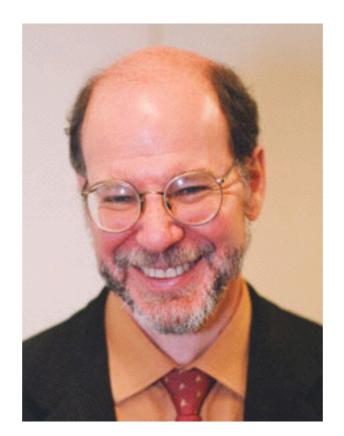
# **Bio393: Genetic Analysis**

Step-wise genetic analysis



**Bob Horvitz** 

# "Model organisms" are everywhere now

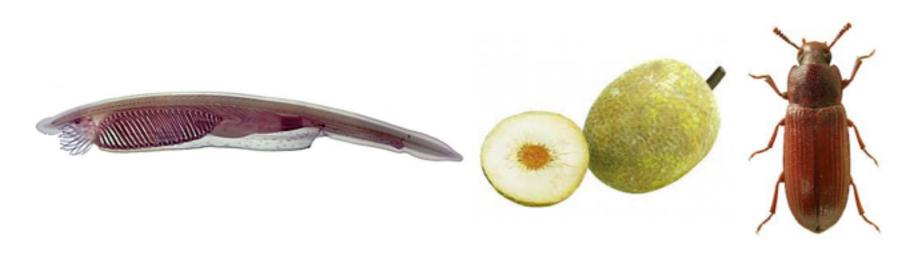














# 1. Define the problem



Let the question choose the organism (not the other way around)

# 2. Choose an organism

Organism	Time to 10 <sup>6</sup>	Space
Bacteriaphage	1 hour	10 nL
Bacteria	15 hours	1 µL
Yeast	1 day	0.1 mL
Worm	10 days	6 cm cube
Fly	6 weeks	0.5 m cube
Mouse	3 years	Half Pancoe
Human	750 years	Evanston + Wilmette

10<sup>6</sup> individuals to study 10<sup>-6</sup> mutation rate

 $2^{20} \approx 10^6$  individuals

# 3. Perform a mutant hunt

To mutagenize or not to mutagenize?

Yes	No	
10-3	10-6	LoF mutation
10-5-10-6	10 <sup>-8</sup> -10 <sup>-9</sup>	Specific mutation

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C. elegans

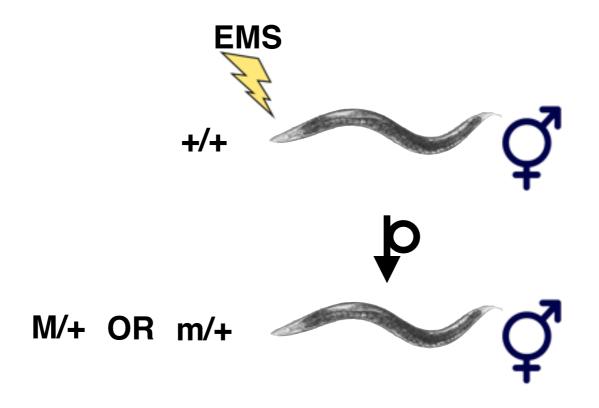
~20,000 genes 20 LoF mutations D. melanogaster



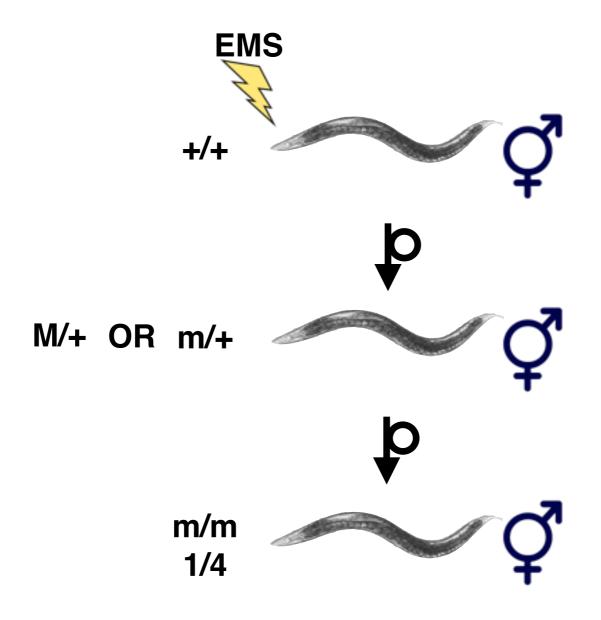
~12,000 genes 12 LoF mutations



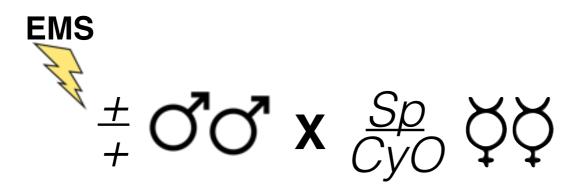
C. elegans screens for dominant or recessive phenotypes

