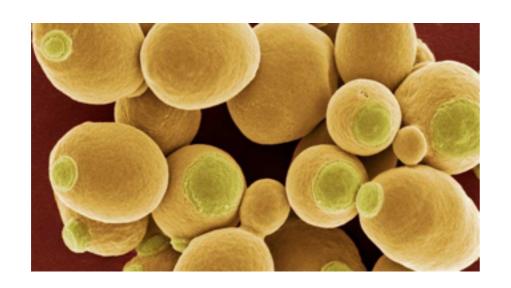
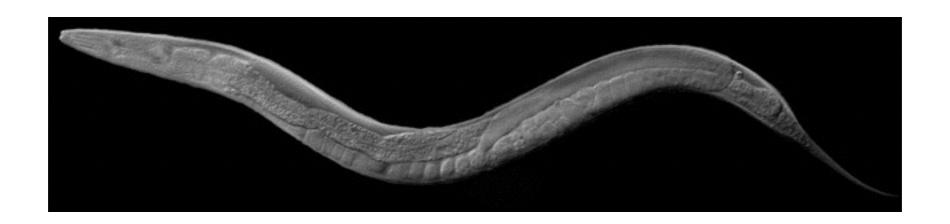
Bio393: Genetic Analysis

Screens, selections, mutants, dosage







Where do all those mutant strains come from?

Natural

- Made by random errors of DNA repair, replication, transcription, recombination, etc.
- Made by natural mutagens (UV, etc.)
- Variants present in a population
- Rare or common

Induced

Made by mutagens (EMS, ENU, X-ray irradiation, etc.)

Where do all those mutant strains come from?

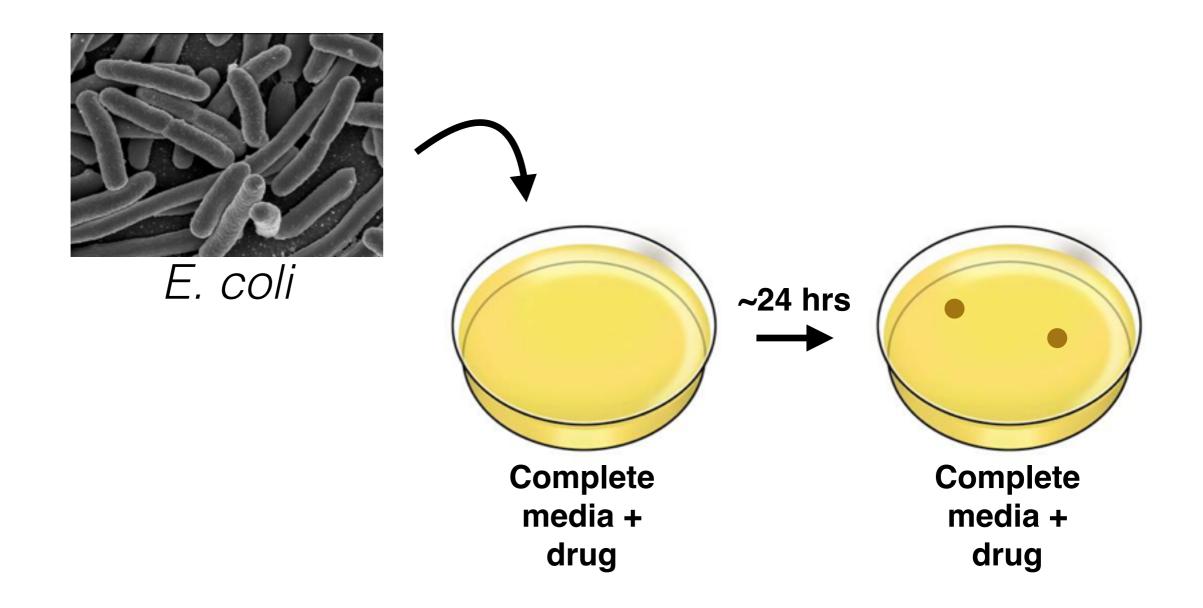
Natural

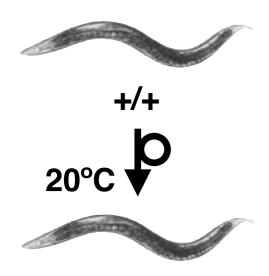
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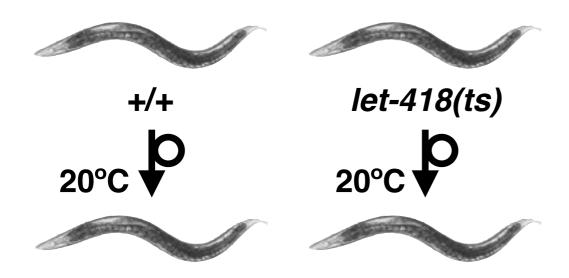
Induced

