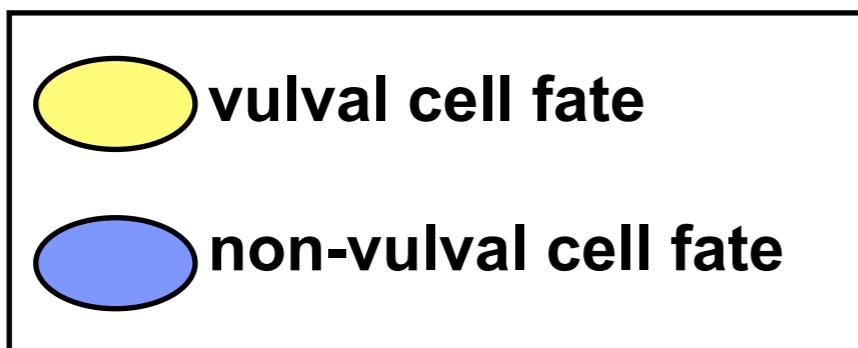
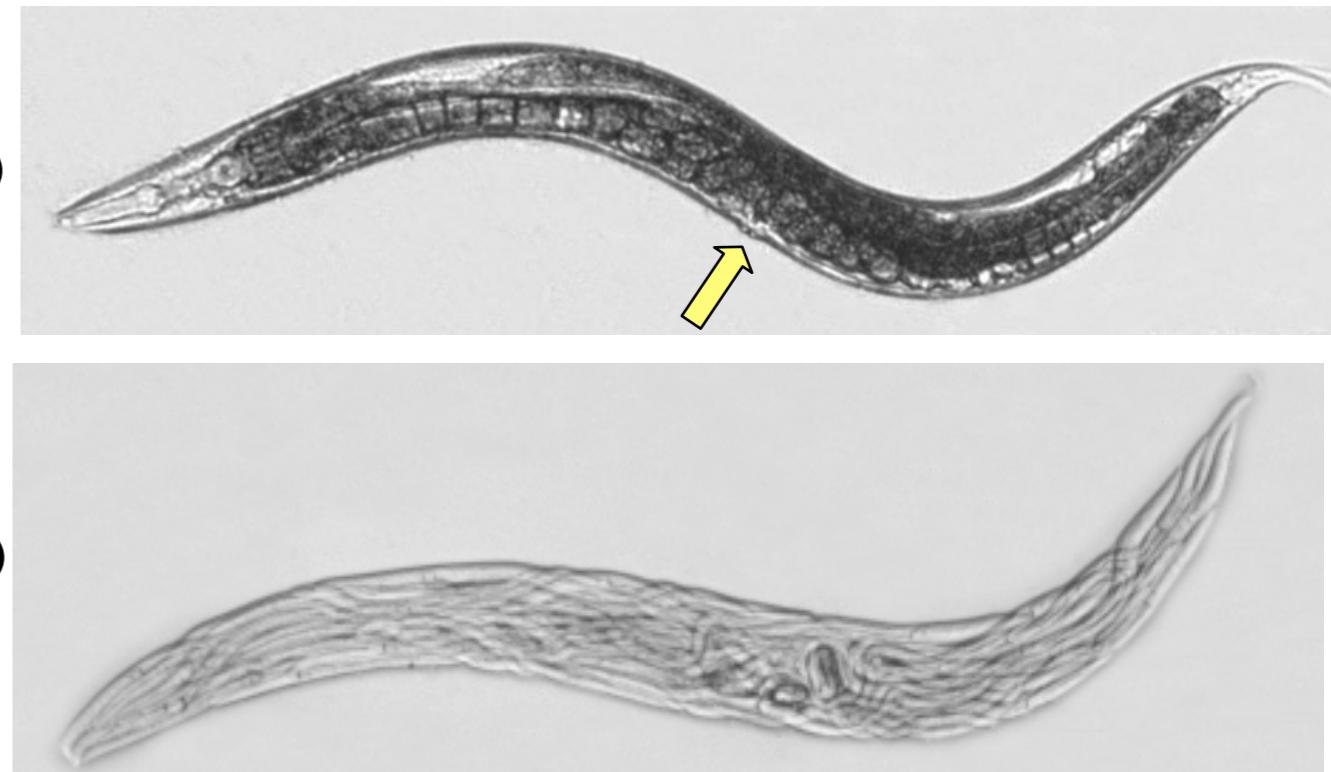
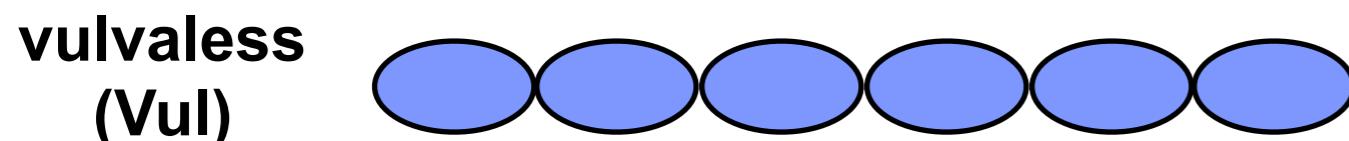
John E. Sulston



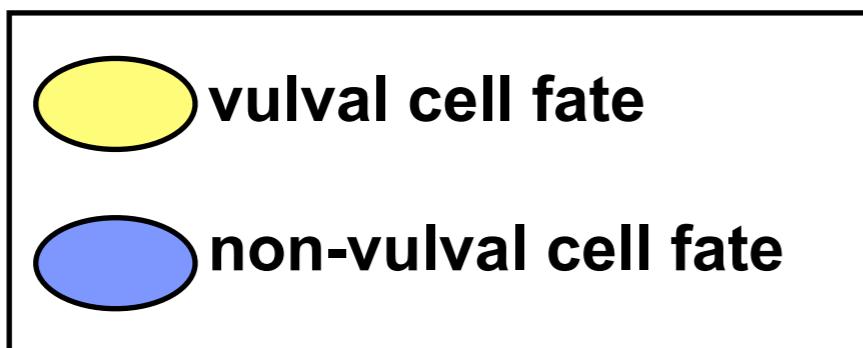
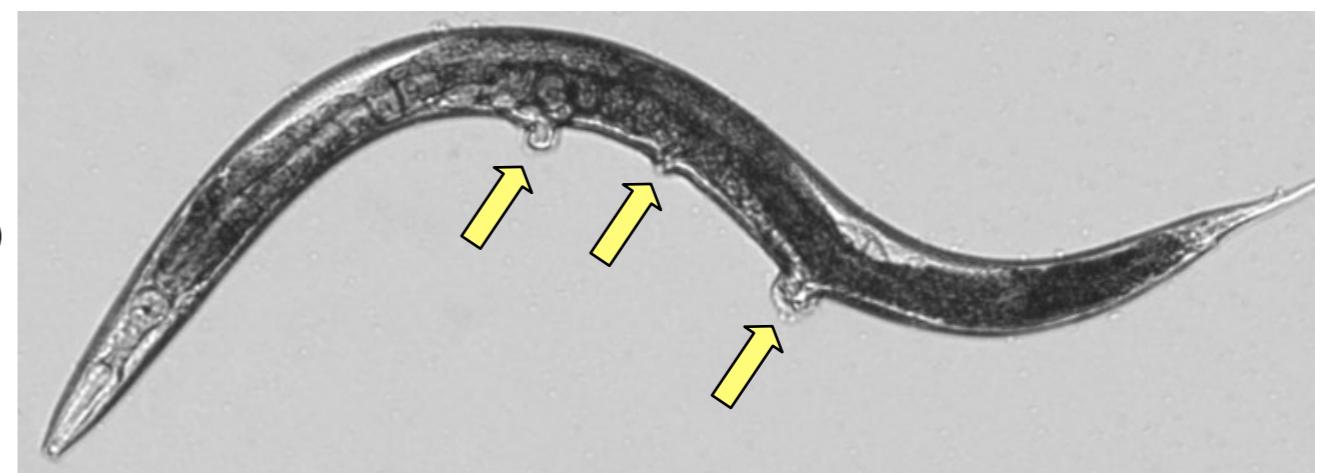
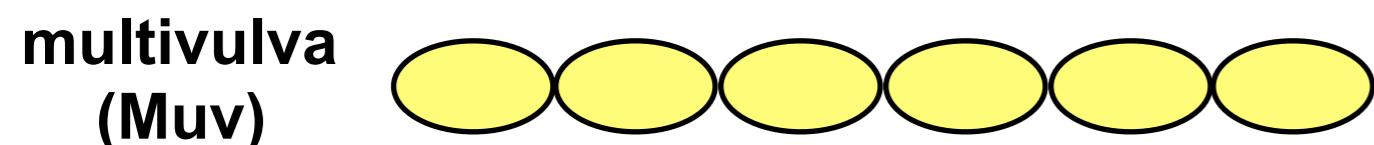
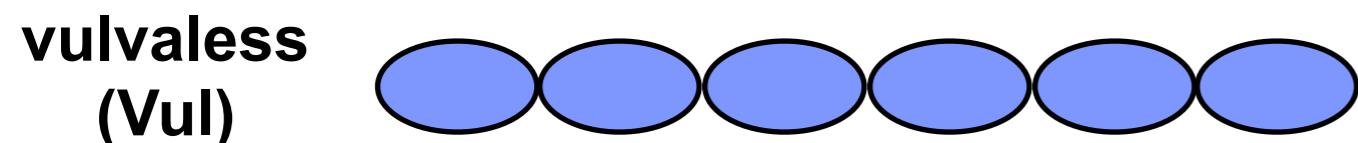
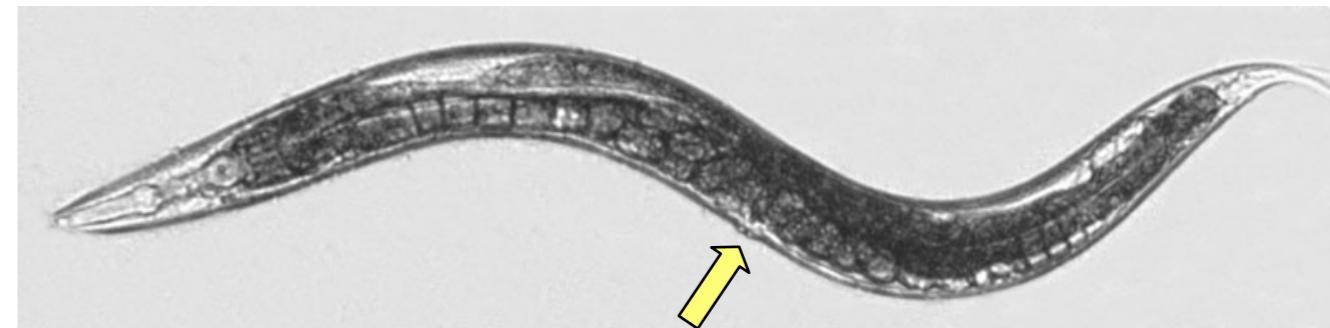
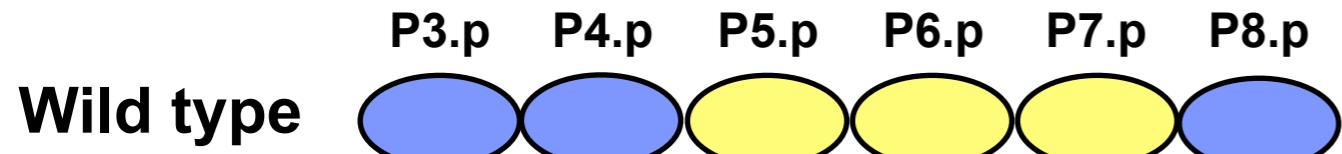
Three cells express vulval fates in wild-type animals



No cells express vulval fates in vulvaless mutants



Six cells express vulval fates in multivulva mutants



Vulval mutants

Mutant	Phenotype
<i>lin-1(0)</i>	Muv
<i>lin-3(0)</i>	Vul
<i>let-60(0)</i>	Vul
<i>let-60(gf)</i>	Muv
<i>let-23(0)</i>	Vul
<i>let-23(gf)</i>	Muv
<i>lin-39(0)</i>	Vul

Double mutants defined the vulval pathway

Mutant genotypes	Phenotype
<i>lin-1(0)</i>	Muv
<i>lin-3(0)</i>	Vul
<i>let-60(0)</i>	Vul
<i>let-60(gf)</i>	Muv
<i>let-23(0)</i>	Vul
<i>let-23(gf)</i>	Muv
<i>lin-39(0)</i>	Vul
<i>lin-3(0); let-23(gf)</i>	Muv
<i>lin-3(0); let-60(gf)</i>	Muv
<i>let-23(0); let-60(gf)</i>	Muv
<i>let-23(gf); let-60(0)</i>	Vul