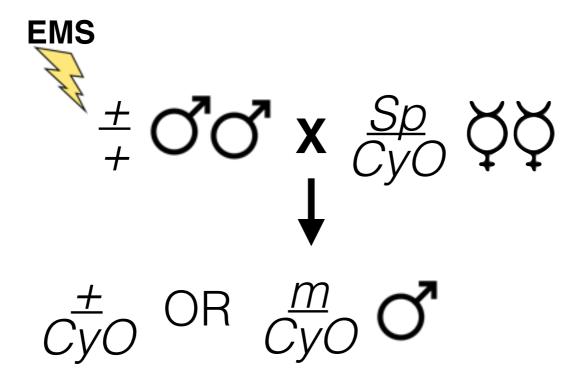


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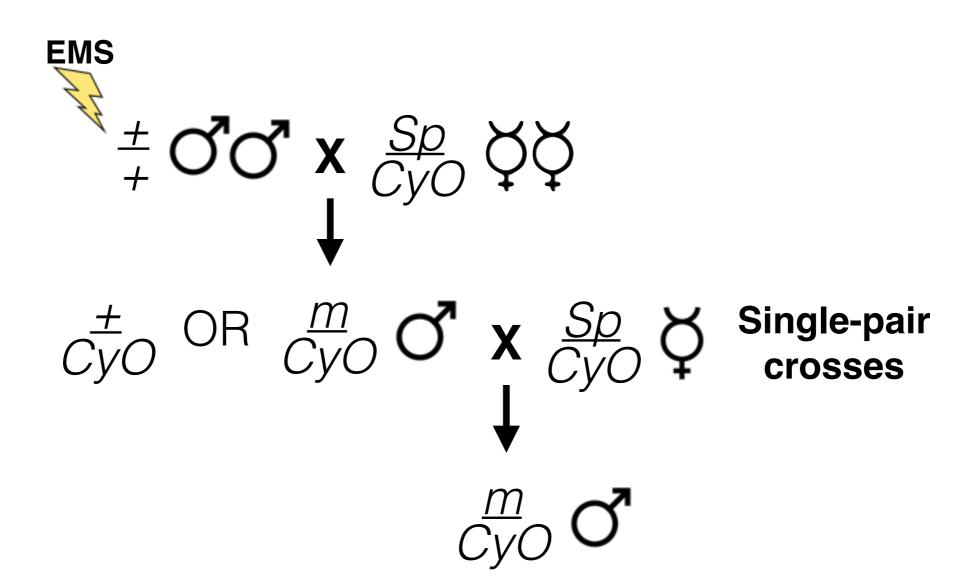


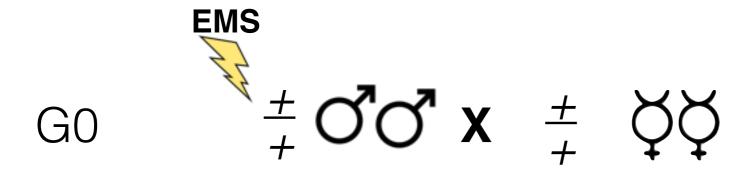
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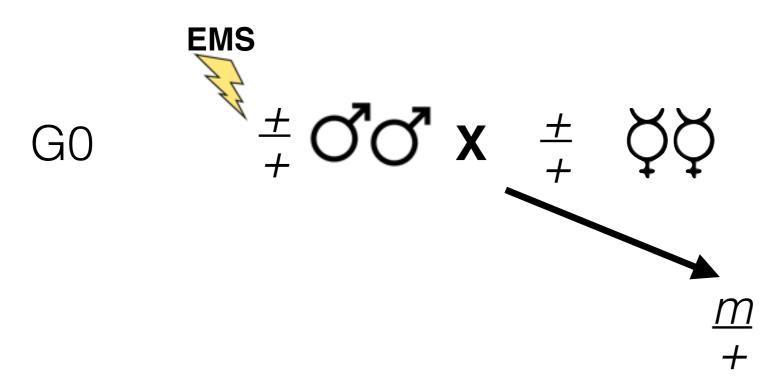