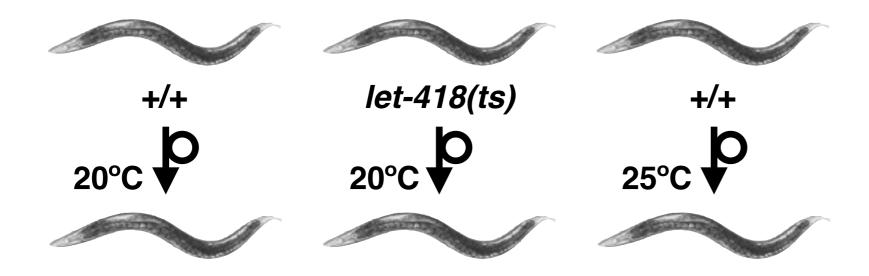
Made by mutagens (EMS, ENU, X-ray irradiation, etc.)

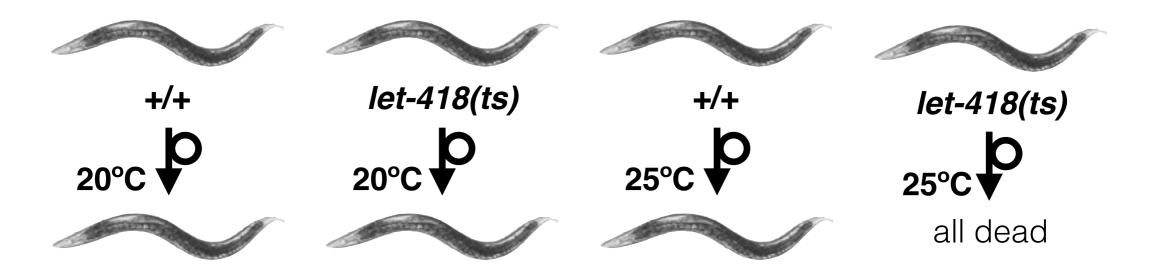
Genomes are full of mutations

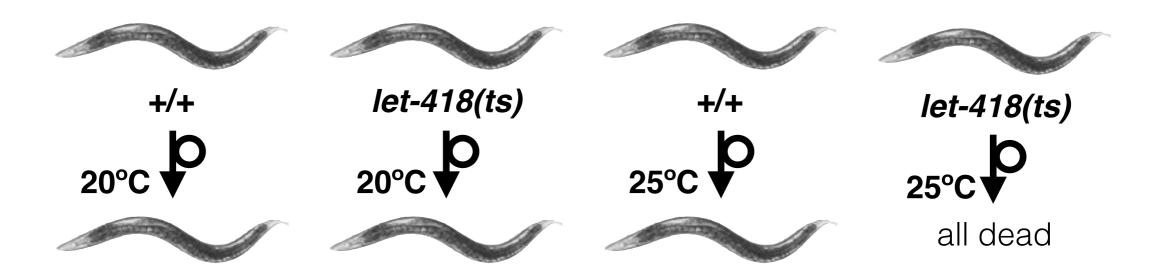