***lin-3* → vulval fate**

***let-60* → vulval fate**

***let-23* → vulval fate**

Double mutants defined the vulval pathway

Mutant genotypes	Phenotype
<i>lin-1(0)</i>	Muv
<i>lin-3(0)</i>	Vul
<i>let-60(0)</i>	Vul
<i>let-60(gf)</i>	Muv
<i>let-23(0)</i>	Vul
<i>let-23(gf)</i>	Muv
<i>lin-39(0)</i>	Vul
<i>lin-3(0); let-23(gf)</i>	Muv
<i>lin-3(0); let-60(gf)</i>	Muv
<i>let-23(0); let-60(gf)</i>	Muv
<i>let-23(gf); let-60(0)</i>	Vul

***lin-3* → vulval fate**

***let-60* → vulval fate**

***let-23* → vulval fate**

***lin-3* → *let-23* → vulval fate**

Double mutants defined the vulval pathway

Mutant genotypes	Phenotype
<i>lin-1(0)</i>	Muv
<i>lin-3(0)</i>	Vul
<i>let-60(0)</i>	Vul
<i>let-60(gf)</i>	Muv
<i>let-23(0)</i>	Vul
<i>let-23(gf)</i>	Muv
<i>lin-39(0)</i>	Vul
<i>lin-3(0); let-23(gf)</i>	Muv
<i>lin-3(0); let-60(gf)</i>	Muv
<i>let-23(0); let-60(gf)</i>	Muv
<i>let-23(gf); let-60(0)</i>	Vul

***lin-3* → vulval fate**

***let-60* → vulval fate**

***let-23* → vulval fate**

***lin-3* → *let-23* → *let-60* → vulval fate**

Double mutants defined the vulval pathway

Mutant genotypes	Phenotype
<i>lin-1(0)</i>	Muv
<i>lin-3(0)</i>	Vul
<i>let-60(0)</i>	Vul
<i>let-60(gf)</i>	Muv
<i>let-23(0)</i>	Vul
<i>let-23(gf)</i>	Muv
<i>lin-39(0)</i>	Vul
<i>lin-3(0); let-23(gf)</i>	Muv
<i>lin-3(0); let-60(gf)</i>	Muv
<i>let-23(0); let-60(gf)</i>	Muv
<i>let-23(gf); let-60(0)</i>	Vul
<i>let-60(0); lin-1(0)</i>	Muv
<i>let-23(0); lin-1(0)</i>	Muv

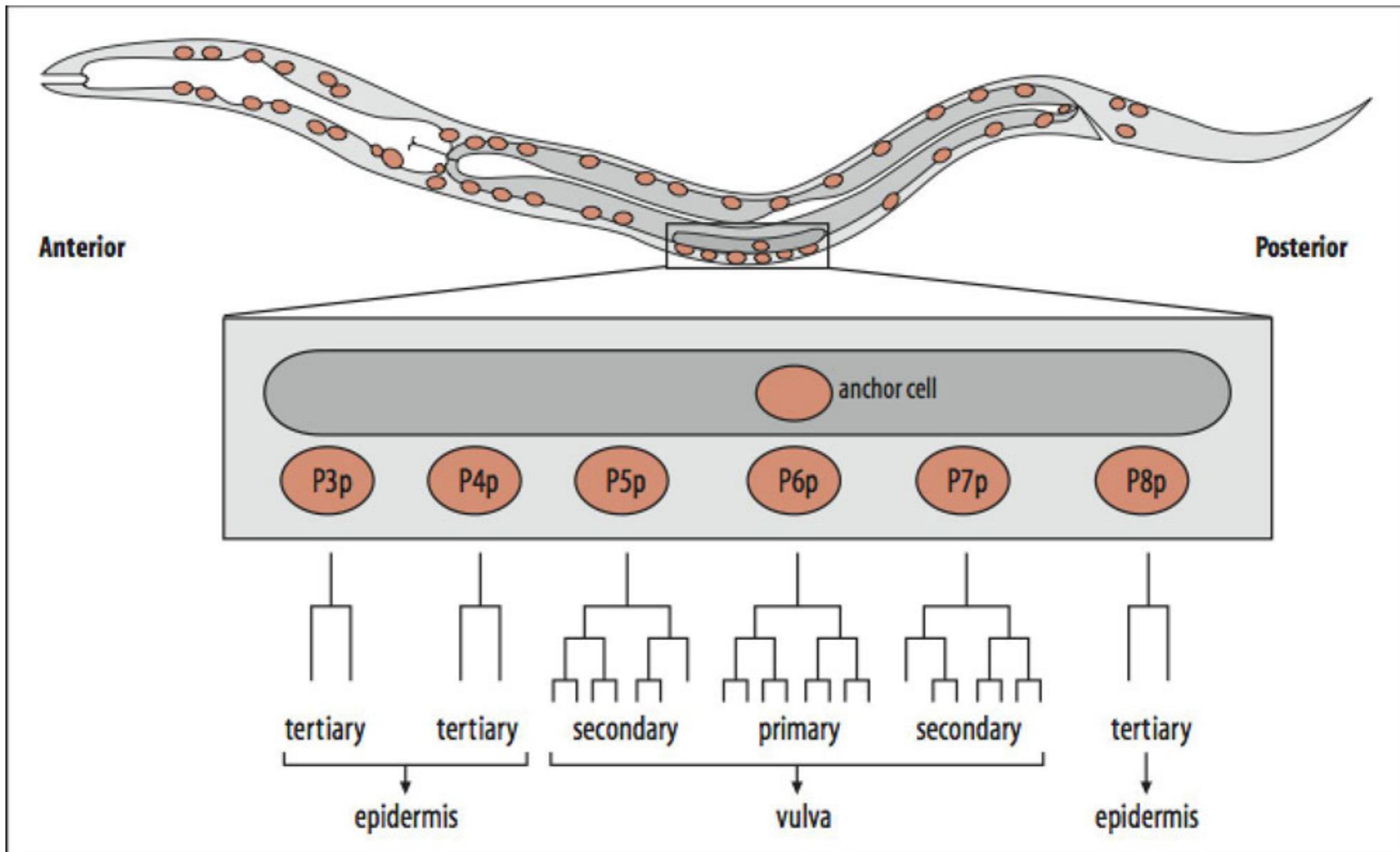
***lin-3* → *let-23* → *let-60* → *lin-1* → vulval fate**

Double mutants defined the vulval pathway

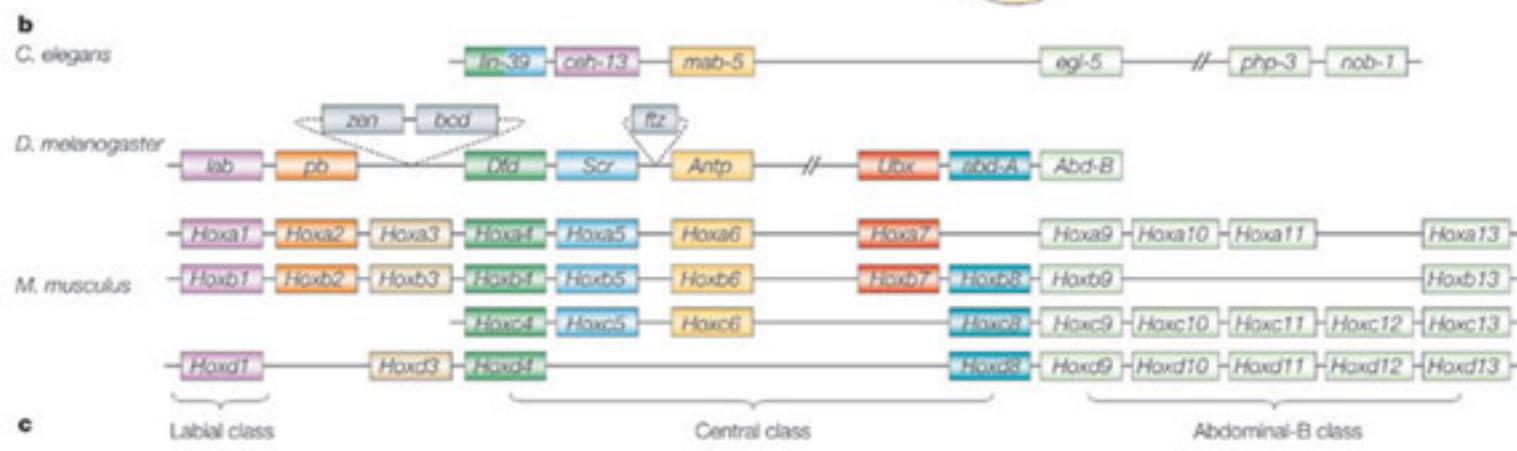
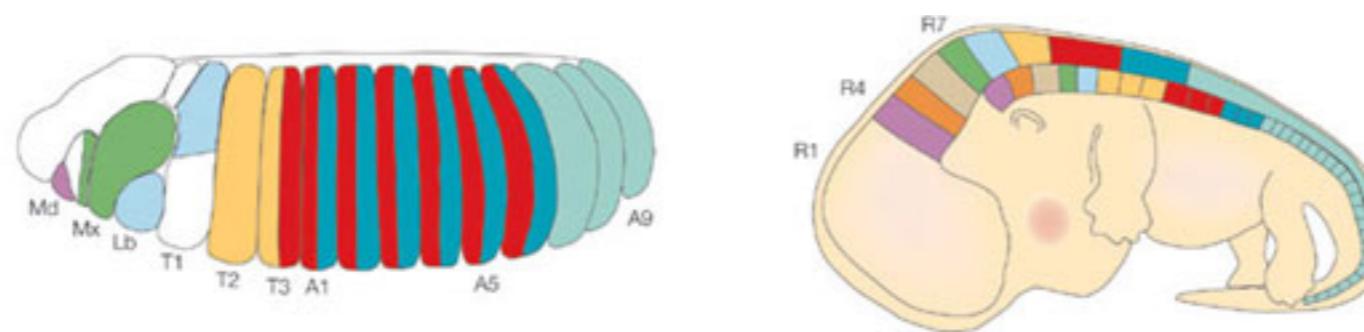
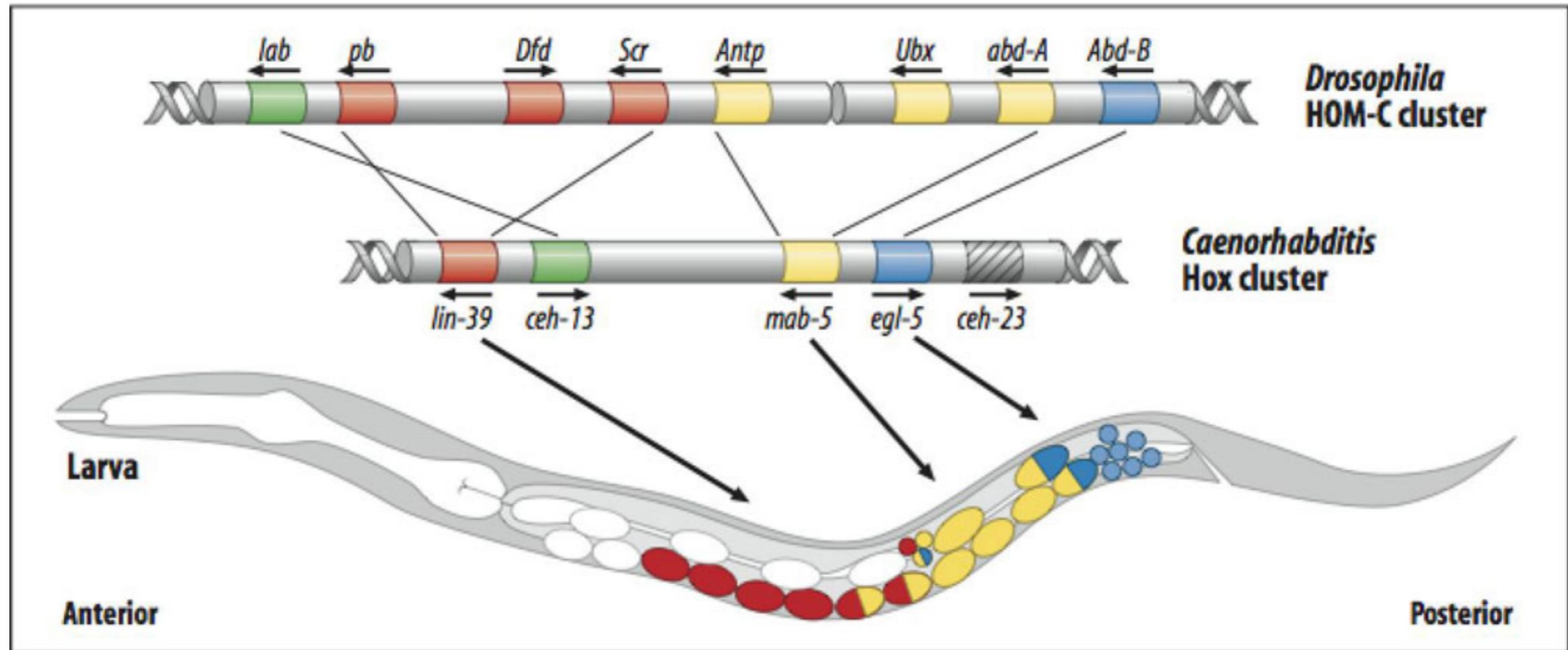
Mutant genotypes	Phenotype
<i>lin-1(0)</i>	Muv
<i>lin-3(0)</i>	Vul
<i>let-60(0)</i>	Vul
<i>let-60(gf)</i>	Muv
<i>let-23(0)</i>	Vul
<i>let-23(gf)</i>	Muv
<i>lin-39(0)</i>	Vul
<i>lin-3(0); let-23(gf)</i>	Muv
<i>lin-3(0); let-60(gf)</i>	Muv
<i>let-23(0); let-60(gf)</i>	Muv
<i>let-23(gf); let-60(0)</i>	Vul
<i>let-60(0); lin-1(0)</i>	Muv
<i>let-23(0); lin-1(0)</i>	Muv
<i>lin-1(0); lin-39(0)</i>	Vul