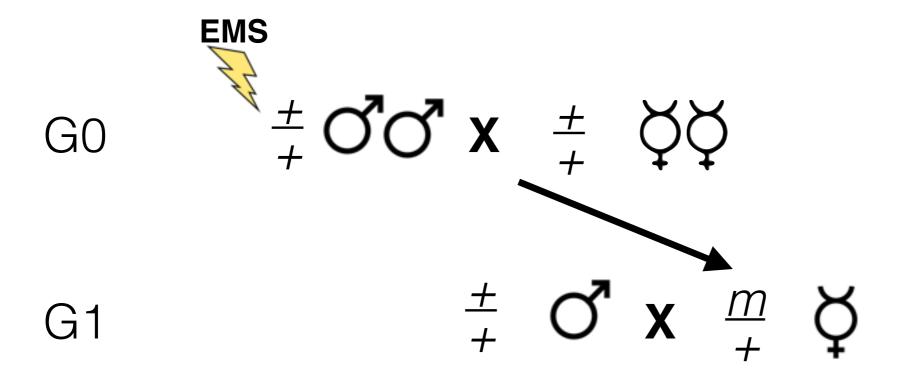


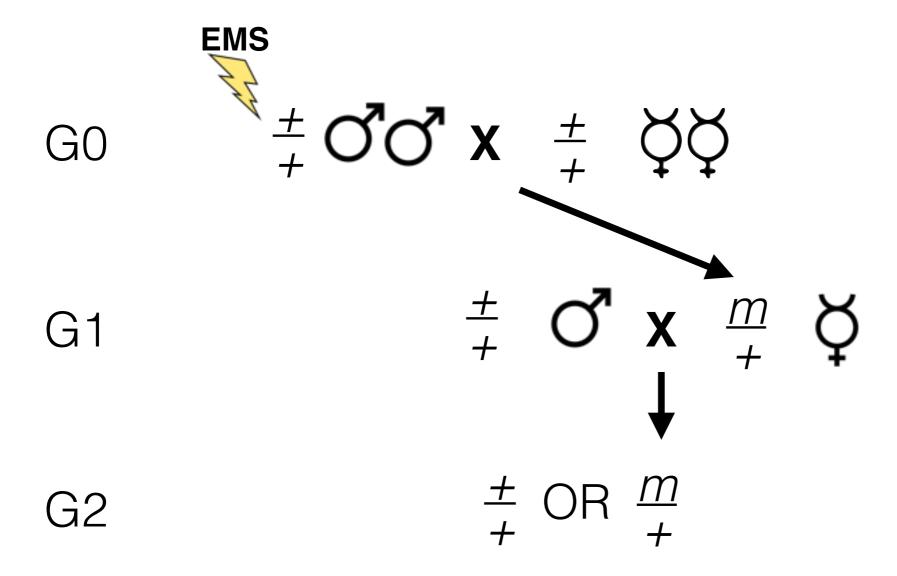
EMS 
$$\frac{\pm}{+}$$
  $\overrightarrow{O}$   $\overrightarrow{O}$   $\overrightarrow{X}$   $\frac{Sp}{CyO}$   $\overleftrightarrow{Q}$   $\overset{+}{\downarrow}$   $\overset{-}{\downarrow}$  OR  $\frac{m}{CyO}$   $\overrightarrow{O}$   $\overset{-}{X}$   $\frac{Sp}{CyO}$   $\overset{-}{\downarrow}$  Single-pair crosses

