

## CONCEPT: TRI-HYBRID CROSS FOR MAPPING

- A tri-hybrid cross, is a mating to look at the inheritance of \_\_\_\_\_ traits in the offspring
  - For unlinked genes, this should follow normal Mendelian inheritance
  - For linked genes, recombination frequencies in the offspring can be used to map all three gene loci

### EXAMPLE:

In a cross of fruit flies, there are three traits of interest.

1. Eye color:  $v^+$  = red,  $v$  = vermillion (like orange/purpleish),
2. Wing veins:  $cv^+$  = presence of wing crossvein,  $cv$  = absence of crossvein
3. Wing shape:  $ct^+$  = normal wing,  $ct$  = cut wing edges

<b>P:</b>	$v^+/v^+ \cdot cv/cv \cdot ct/ct$	x	$v/v \cdot cv^+/cv^+ \cdot ct^+/ct^+$
	<b>Parental phenotypes:</b> Red, crossveinless, cut wings	x	vermillion, crossvein, normal wing
Gametes:	$v^+ \cdot cv \cdot ct$		$v \cdot cv^+ \cdot ct^+$
<b>F<sub>1</sub> trihybrid</b>	$v^+/v \cdot cv/cv^+ \cdot ct/ct^+$		
F <sub>1</sub> test cross	$v^+/v \cdot cv/cv^+ \cdot ct/ct^+$	x	$v/v \cdot cv/cv \cdot ct/ct$
	<b>F<sub>1</sub> phenotypes:</b> Red, crossvein, normal wing		vermillion, crossveinless, cut wings

	$v^+$	$cv$	$ct$
Parental gametes:	$v$	$cv^+$	$ct^+$

Gametes	Phenotypes	Offspring #	Recomb. $v$ and $cv$	Recomb $v$ and $ct$	Recomb $cv$ and $ct$
$v \cdot cv^+ \cdot ct^+$	Parental	580			
$v^+ \cdot cv \cdot ct$	Parental	592			
$v \cdot cv \cdot ct^+$	Recombinant	45			
$v^+ \cdot cv^+ \cdot ct$	Recombinant	40			
$v \cdot cv \cdot ct$	Recombinant	89			
$v^+ \cdot cv^+ \cdot ct^+$	Recombinant	94			
$v \cdot cv^+ \cdot ct$	Recombinant	3			
$v^+ \cdot cv \cdot ct^+$	Recombinant	5			
	<b>Total</b>	1448	268	191	93

Now, you determine the recombination frequencies:

1. For v and cv:  $RF = 268/1448 = 18.5$
2. For v and ct:  $RF = 191/1448 = 13.2$
3. For cv and ct:  $RF = 93/1448 = 6.4$

What can we determine from these \_\_\_\_\_ frequencies?

1. All three genes are linked, because their RFs are less than 50%
2. The orientation and distances of the gene loci

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□ Why does  $13.2 + 6.4 = 19.6$  and not the 18.5 (RF for v and cv)?

- We should have counted the double cross overs twice, instead of once!

v+	ct	cv
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v	ct+	cv+

Gametes	Phenotypes	Offspring #	Recomb. v and cv
v · ct <sup>+</sup> · cv <sup>+</sup>	Parental	580	
v <sup>+</sup> · ct · cv	Parental	592	
v · ct <sup>+</sup> · cv	Recombinant	45	R
v <sup>+</sup> · ct · cv <sup>+</sup>	Recombinant	40	R
v · ct · cv	Recombinant	89	R
v <sup>+</sup> · ct <sup>+</sup> · cv <sup>+</sup>	Recombinant	94	R
v · ct · cv <sup>+</sup>	Recombinant	3	R x2
v <sup>+</sup> · ct <sup>+</sup> · cv	Recombinant	5	R x2
	<b>Total</b>	1448	284

$$RF = 284/1448 = 19.6\%$$

## PRACTICE

1. The following table shows data from a cross (ABC x abc) examining three genes (a, b, and c). Calculate the recombination frequency for A and B

- a. 20%
- b. 32%
- c. 37%
- d. 9.8%

Genotype	Offspring	A C	A B	B C
A B C	320			
a b c	276			
a B C	145			
A b c	152			
A b C	43			
a B c	34			
A B c	9			
a b C	12			

2. The following table shows data from a cross examining three genes (a, b, and c). Calculate the recombination frequency for A and C

- a. 32%
- b. 37%
- c. 20%
- d. 9.8%

Genotype	Offspring	A C	A B	B C
A B C	320			
a b c	276			
a B C	145			
A b c	152			
A b C	43			
a B c	34			
A B c	9			
a b C	12			

3. The following table shows data from a cross examining three genes (a, b, and c). Calculate the recombination frequency for B and C

- a. 20%
- b. 32%
- c. 37%
- d. 9.8%

Genotype	Offspring	A C	A B	B C
A B C	320			
a b c	276			
a B C	145			
A b c	152			
A b C	43			
a B c	34			
A B c	9			
a b C	12			

4. The following table shows data from a cross examining three genes (a, b, and c). Determine the order of genes

- a. A B C
- b. B C A
- c. C A B

Genotype	Offspring	A C	A B	B C
A B C	320			
a b c	276			
a B C	145			
A b c	152			
A b C	43			
a B c	34			
A B c	9			
a b C	12			