***lin-3* → *let-23* → *let-60* → *lin-1* → *lin-39* → vulval fate**

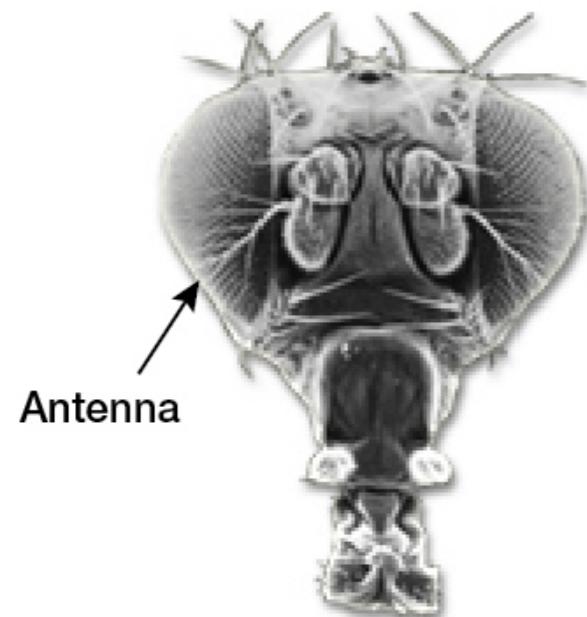
Careful observation of *lin-39(0)* mutants showed that P3p through P8p were not made correctly



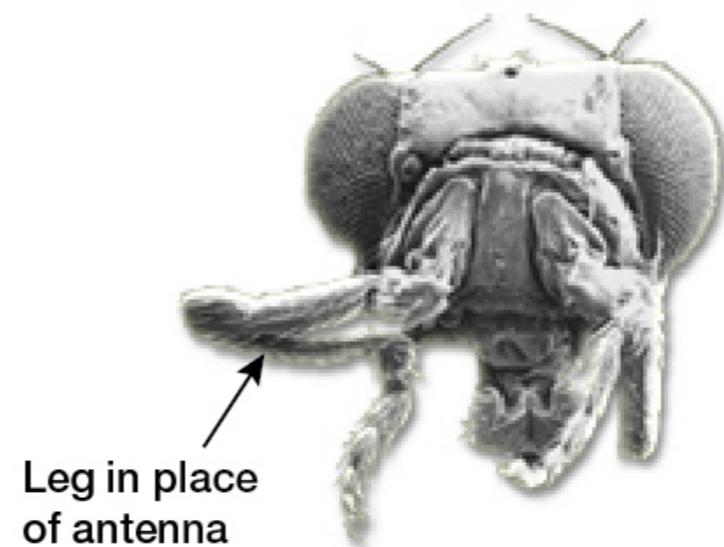
lin-39 is a Hox gene



Regional identity in the animal is deficient in Hox mutants

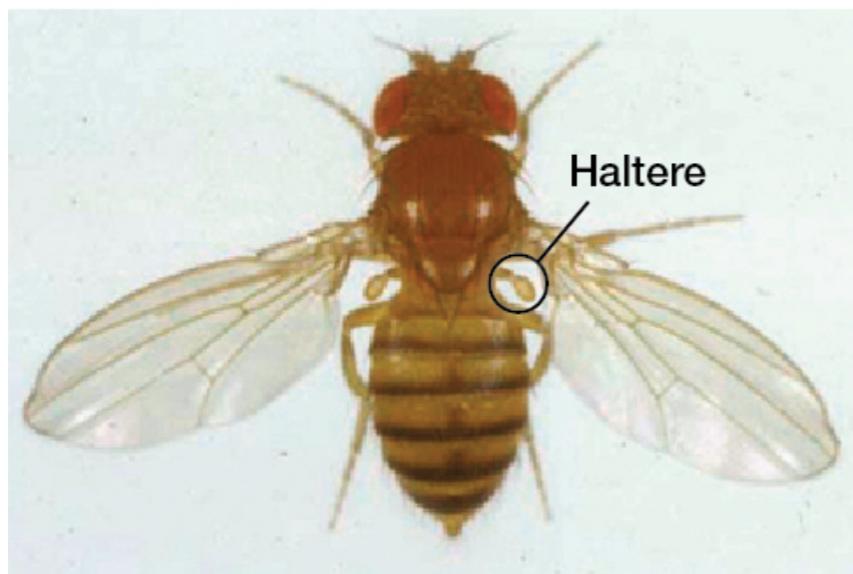


Antenna



Leg in place
of antenna

Antenna to leg

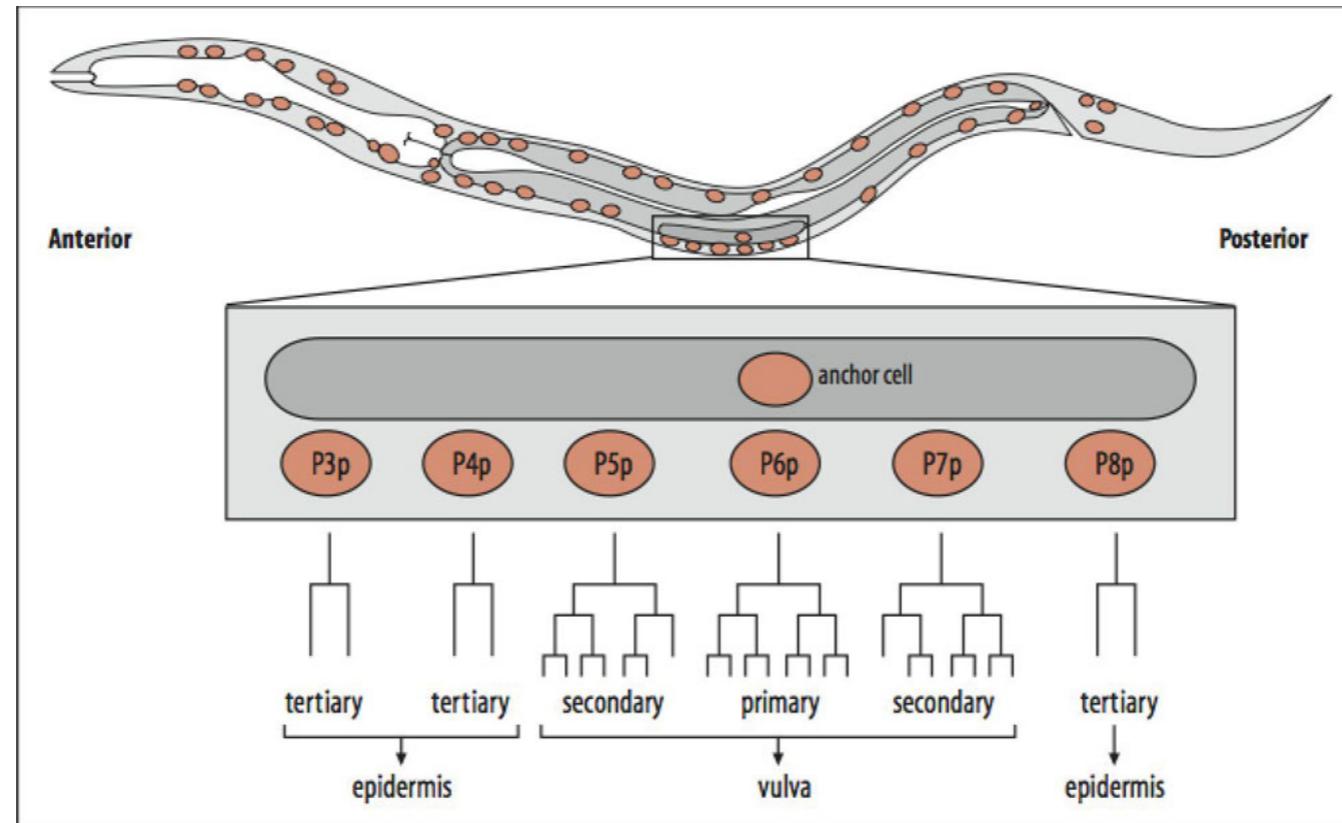


Haltere



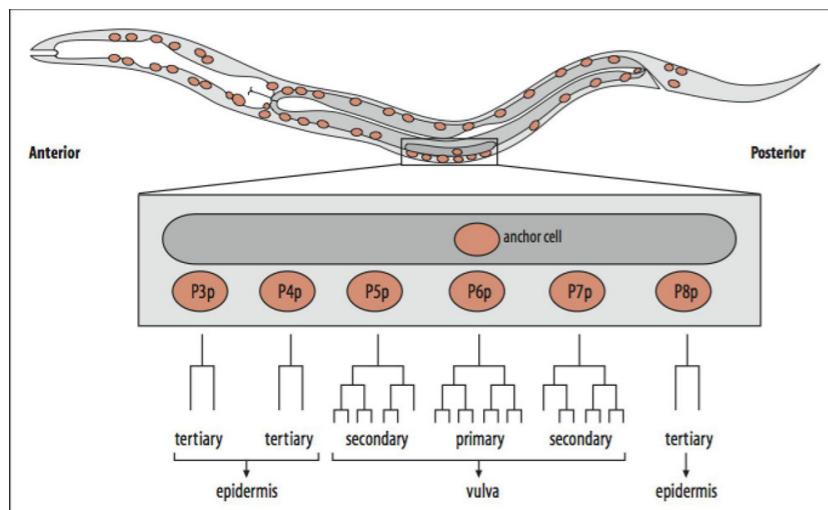
Haltere to wing

lin-39 is a Hox gene



Generation of the vulval precursor cells is epistatic to the fates of those cells