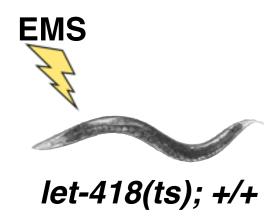


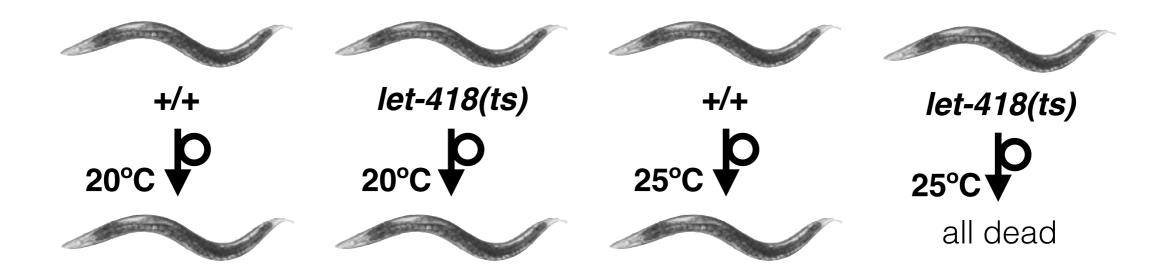


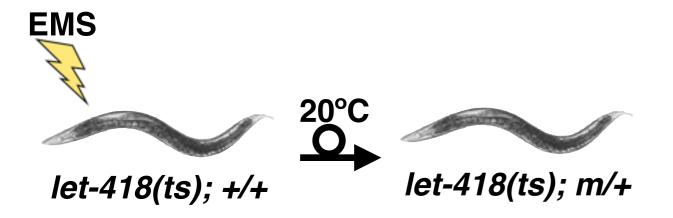


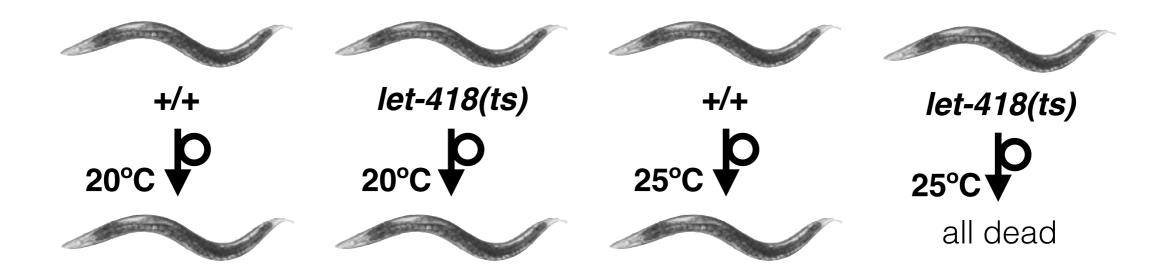


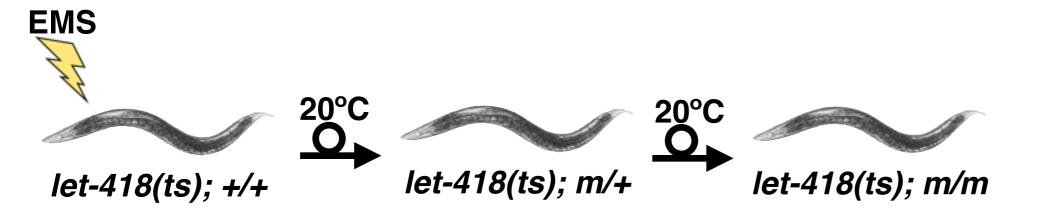


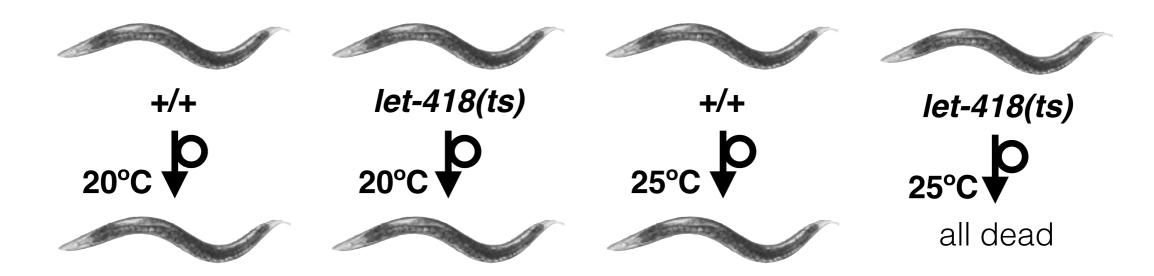