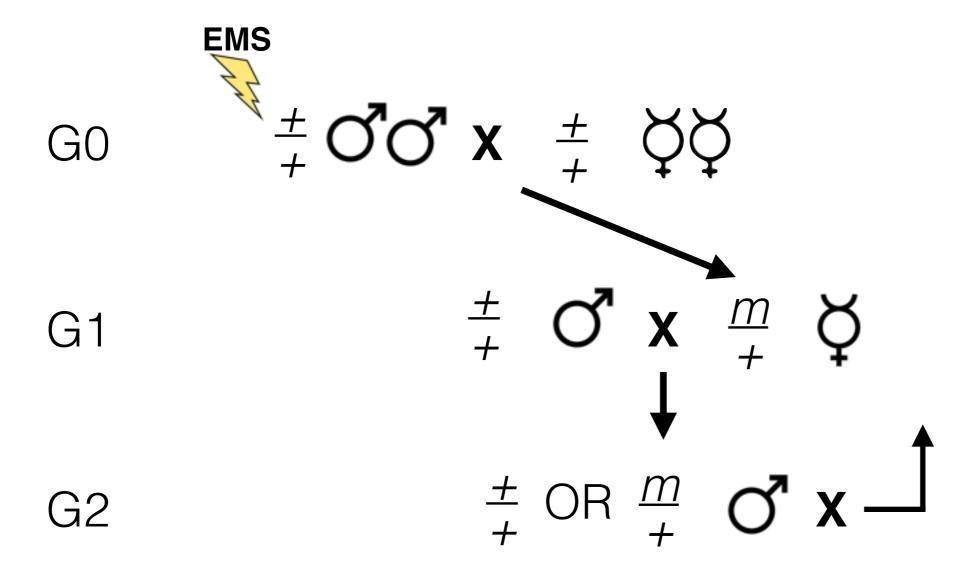


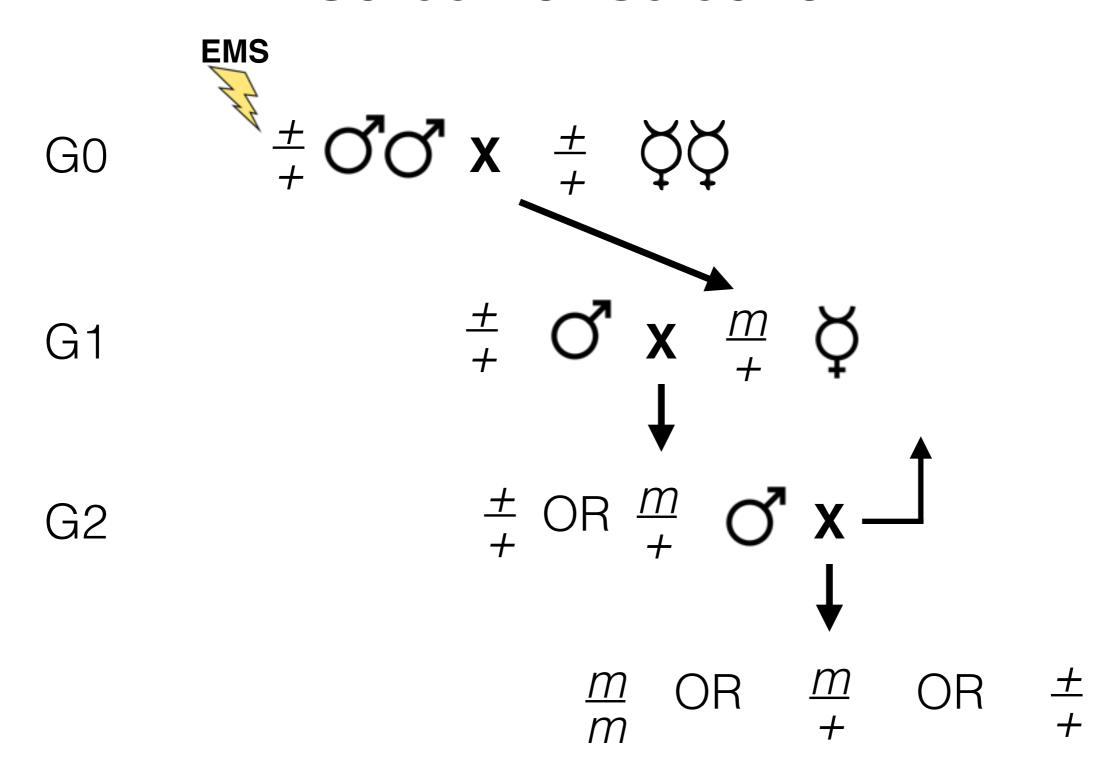












Remember hemizygous screens too

Use Poisson sampling and common sense

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Saturation of the investigators patience

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Change mutagens

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Why might we miss genes?

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#### Why might we miss genes?

Numbers are too small

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#### Why might we miss genes?

Numbers are too small Pleiotropy

Use Poisson sampling and common sense

Saturation of the investigators patience

Change mutagens

#### Why might we miss genes?

Numbers are too small Pleiotropy Redundancy

True-breeding stocks

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Balancers, balanced stocks

#### 5. Establish a strain

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Freeze organisms

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True-breeding stocks

Balancers, balanced stocks

Freeze organisms

The most common phenotypes are sterile or dead!

Mutagenesis adds hundreds of mutations randomly throughout the genome.

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Backcross = cross to parent used in the screen/selection

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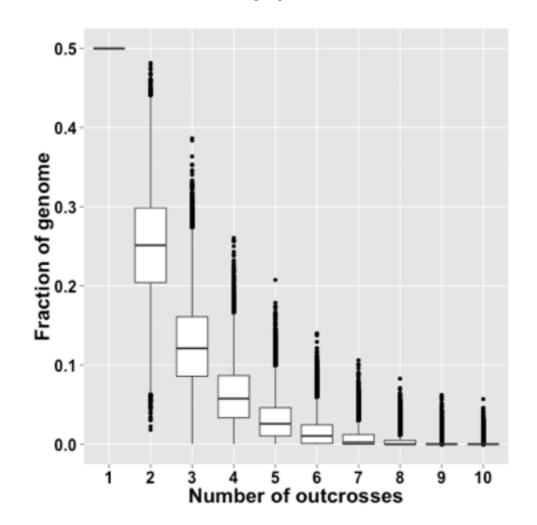
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Outcross = cross to a wild-type strain

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8. Single-gene phenotype?