lin-39 is a Hox gene

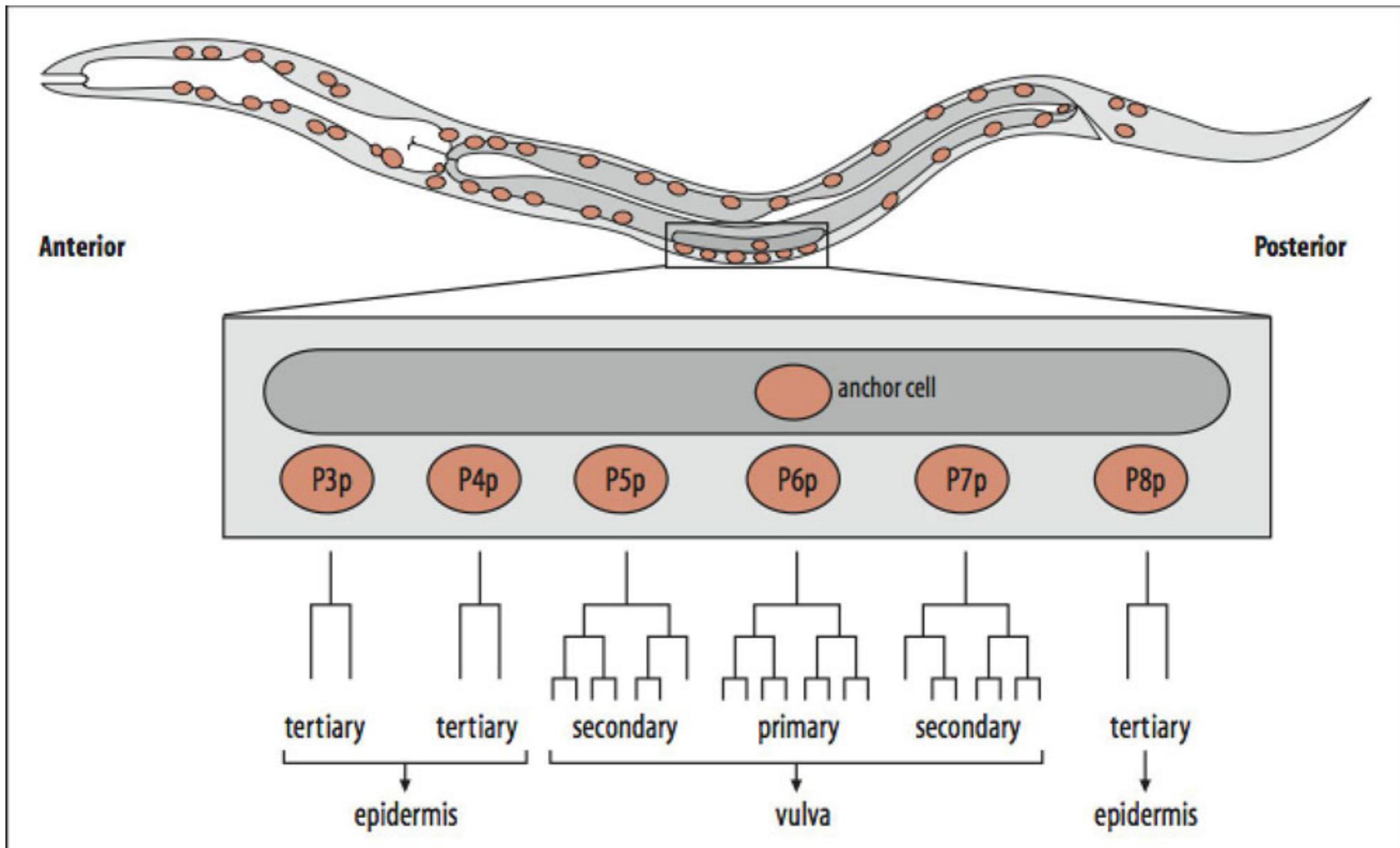


Generation of the vulval precursor cells is epistatic to the fates of those cells

lin-39 → *lin-3* → *let-23* → *let-60* → *lin-1* → vulval fate

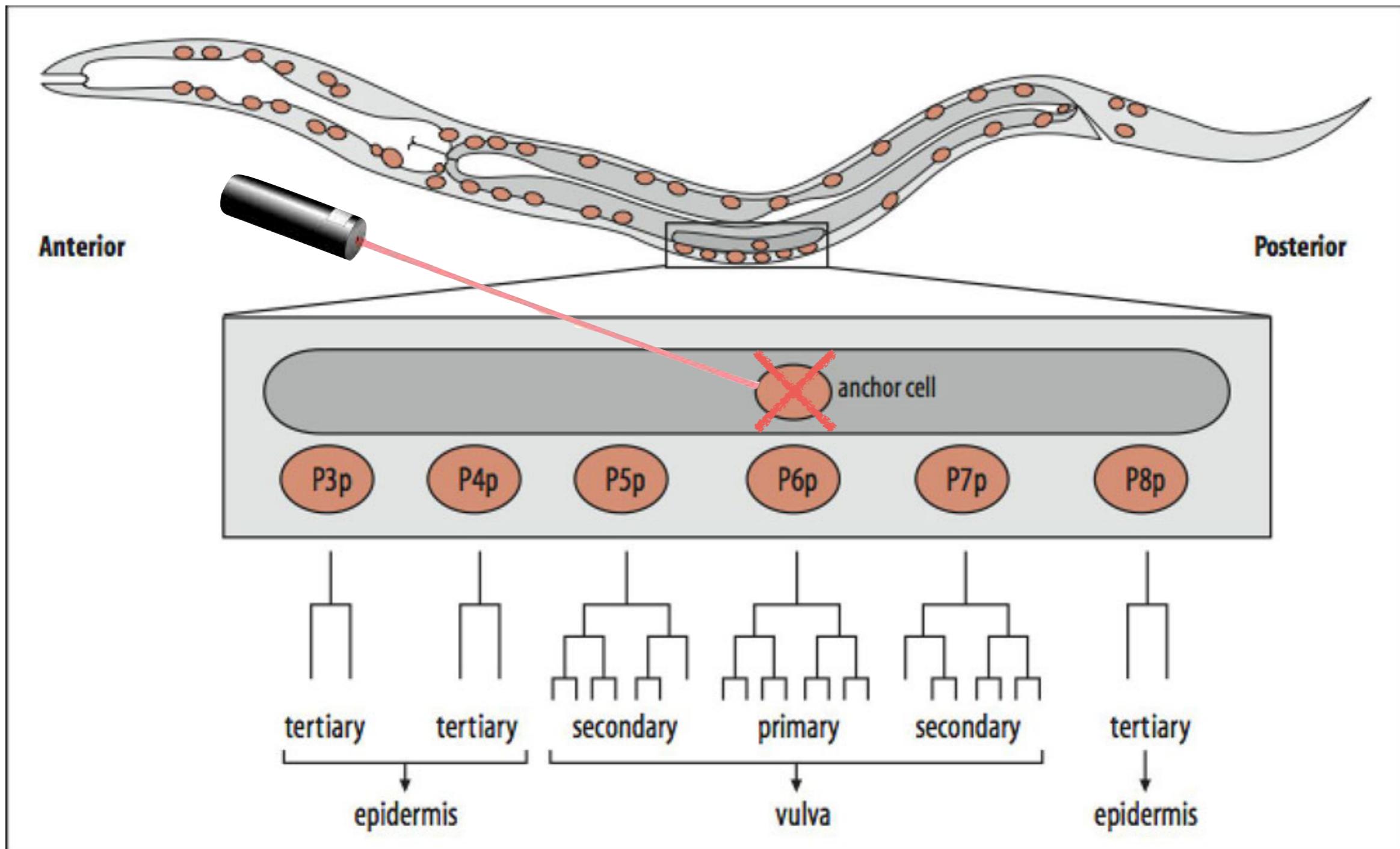
Mutant genotypes	Phenotype
<i>lin-1(0)</i>	Muv
<i>lin-3(0)</i>	Vul
<i>lin-39(0)</i>	Vul
<i>lin-1(0); lin-3(0)</i>	Muv
<i>lin-1(0); lin-39(0)</i>	Vul

What is the source of the inductive signal?



How do we determine that the AC is necessary and sufficient for vulval development (primary or secondary cells)?

Ablation removes cells (necessary)

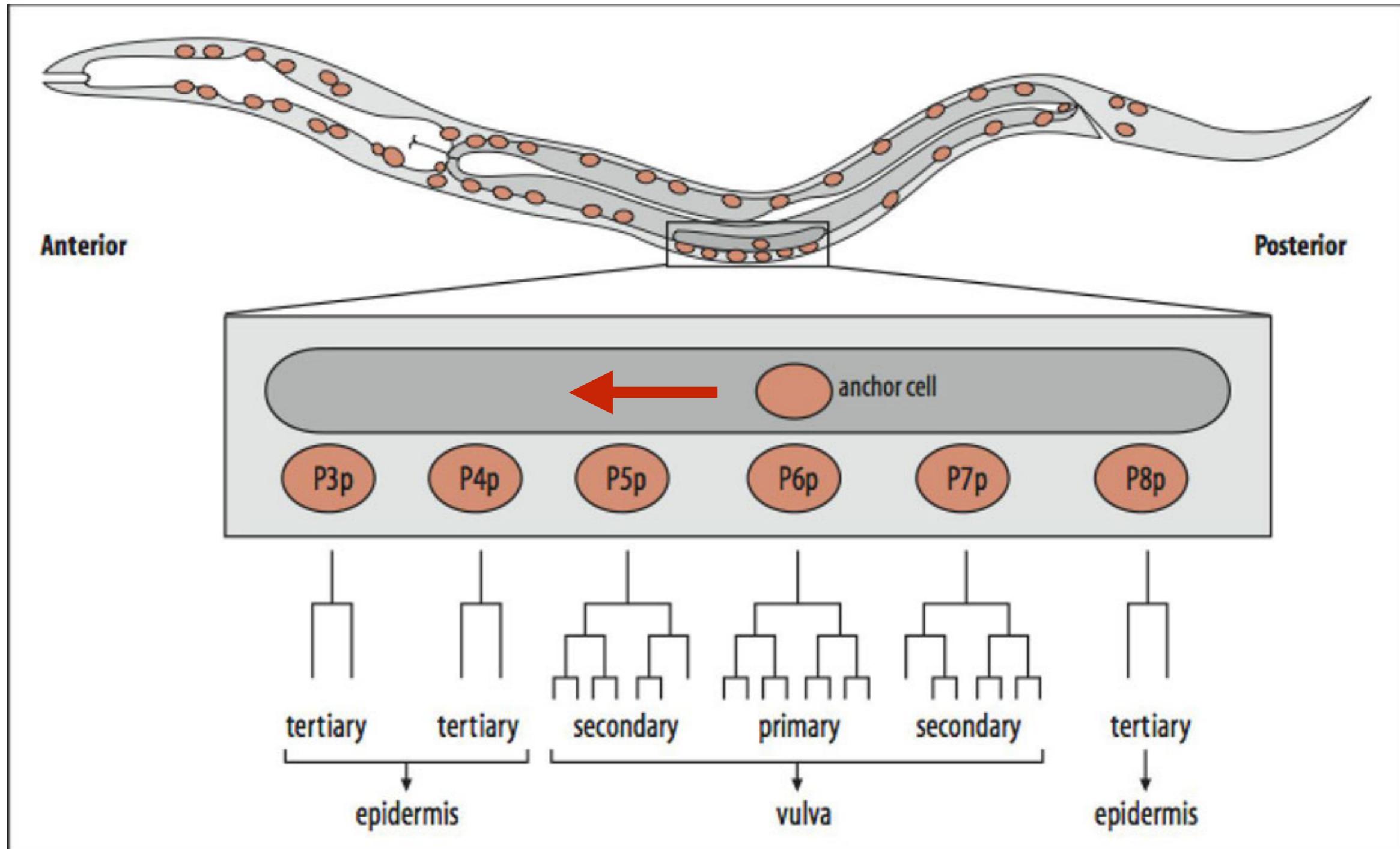


No AC leads to no vulval cell specification
and a vulvaless phenotype

All other vulval mutants are epistatic to AC ablation

Mutant genotypes	Phenotype
AC ablation	Vul
<i>lin-1(0)</i>	Muv
<i>lin-3(0)</i>	Vul
<i>let-60(gf)</i>	Muv
<i>let-23(gf)</i>	Muv
AC ablation; <i>let-23(gf)</i>	Muv
AC ablation; <i>let-60(gf)</i>	Muv
AC ablation; <i>lin-1(0)</i>	Muv

Moving the AC moves the vulva (sufficient)