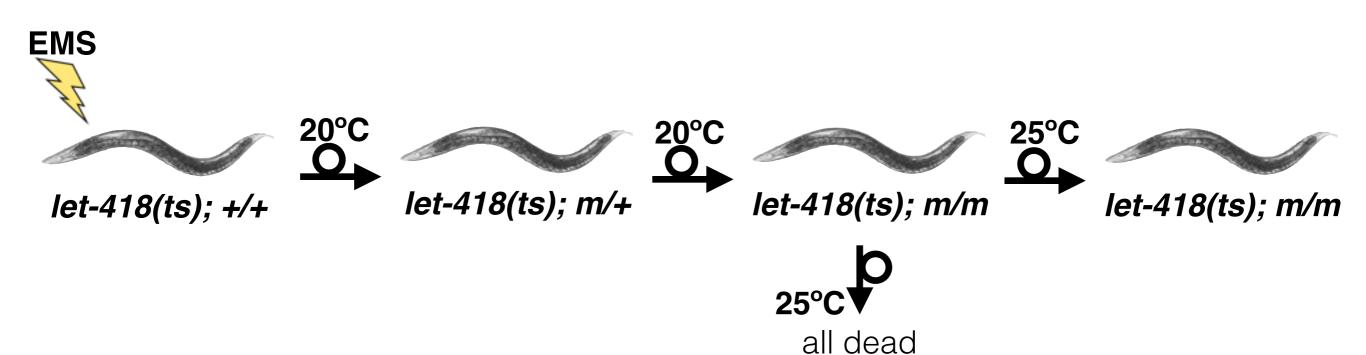


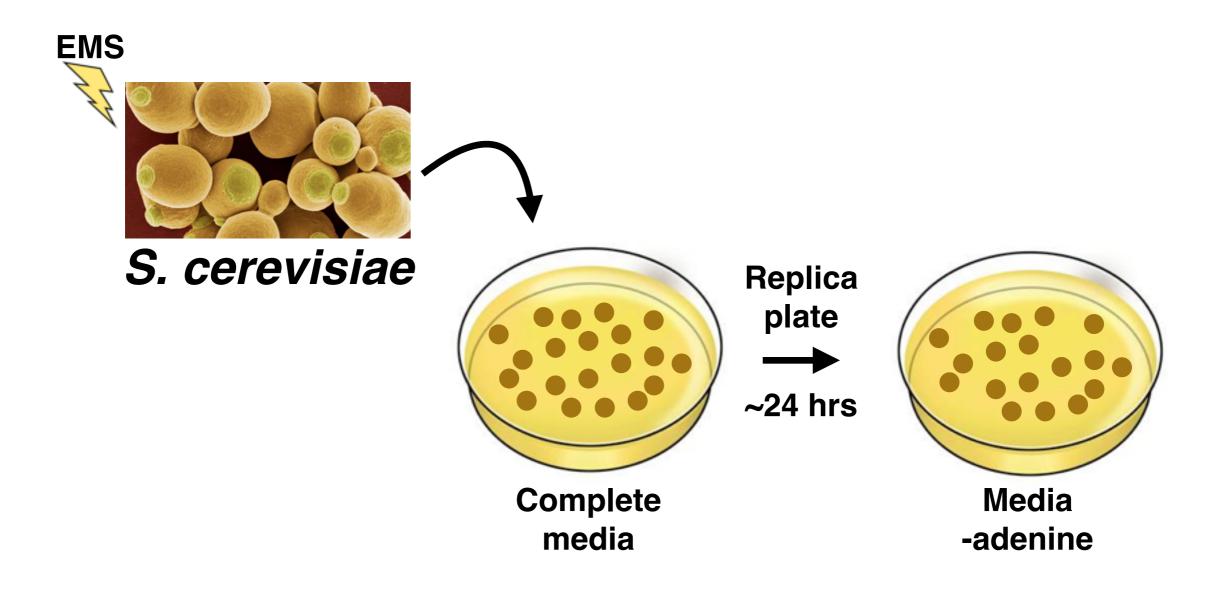


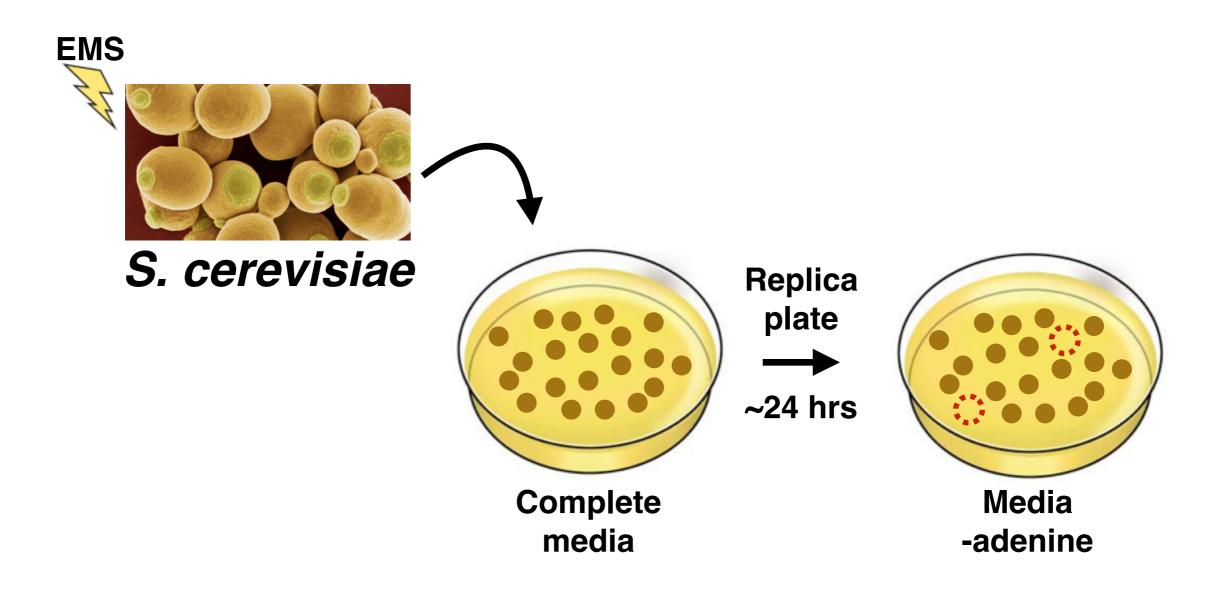


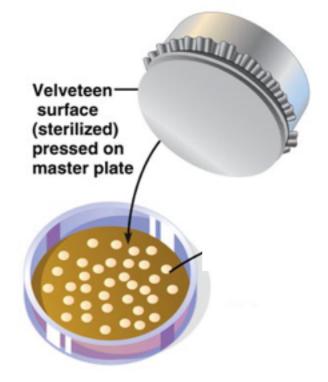


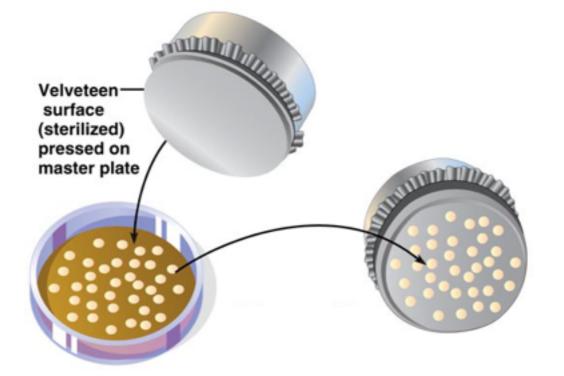