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9. Mapping and complementation

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9. Mapping and complementation

What have we discovered so far?

## 10. Characterize the phenotype

Look at the wild-type and mutant organisms in detail

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

What could be mutated?

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Look at the wild-type and mutant organisms in detail



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

What could be mutated?

No embryos

No vulva

No vulval muscles

No neurons

Or malfunction of any vulva, muscle, or neuron

11. Define the nature of the mutant allele(s)

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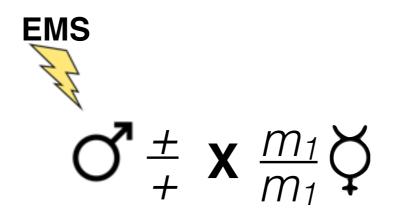
Loss of function more likely than gain of function

## 11. Define the nature of the mutant allele(s)

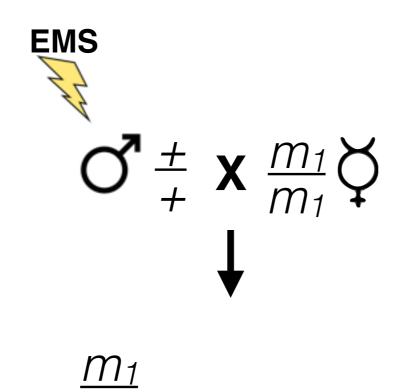
Loss of function more likely than gain of function

What if you only have one mutant?

## 12. Perform non-complementation screens

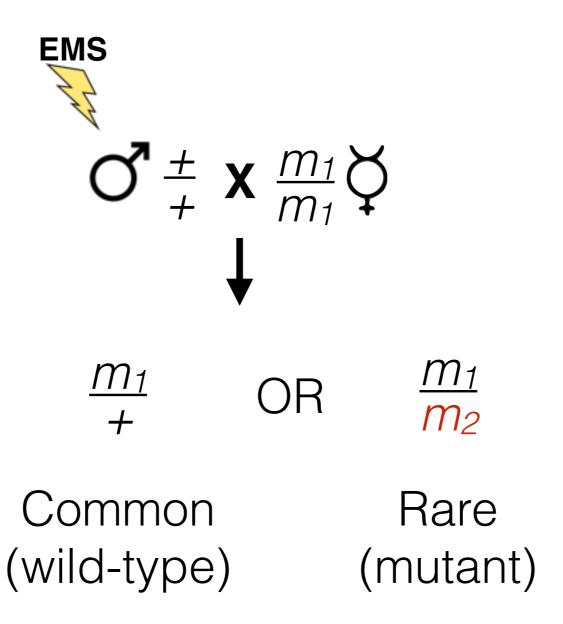


## 12. Perform non-complementation screens



Common (wild-type)

## 12. Perform non-complementation screens



We know the location of the gene

Use deficiencies and duplications to study dosage

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1. Is the phenotype dominant or recessive?

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We know the location of the gene

Use deficiencies and duplications to study dosage

- 1. Is the phenotype dominant or recessive?
- 2. Recessive: hypomorph or null
- 3. Dominant: hypermorph, neomorph, antimorph

## 14. Define the null phenotype

What happens with a complete loss of gene function?

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What happens with a complete loss of gene function?

Dosage studies, non-complementation screens, and characterization of the mutant phenotype tell you about the null phenotype