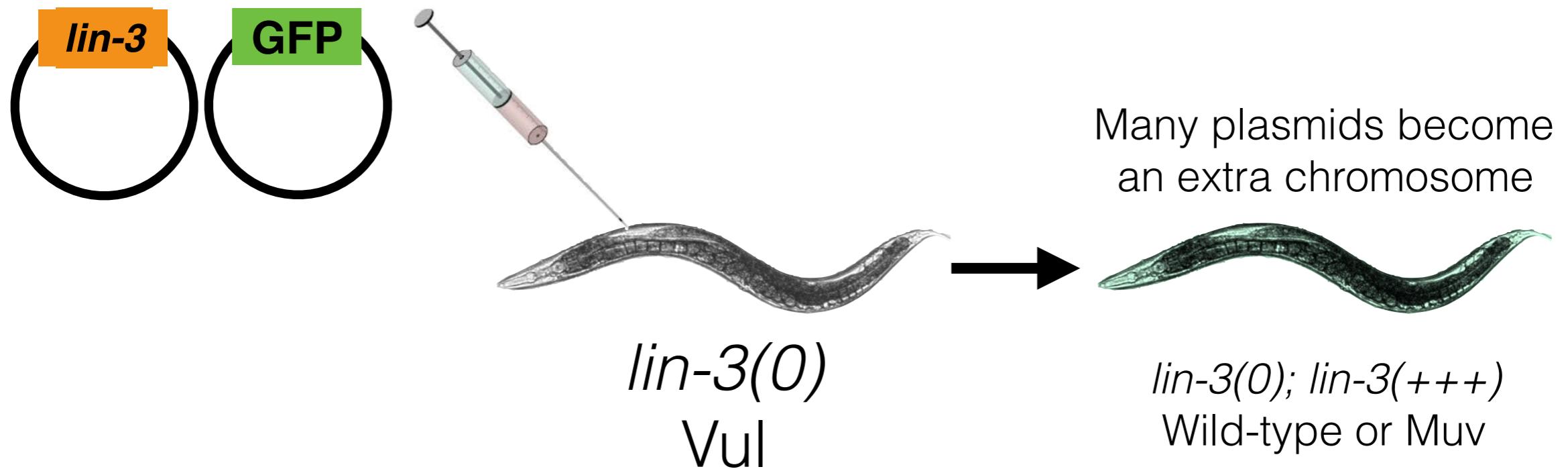


dig-1 displaced gonad mutants

AC ablation and loss of *lin-3* have the same phenotype and epistatic relationships

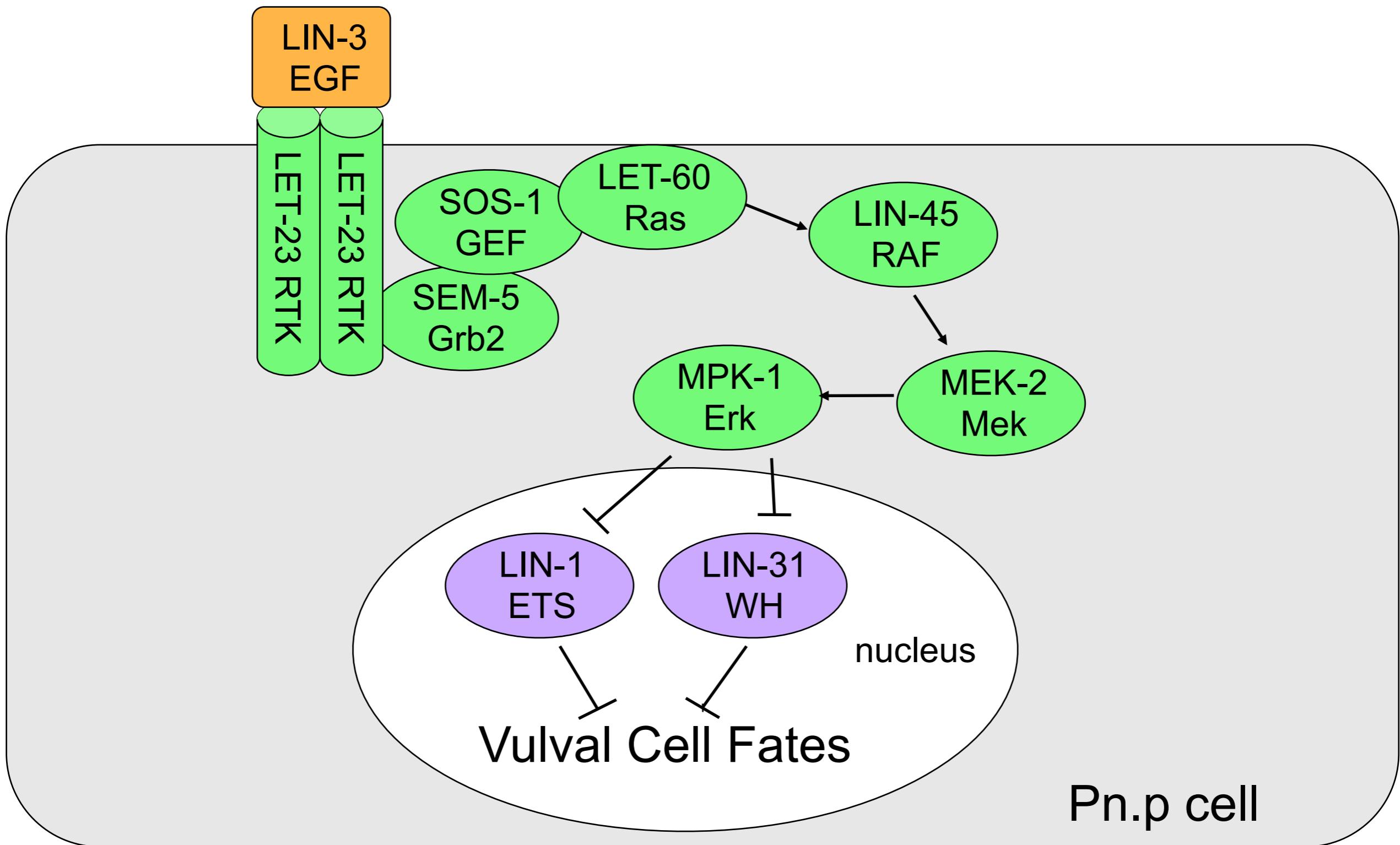
Mutant genotypes	Phenotype
AC ablation	Vul
<i>lin-3(0)</i>	Vul
<i>lin-1(0)</i>	Muv
<i>let-60(gf)</i>	Muv
<i>let-23(gf)</i>	Muv
AC ablation; <i>let-23(gf)</i>	Muv
AC ablation; <i>let-60(gf)</i>	Muv
AC ablation; <i>lin-1(0)</i>	Muv
<i>lin-3(0); let-23(gf)</i>	Muv
<i>lin-3(0); let-60(gf)</i>	Muv
<i>lin-3(0); lin-1(0)</i>	Muv

LIN-3 is expressed in the AC; does it function there?

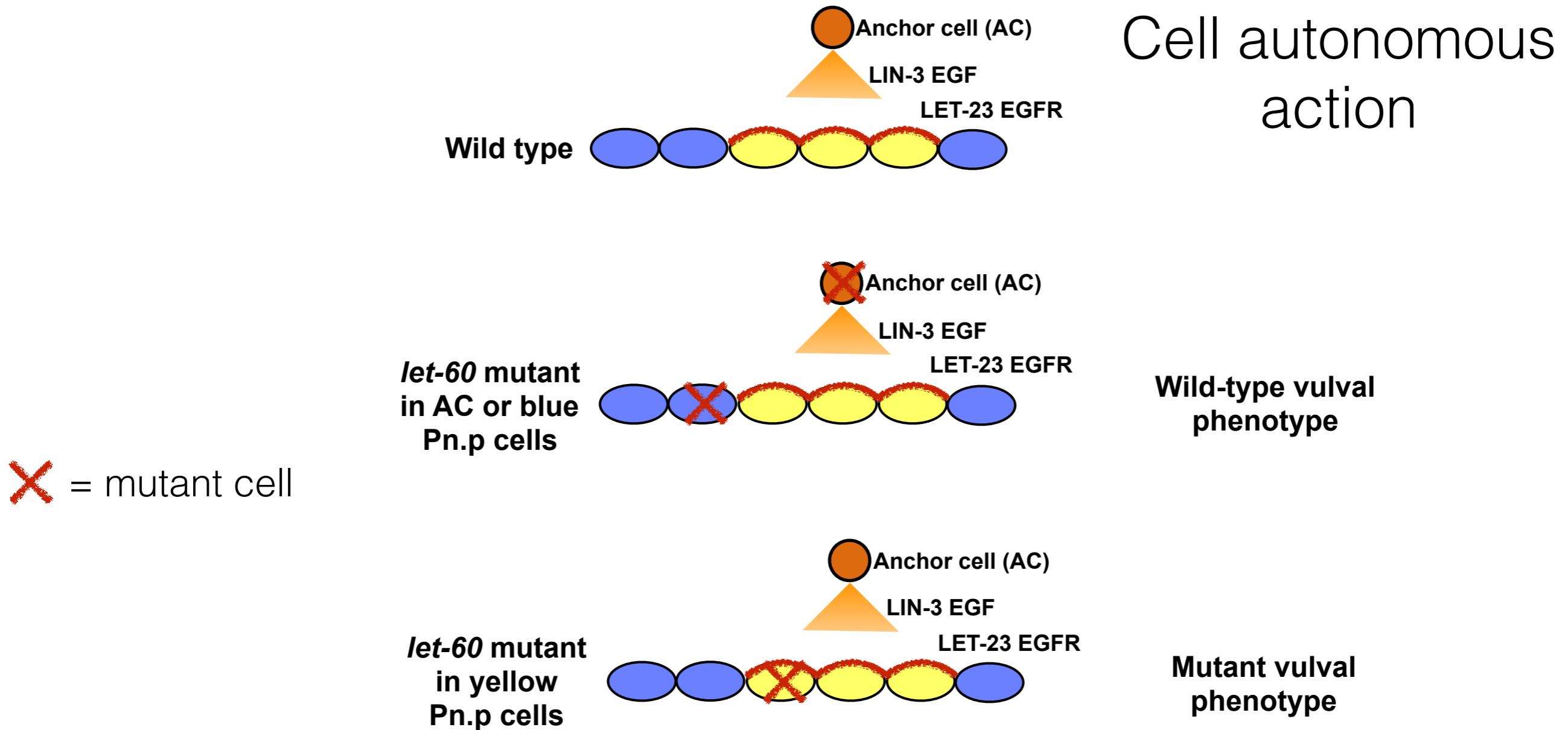


Genotype	Phenotype	
<i>lin-3(0)</i>	Vul	
<i>lin-3(0); lin-3(+++)</i>	Wild-type	
<i>lin-3(0); vulval cell:lin-3(+++)</i>	Vul	Expression of <i>lin-3</i> in vulval cells
<i>lin-3(0); intestine:lin-3(+++)</i>	Vul	Expression of <i>lin-3</i> in the intestine
<i>lin-3(0); neurons:lin-3(+++)</i>	Vul	Expression of <i>lin-3</i> in the neurons
<i>lin-3(0); AC:lin-3(+++)</i>	Wild-type	Expression of <i>lin-3</i> in the AC