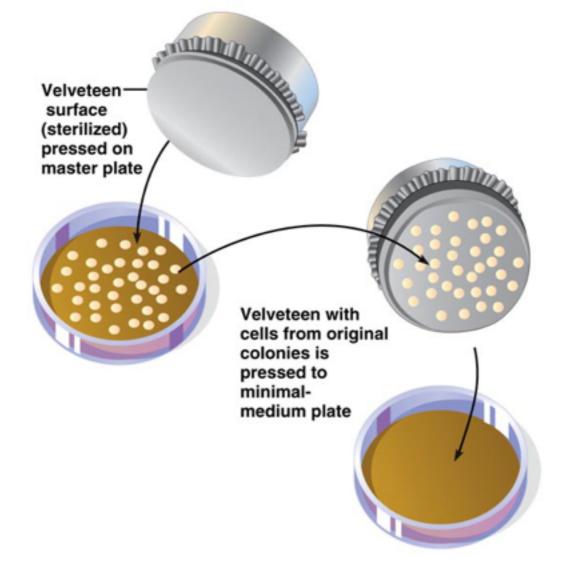


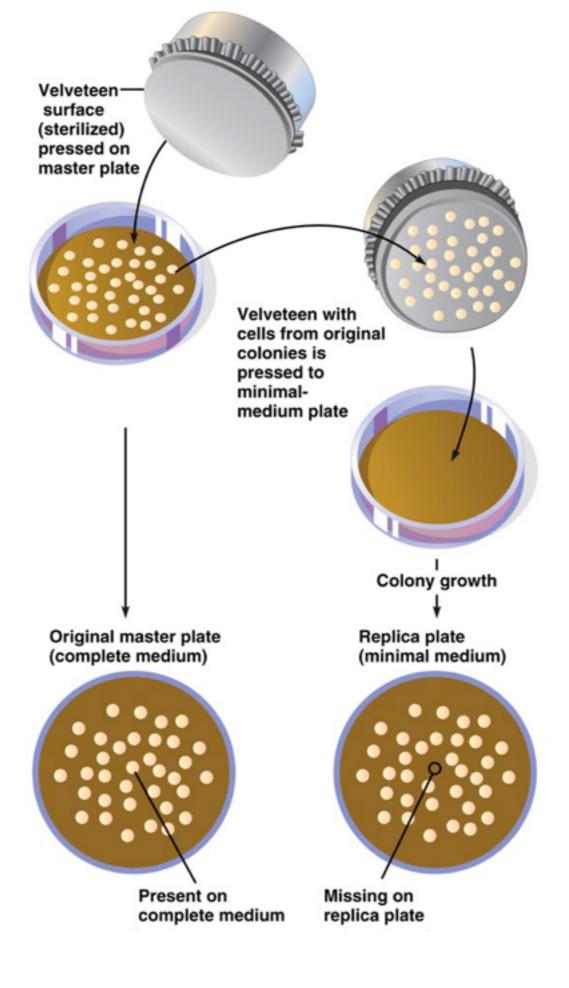










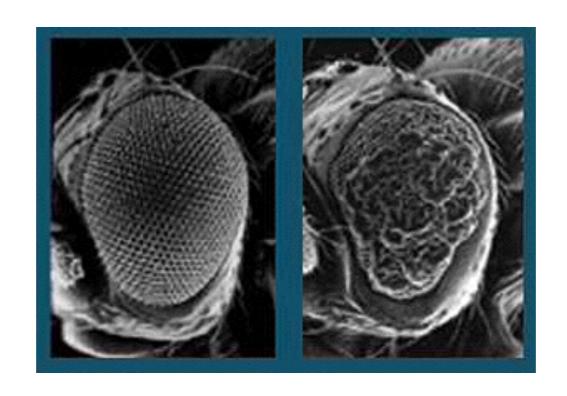




D. melanogaster



D. melanogaster



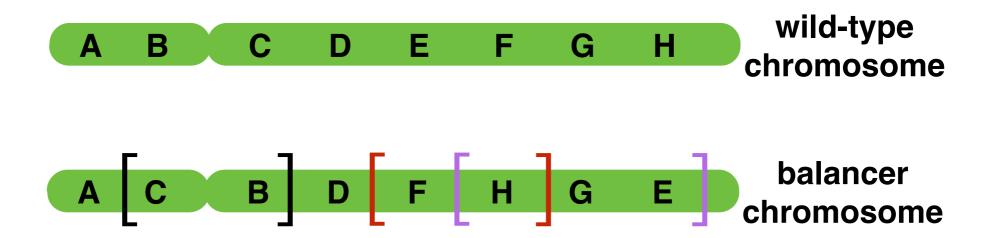