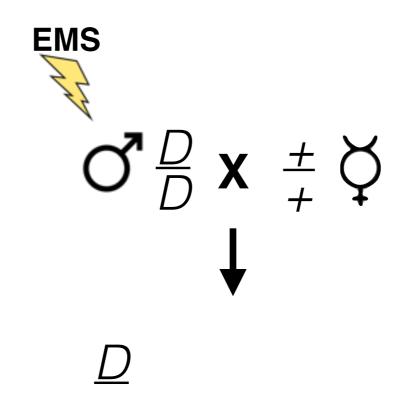
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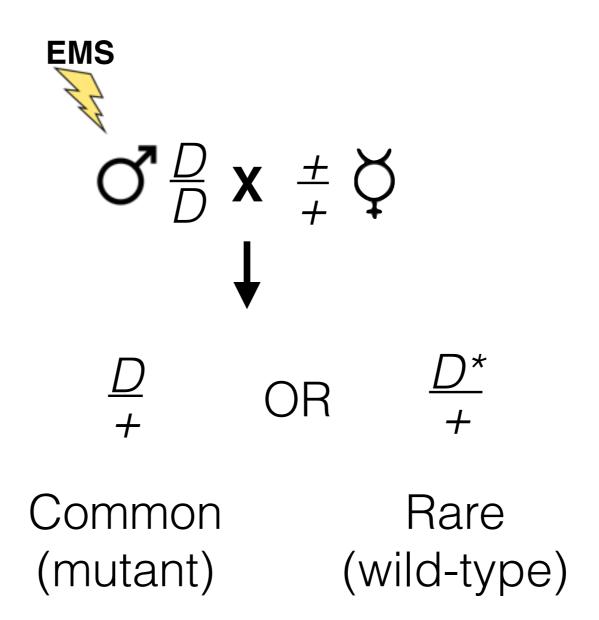
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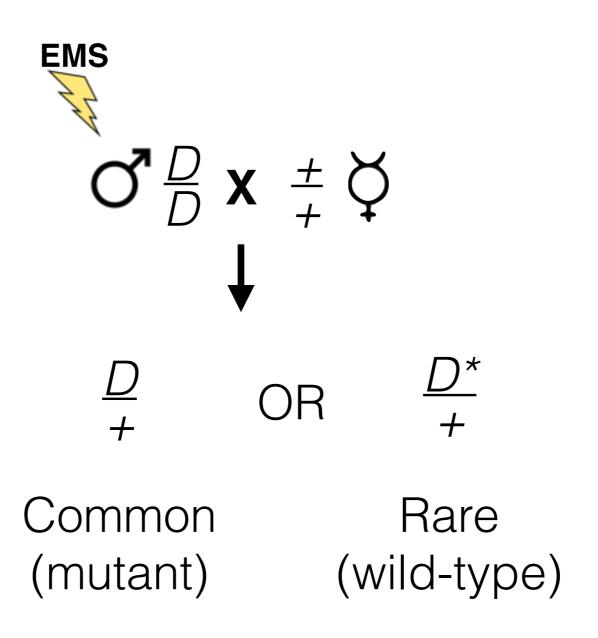
What if you have a mutant with a dominant gain-of-function phenotype?

EMS
$$O^{T} \frac{D}{D} \mathbf{x} \stackrel{+}{\leftarrow} \mathbf{\nabla}$$

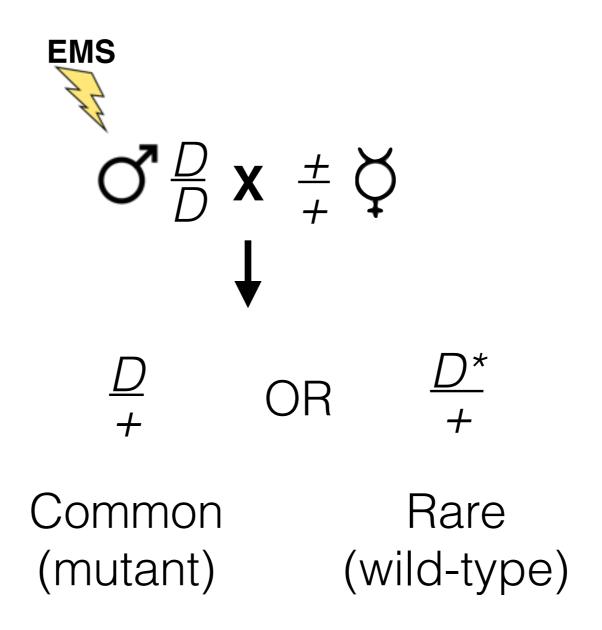


Common (mutant)





What could  $D^*$  be?



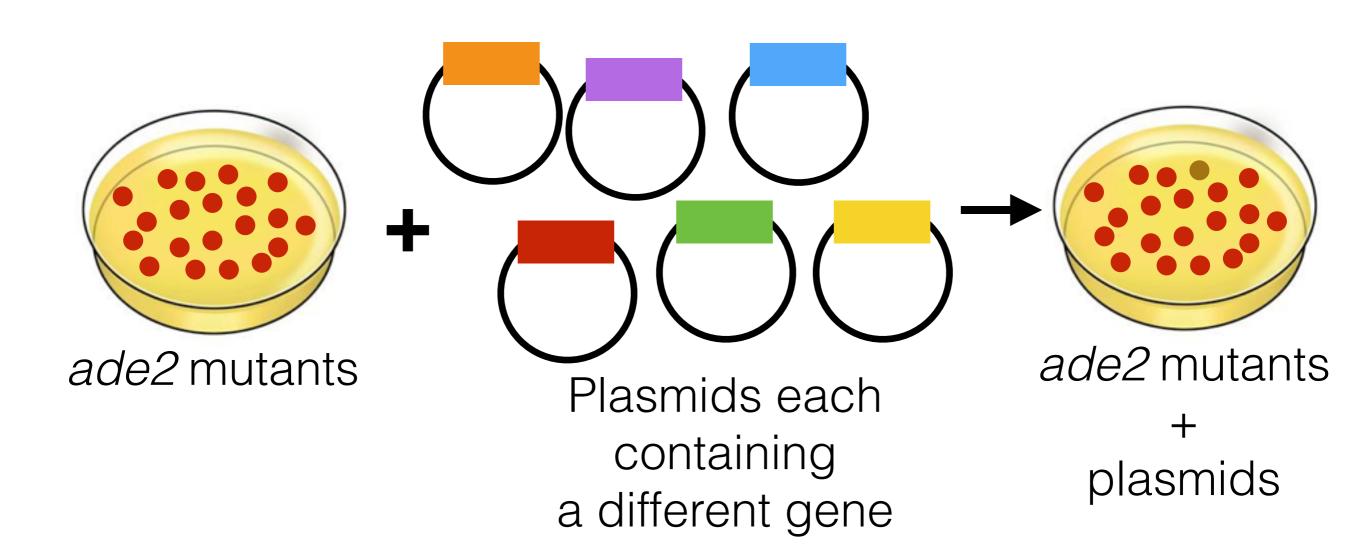
What could  $D^*$  be?