A Ras pathway promotes vulval fates

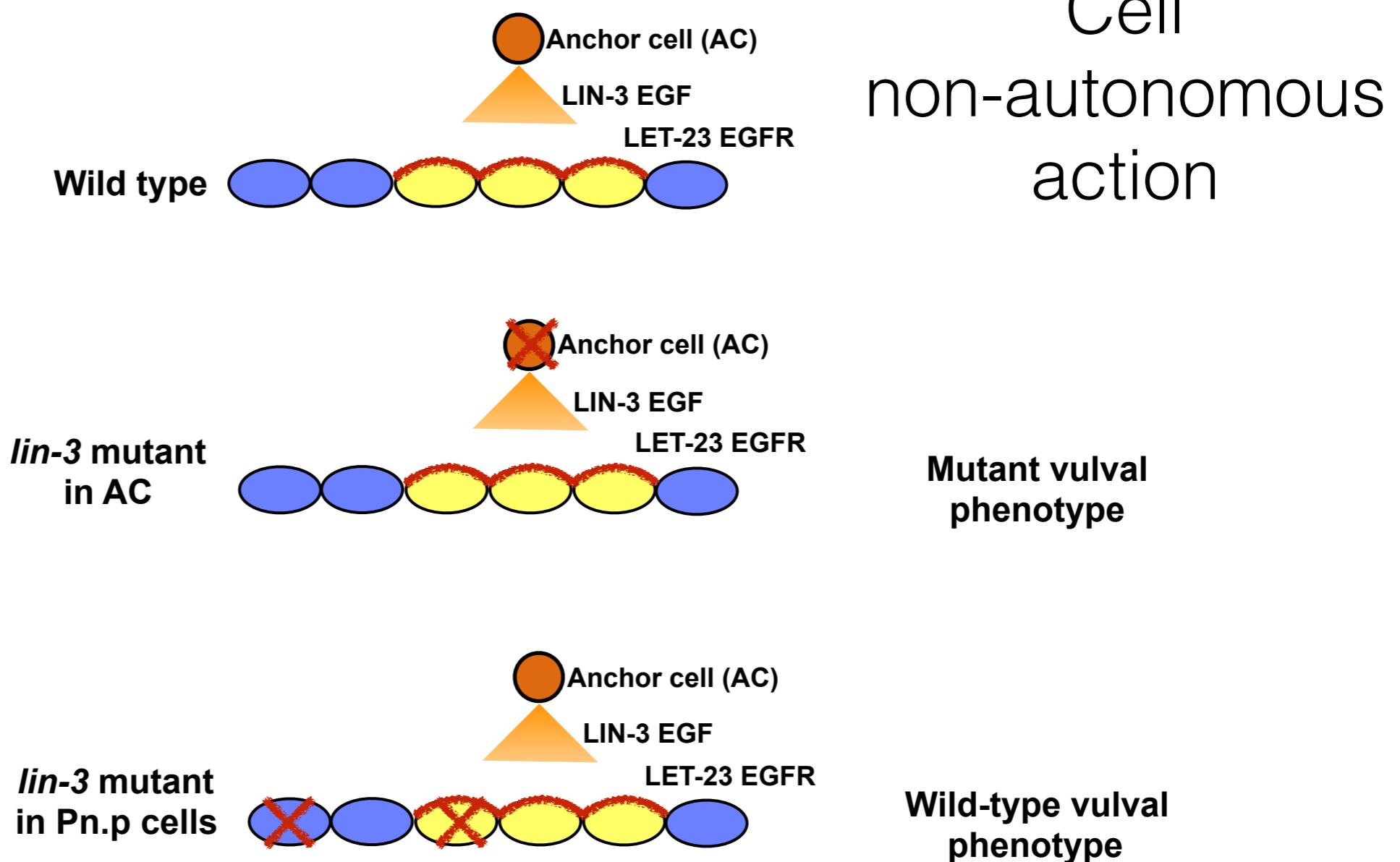


Cell autonomy of *let-60*



A cell autonomous trait is a trait in a multicellular organism in which only genotypically mutant cells have the mutant phenotype

Cell autonomy of *lin-3*



A cell autonomous trait is a trait in a multicellular organism in which only genotypically mutant cells have the mutant phenotype

Vulval mutants

Mutant	Phenotype
<i>lin-1(0)</i>	Muv
<i>lin-3(0)</i>	Vul
<i>let-60(0)</i>	Vul
<i>let-60(af)</i>	Muv
<i>let-23(0)</i>	Vul
<i>let-23(af)</i>	Muv
<i>lin-2(0)</i>	a little Vul
<i>lin-7(0)</i>	a little Vul
<i>lin-10(0)</i>	a little Vul

Vulval mutants

Mutant	Phenotype
<i>lin-1(0)</i>	Muv
<i>lin-3(0)</i>	Vul
<i>let-60(0)</i>	Vul
<i>let-60(af)</i>	Muv
<i>let-23(0)</i>	Vul
<i>let-23(af)</i>	Muv
<i>lin-2(0)</i>	a little Vul
<i>lin-7(0)</i>	a little Vul
<i>lin-10(0)</i>	a little Vul
<i>lin-2(0); lin-7(0)</i>	more Vul
<i>lin-2(0); lin-10(0)</i>	more Vul
<i>lin-7(0); lin-10(0)</i>	more Vul

Vulval mutants

Mutant	Phenotype
<i>lin-1(0)</i>	Muv
<i>lin-3(0)</i>	Vul
<i>let-60(0)</i>	Vul
<i>let-60(af)</i>	Muv
<i>let-23(0)</i>	Vul
<i>let-23(af)</i>	Muv
<i>lin-2(0)</i>	a little Vul
<i>lin-7(0)</i>	a little Vul
<i>lin-10(0)</i>	a little Vul
<i>lin-2(0); lin-7(0)</i>	more Vul
<i>lin-2(0); lin-10(0)</i>	more Vul
<i>lin-7(0); lin-10(0)</i>	more Vul
<i>lin-2(0); lin-7(0); lin-10(0)</i>	Vul

Told the geneticists that the three genes act in a redundant process,
but how?

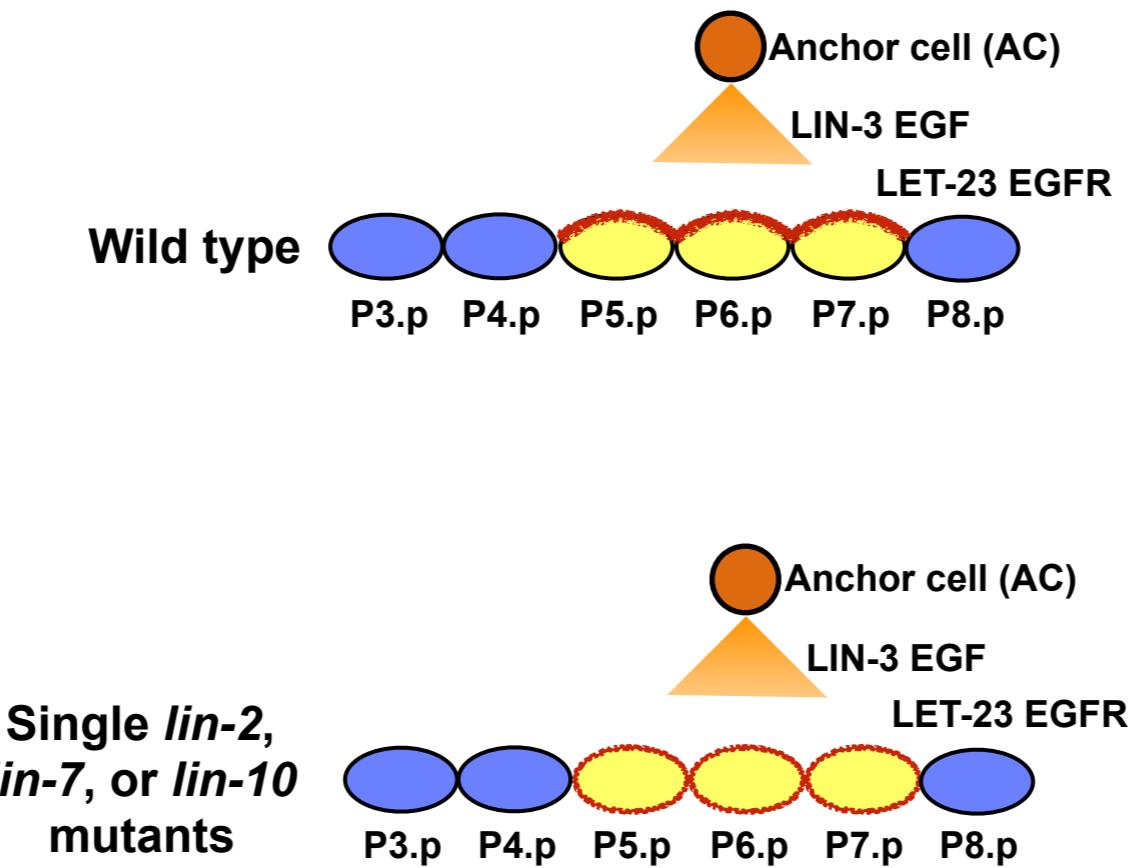
**Told the geneticists that the three genes
act in a redundant process
But how?**

Act downstream of *lin-3*

Act upstream of *let-23*

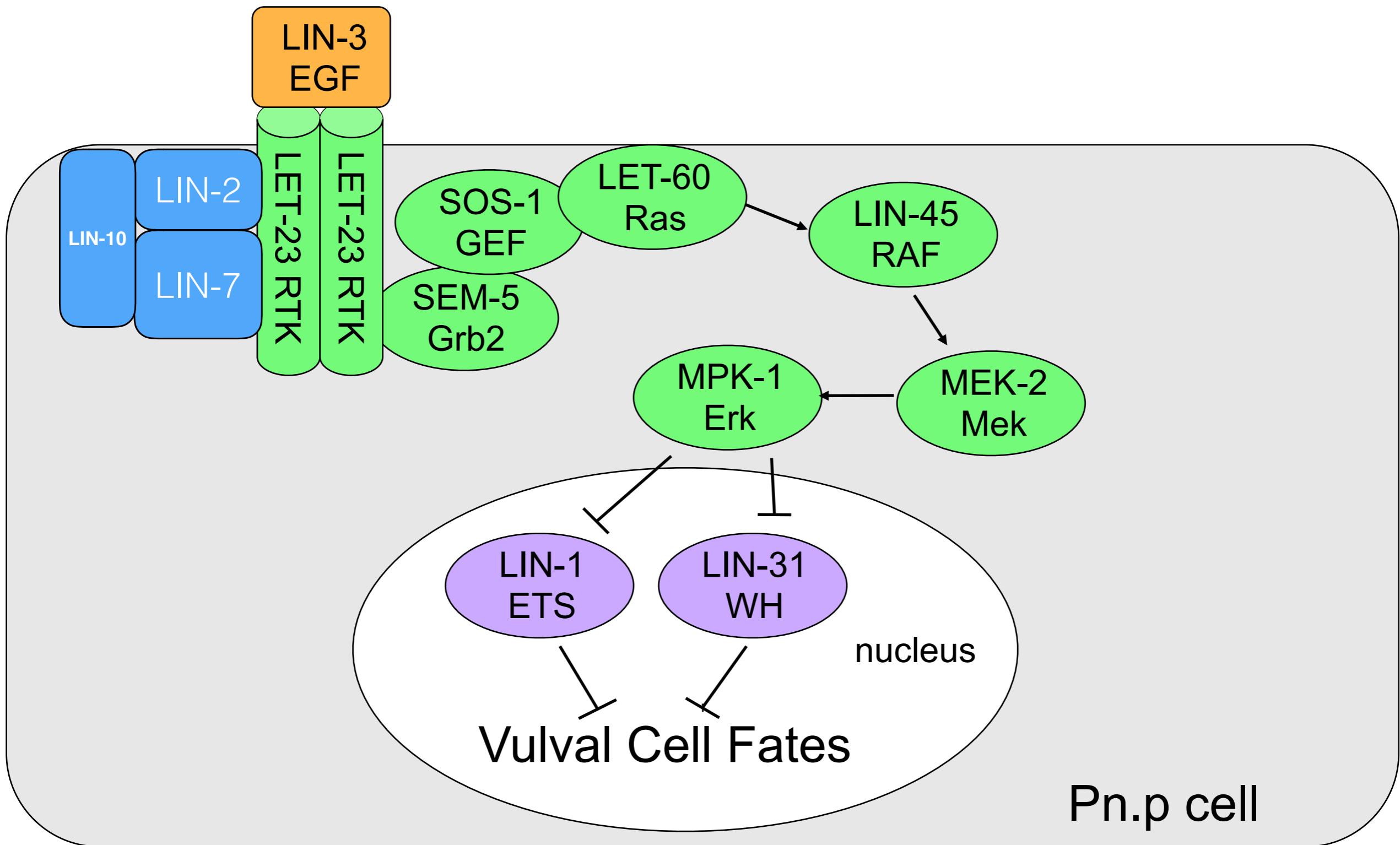
Expressed in vulval cells

lin-2, *lin-7*, and *lin-10* act redundantly to localize LET-23 RTK



lin-3
↓
lin-2/7/10 → *let-23* → *let-60* → *lin-1* → vulval fate

A Ras pathway promotes vulval fates



Vulval mutants

Mutant	Phenotype
<i>lin-1(0)</i>	Muv
<i>lin-3(0)</i>	Vul
<i>let-60(0)</i>	Vul
<i>let-60(gf)</i>	Muv
<i>let-23(0)</i>	Vul
<i>let-23(gf)</i>	Muv
<i>lin-2(0)</i>	~Vul
<i>lin-7(0)</i>	~Vul
<i>lin-10(0)</i>	~Vul
<i>lin-8(0); lin-9(0)</i>	synMuv