A B C D E F G H chromosome





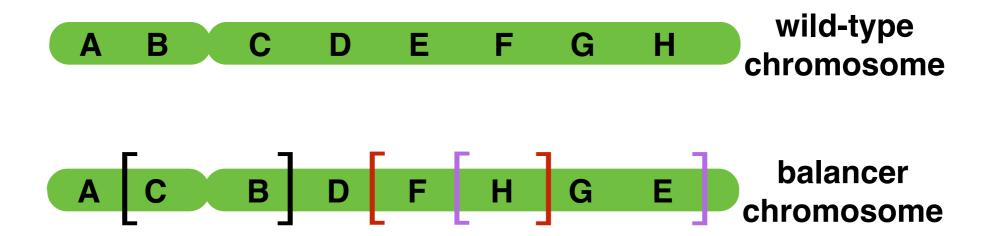
Every balancer chromosome:

1. has many inversions to eliminate recombination

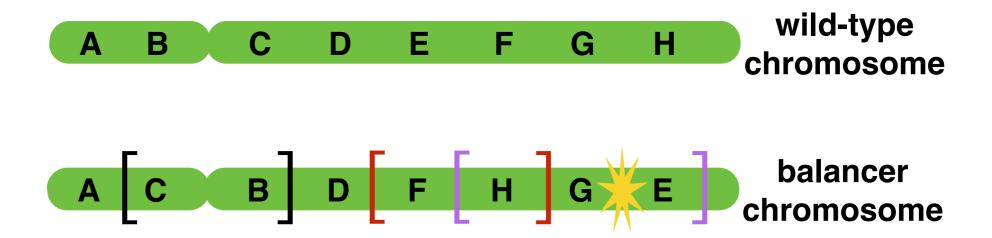


Every balancer chromosome:

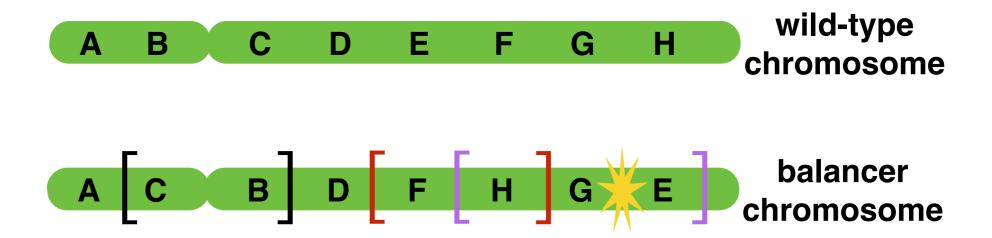
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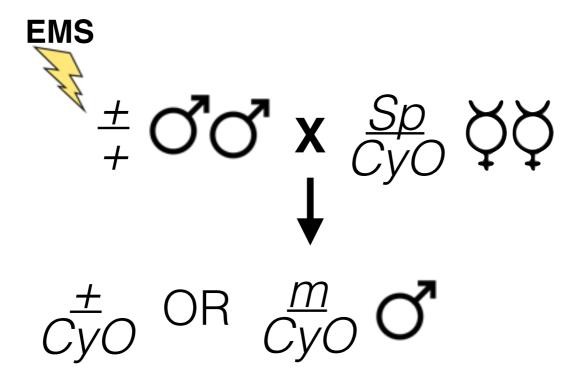


- 1. has many inversions to eliminate recombination
- 2. confers an easily scored dominant phenotype
- 3. is recessive lethal



<u>Sp</u> CyO

EMS
$$\pm \sigma \sigma x \frac{Sp}{CyO} \circlearrowleft$$



Screen: $\frac{\pm}{+}$ $\frac{d}{d}$ \mathbf{x} $\frac{Sp}{CyO}$ $\mathbf{\nabla}$ C_{yO}^{\pm} OR C_{yO}^{m} O' X_{CyO}^{∞} Single-pair crosses

Screen: $\frac{\pm}{+}$ $\frac{d}{d}$ \mathbf{x} $\frac{Sp}{CyO}$ $\mathbf{\nabla}$ C_{yO}^{\pm} OR C_{yO}^{m} O' X_{CyO}^{∞} Single-pair crosses C_{VO} \mathbf{X} C_{VO} $\mathbf{\hat{Q}}$

Screen: **EMS** $\frac{\pm}{+}$ $\frac{d}{d}$ \mathbf{x} $\frac{Sp}{CyO}$ $\mathbf{\nabla}$ C_{yO}^{\pm} OR C_{yO}^{m} O' X_{CyO}^{∞} C_{yO}^{∞} Single-pair crosses C_{yO}^{m} C_{yO}^{m} C_{yO}^{m} dead **Mutant?** Curly wings

How would you screen or select for mutants that cause a dominant or recessive phenotype in yeast, *C. elegans*, *Drosophila*, and mice?

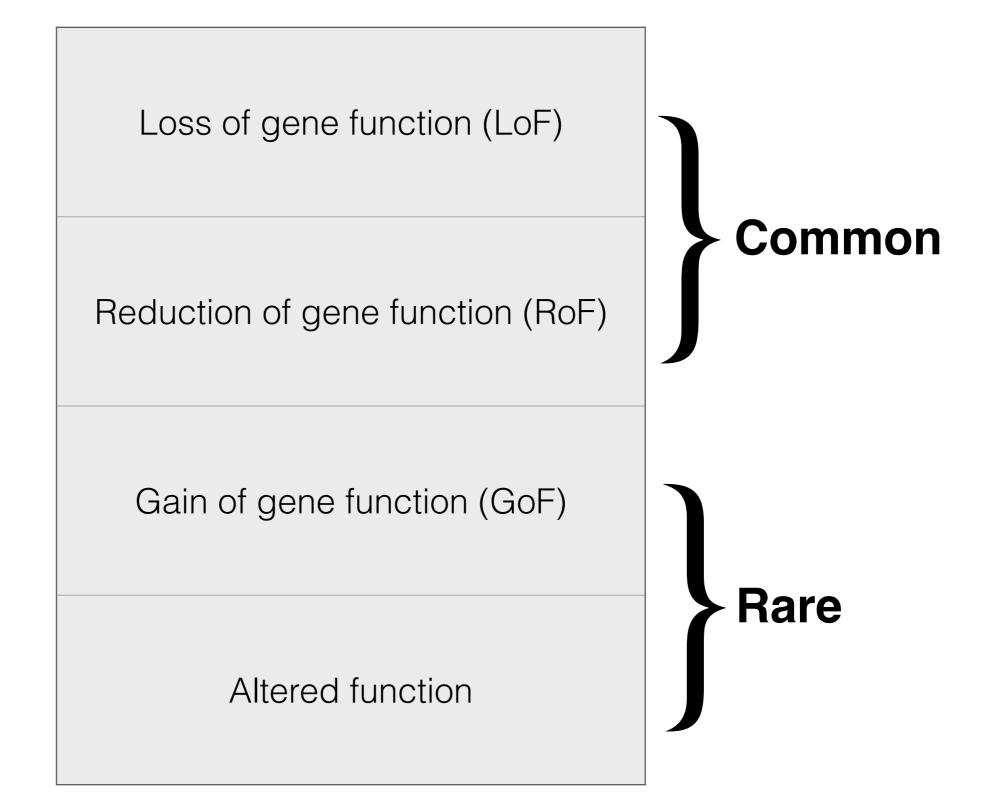
What does a mutation do to gene function?