Revertant, extragenic suppressor, intragenic suppressor, or null mutant

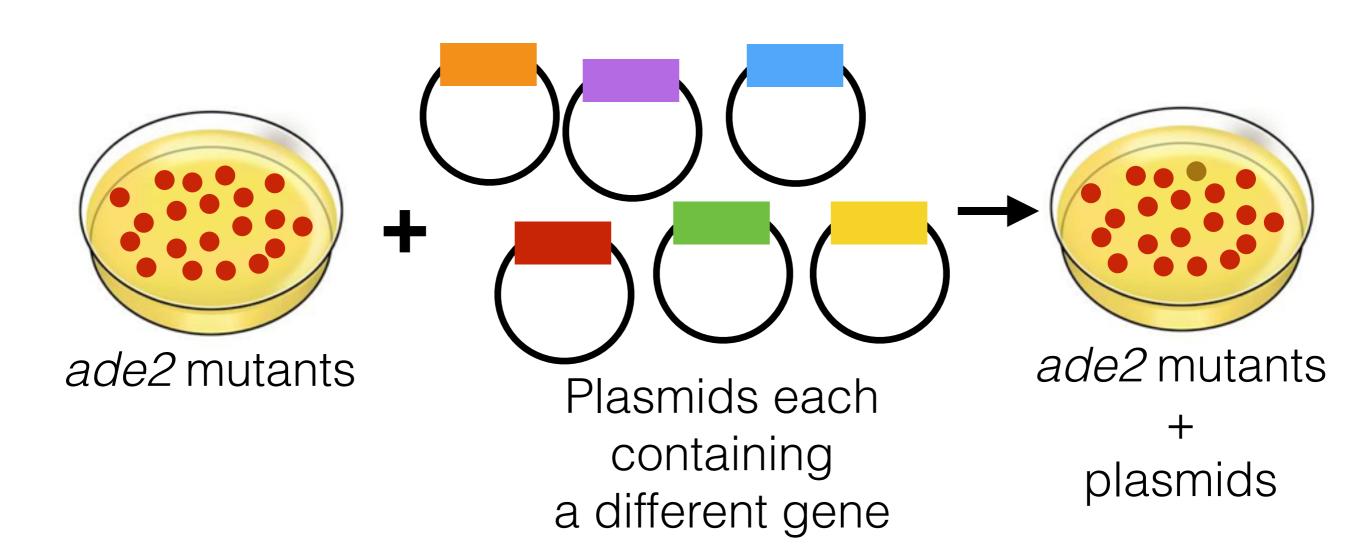
## 15. Clone the gene

- 1. Clone by complementation
- 2. Clone by phenocopy
- 3. Clone by sequencing

# Cloning by complementation in bacteria and yeast

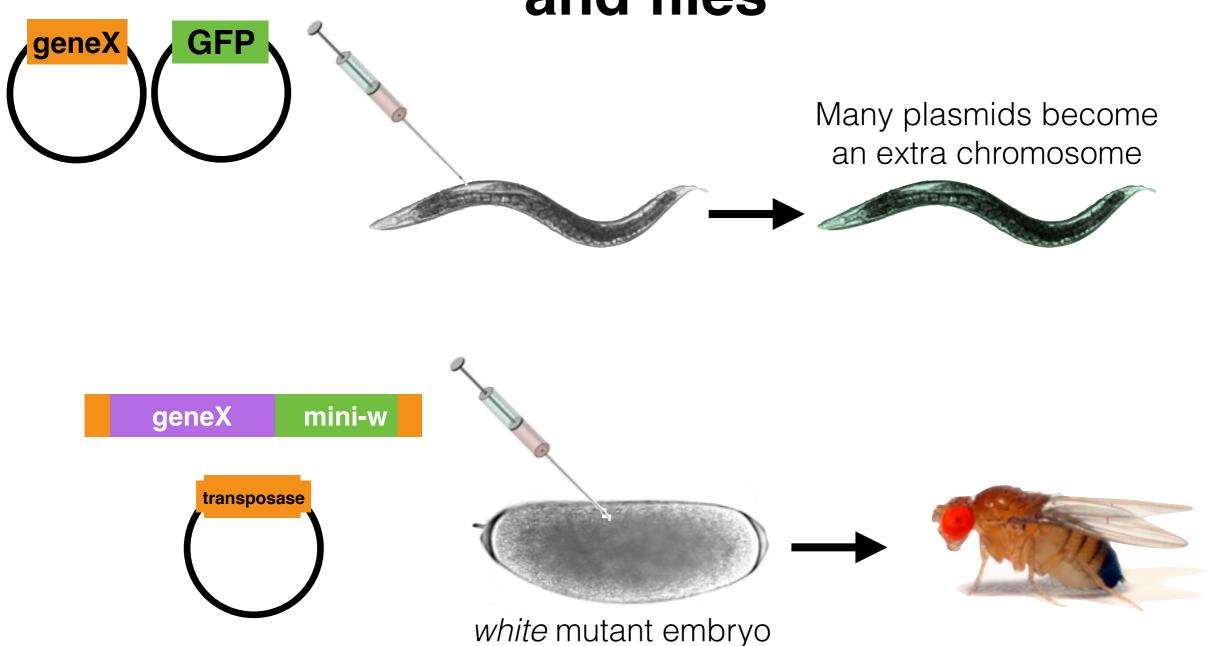


# Cloning by complementation in bacteria and yeast



Caveat: overexpression bypass suppressors

Cloning by complementation in worms and flies