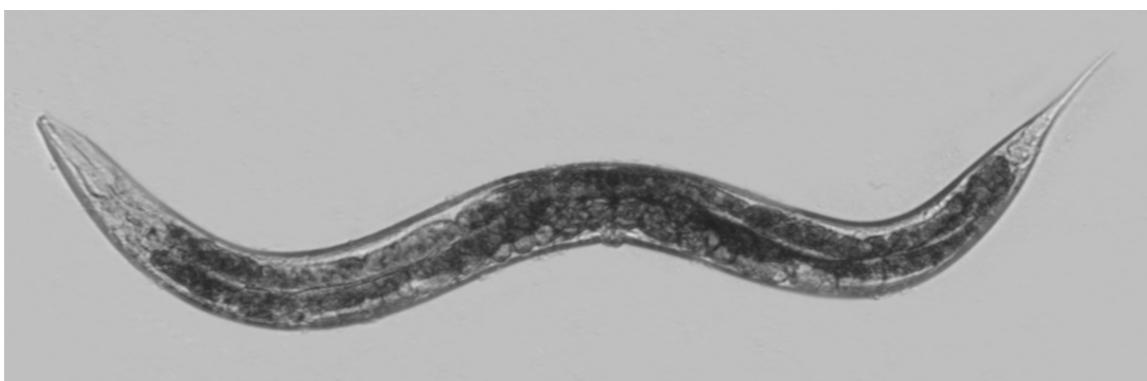
synMuv = synthetic Multivulva

Two parallel pathways

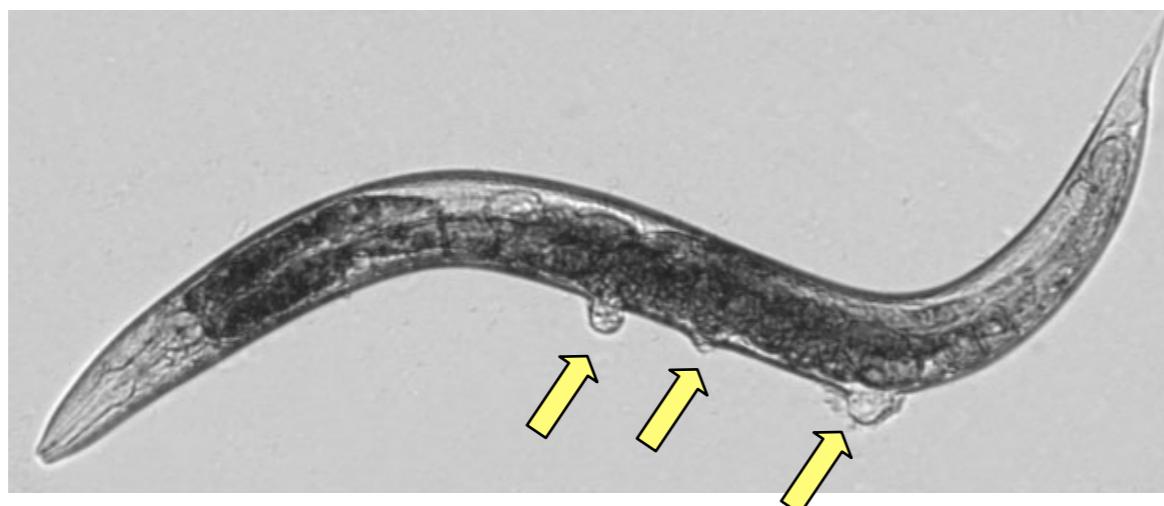
The synMuv phenotype is caused by mutations in both class A and B genes



class A single mutant



class B single mutant

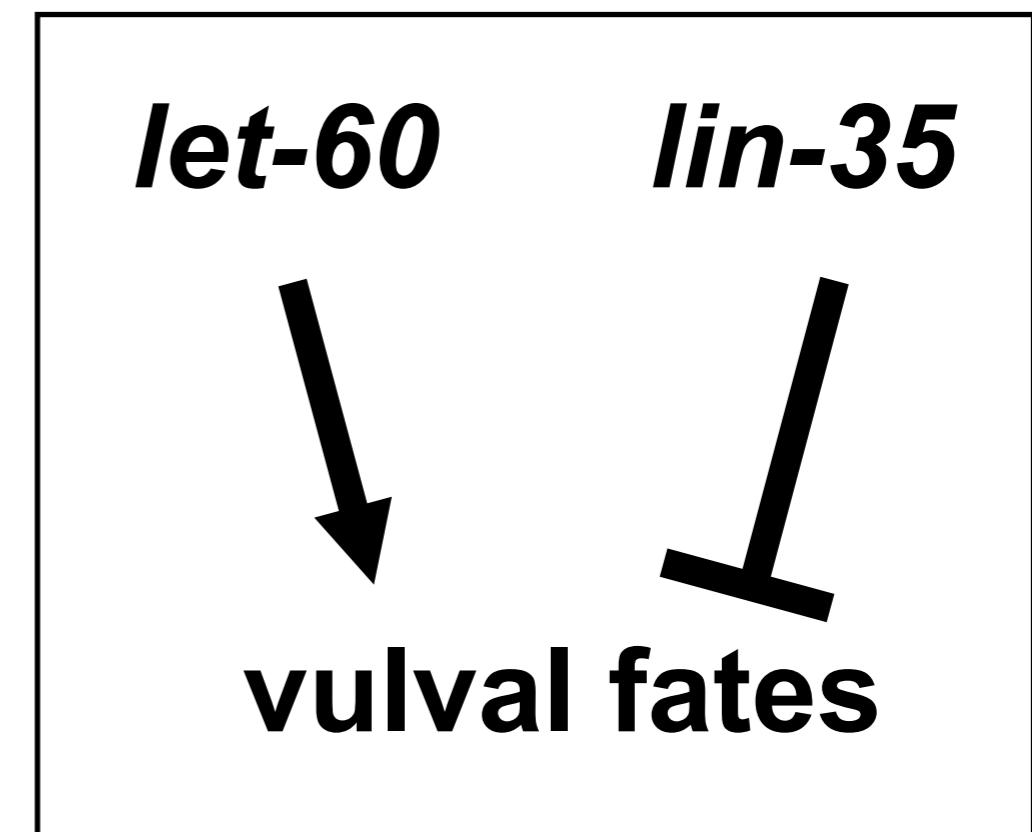
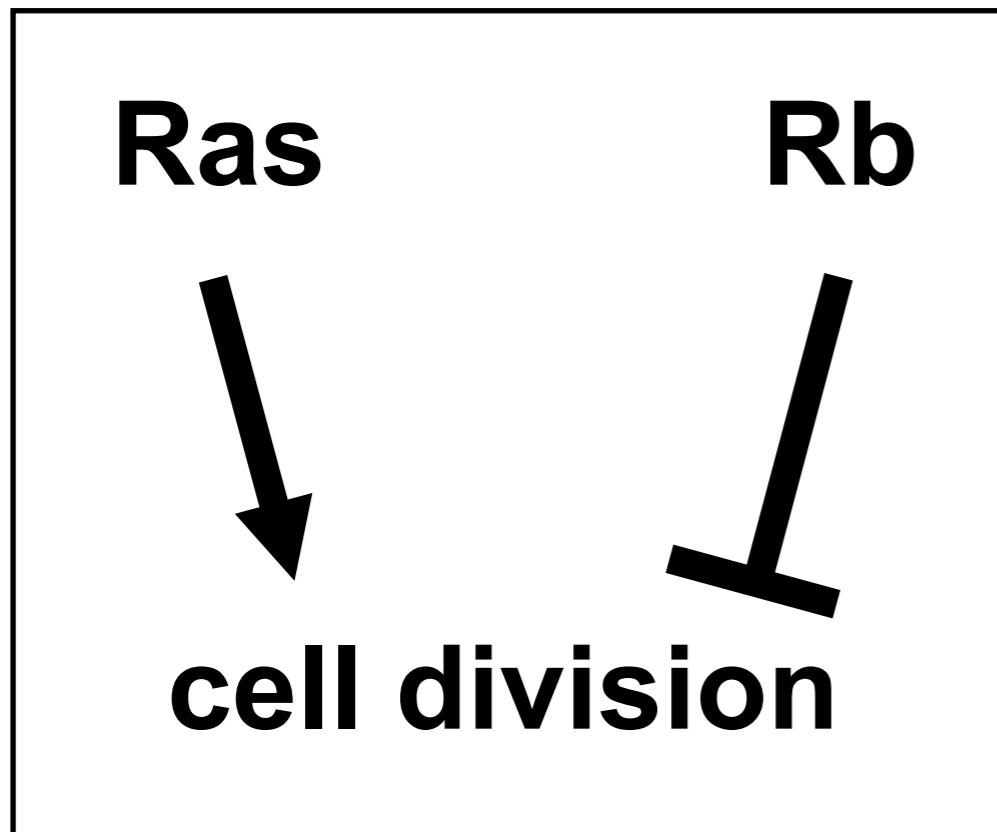


class AB double mutant

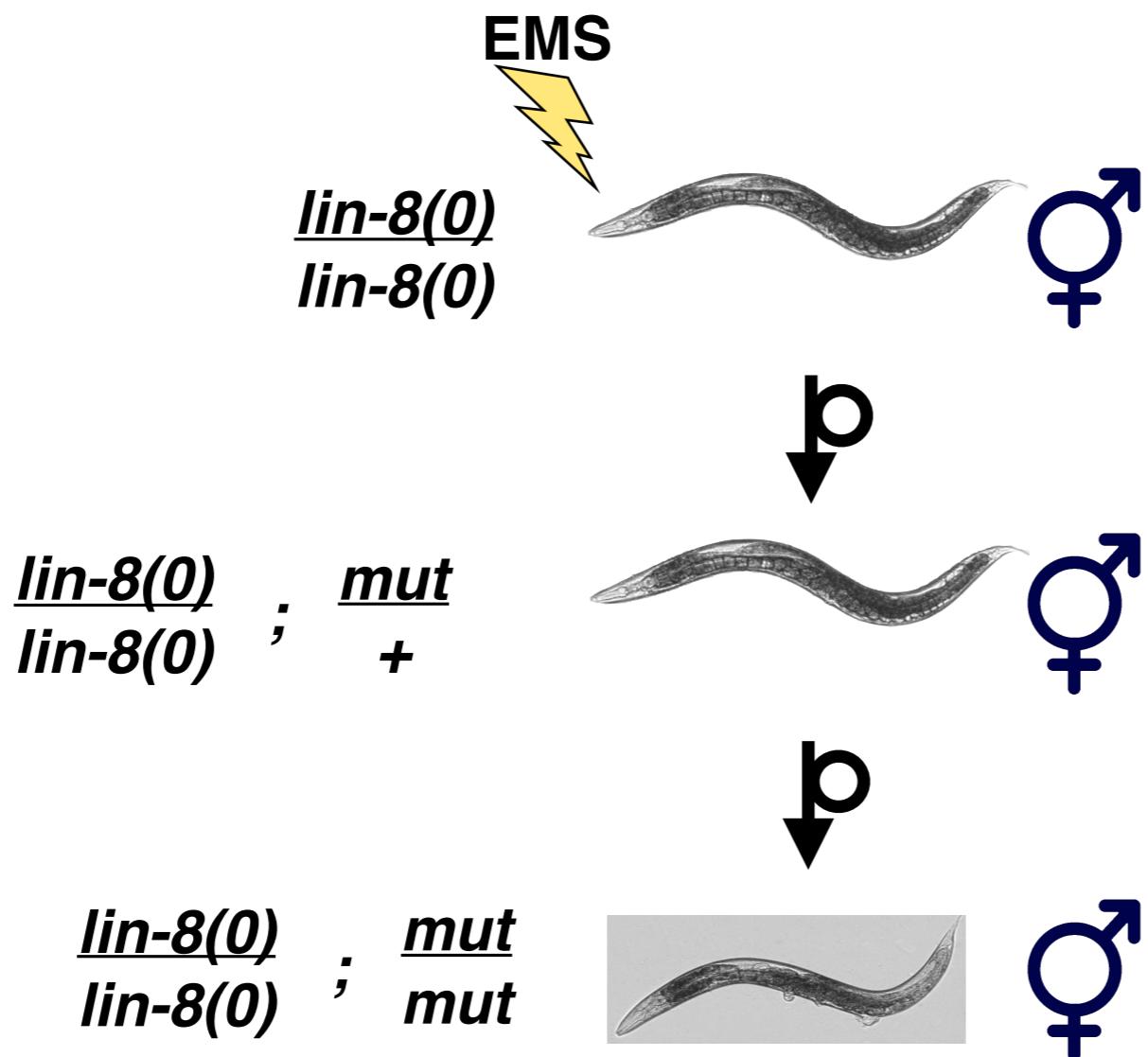
Double mutants within the same class are not enhanced and not multivulva

	Class A	Class B
Class A	non-Muv	Muv
Class B		non-Muv

The vulval cell-fate decision models cell-fate decisions involved in carcinogenesis