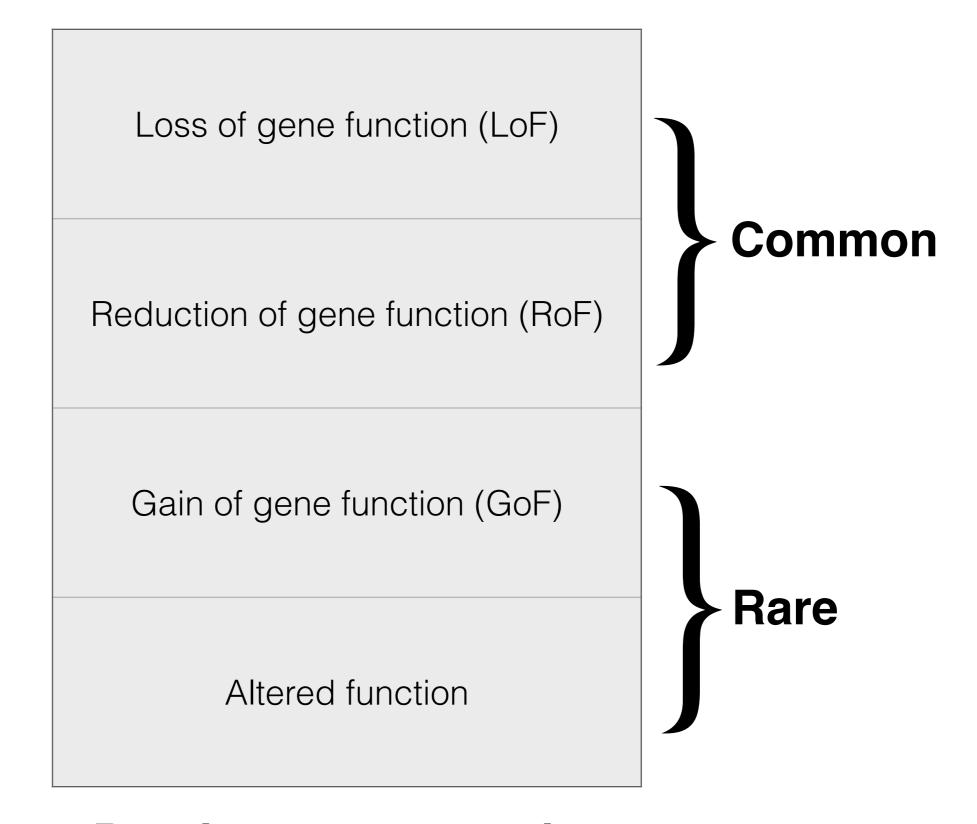
Loss of gene function (LoF)

Reduction of gene function (RoF)

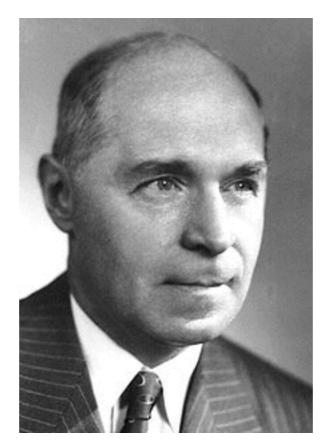
Gain of gene function (GoF)

Altered function





Dominant or recessive correlates with mutation type most times



Hermann Muller



Muller's morphs - gene dosage tests

Loss of gene function (LoF)	amorph, nullomorph
Reduction of gene function (RoF)	hypomorph
Gain of gene function (GoF)	hypermorph
Altered function	neomorph, antimorph

```
m = mutation of gene
```

```
\triangle = deletion of gene
```

+ = normal copy of gene

```
= Phenotype is equivalent
```

```
> = Phenotype is more mutant than
```

< = Phenotype is less mutant than</pre>