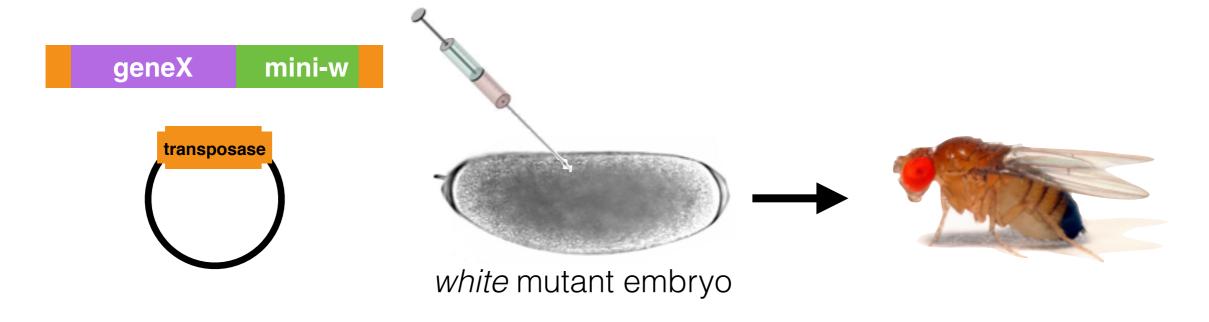


#### Transgenesis and rescue

## Cloning by complementation in worms and flies



Caveat: overexpression bypass suppressors and not stable

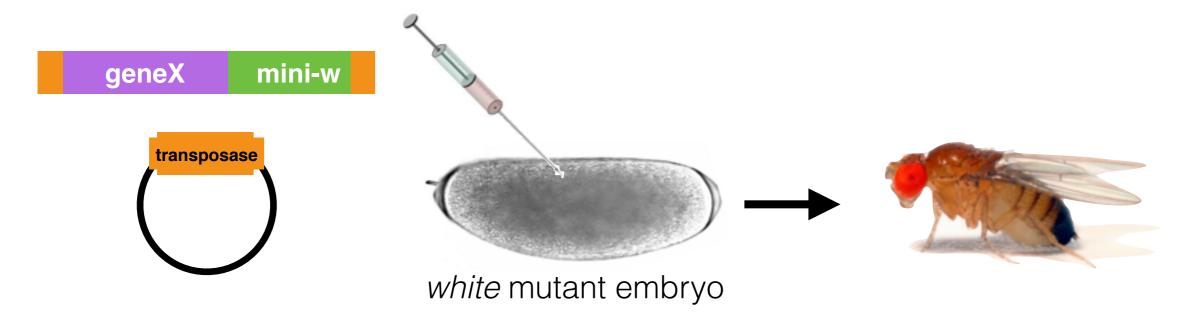


#### Transgenesis and rescue

## Cloning by complementation in worms and flies



Caveat: overexpression bypass suppressors and not stable



Caveat: overexpression bypass suppressor and variable expression

#### Transgenesis and rescue