Enhancer screens to find more synthetic multivulva genes



The synMuv genes

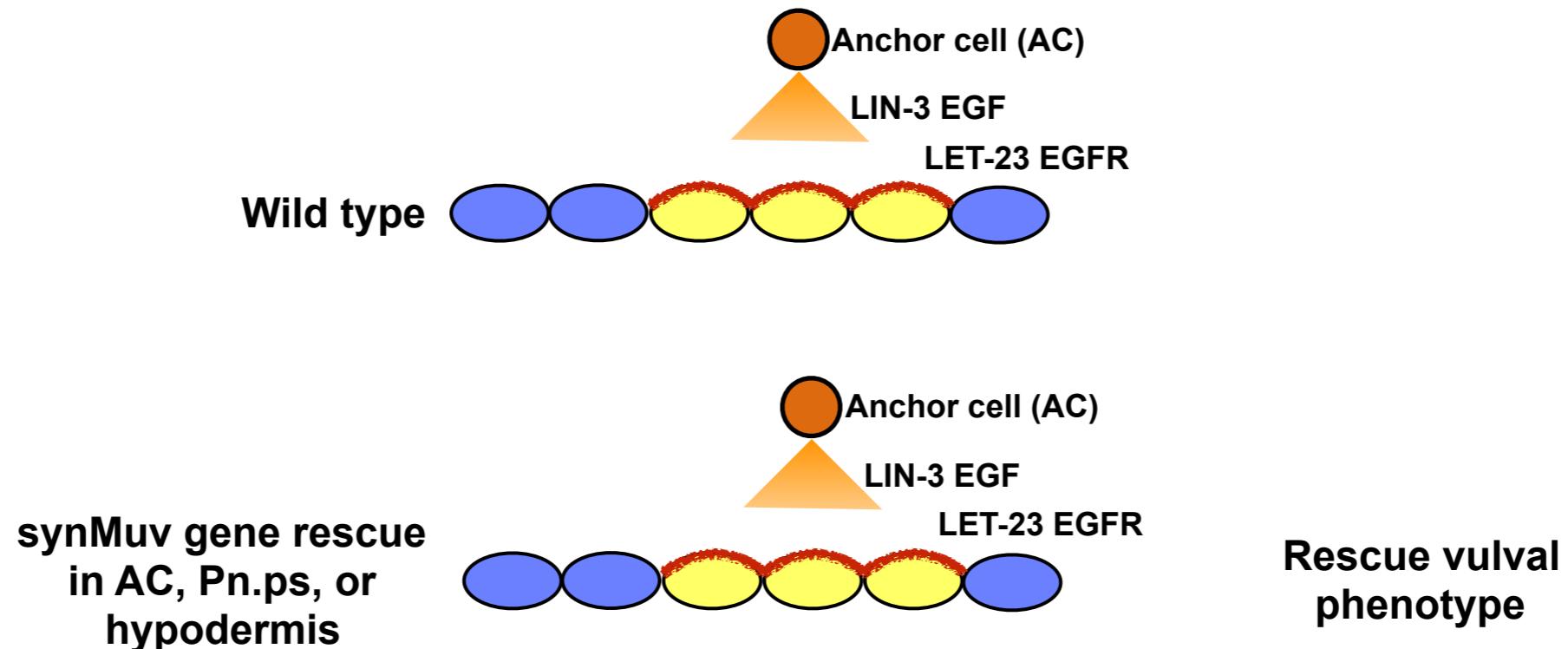
class A

<i>lin-8</i>	novel
<i>lin-15A</i>	THAP
<i>lin-38</i>	Zn finger
<i>lin-56</i>	THAP

class B

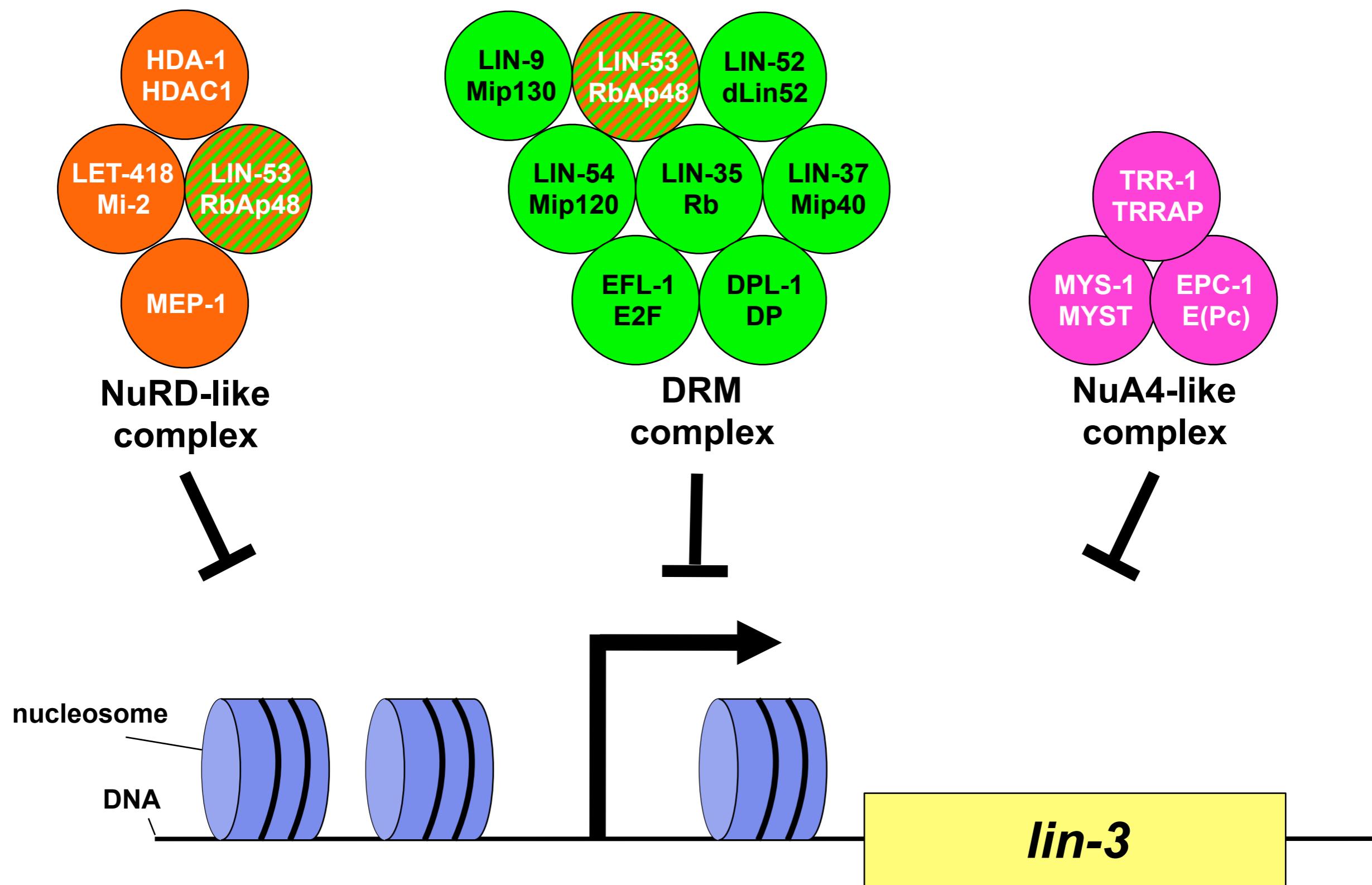
<i>lin-9</i>	Mip130/ALY	<i>dpl-1</i>	DP	<i>trr-1</i>	TRRAP
<i>lin-13</i>	Zn fingers	<i>efl-1</i>	E2F4	<i>mys-1</i>	HAT
<i>lin-15B</i>	THAP	<i>lin-35</i>	Rb	<i>epc-1</i>	E(Pc)
<i>lin-36</i>	THAP	<i>lin-53</i>	RbAp48		
<i>lin-37</i>	Mip40	<i>hda-1</i>	HDAC1		
<i>lin-52</i>	dLin52	<i>let-418</i>	Mi-2		
<i>lin-54</i>	Mip120	<i>mep-1</i>	Zn fingers		
<i>lin-61</i>	I(3)MBT	<i>hpl-2</i>	HP1		
<i>lin-65</i>	novel	<i>gap-1</i>	RasGAP		
<i>tam-1</i>	RING finger	<i>sli-1</i>	c-Cbl		
		<i>ark-1</i>	Ack		

Cell autonomy of synMuv genes

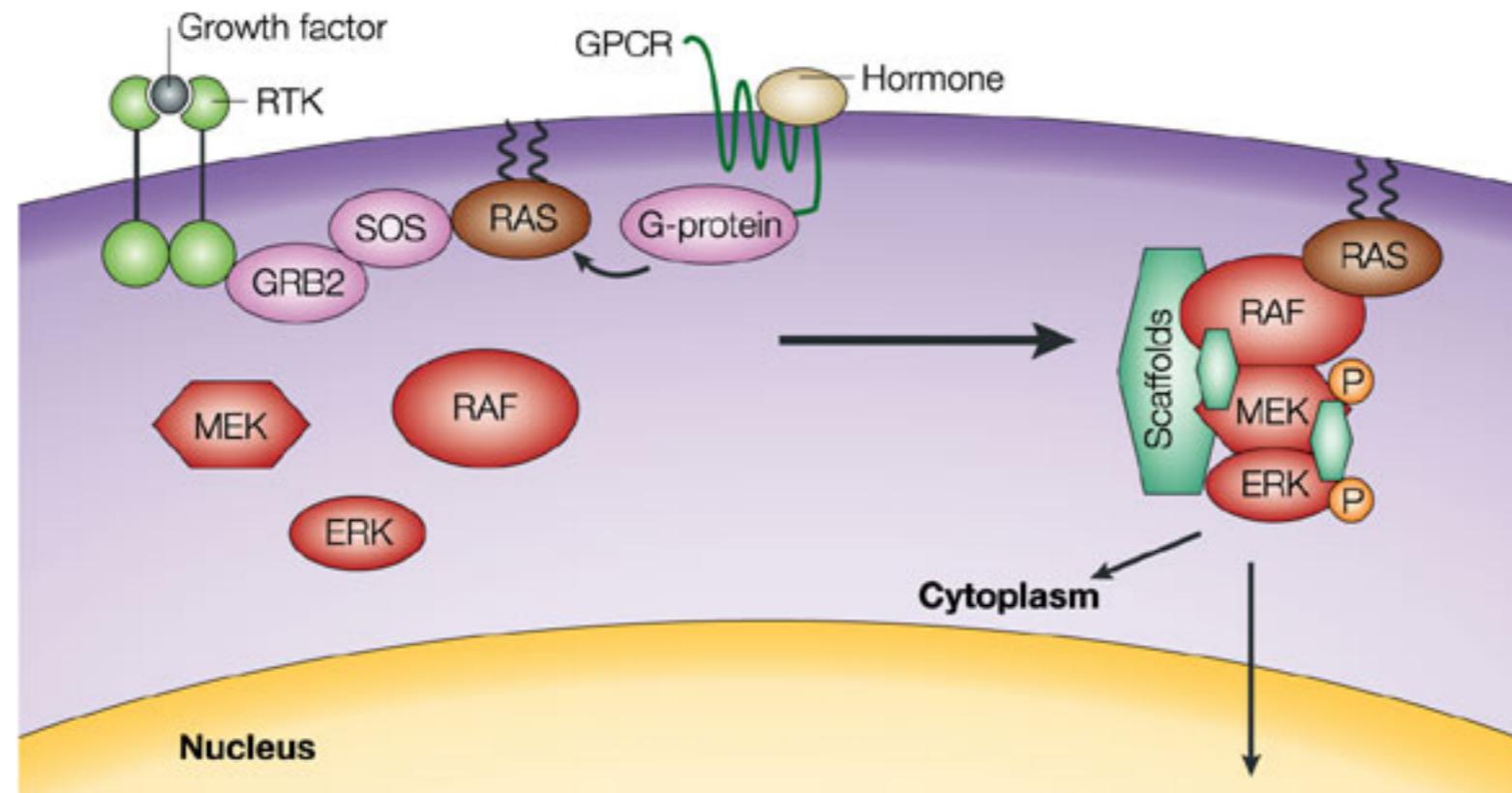


Cell
non-autonomous
action

NuRD-like, DRM, and NuA4-like complexes affect the transcription of vulval genes



Two decades of research in *Drosophila* and *C. elegans* led to these pathways



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We NEED basic research for this reason!