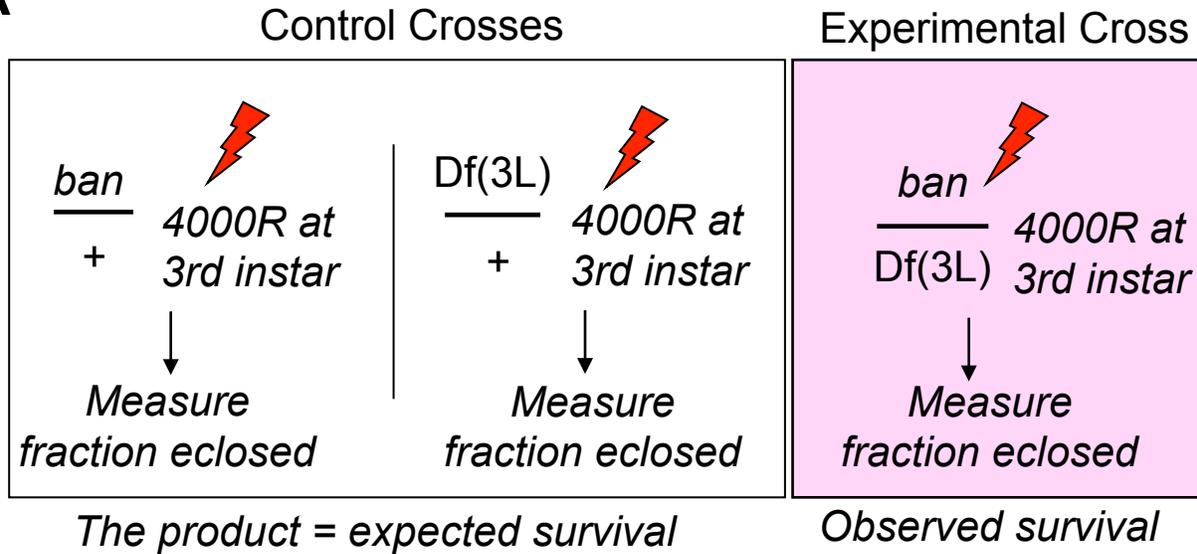
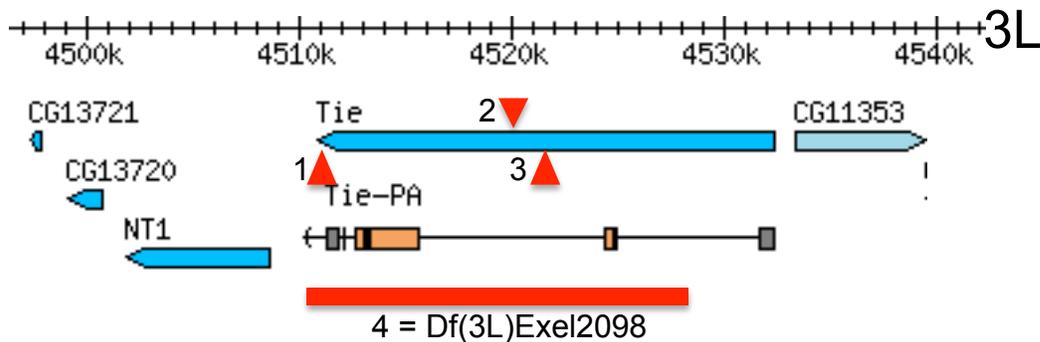


**A**



**B**



- 1 = PBAC{3HPy<sup>+</sup>}Tie<sup>e098</sup>, inserted in predicted 3'UTR
- 2 = PBAC{RB}Tie<sup>e03394</sup>, inserted in the second intron
- 3 = PBAC{RB}Tie<sup>e02680</sup>, inserted in the second intron
- 4 = Df(3L)Exel2098, predicted to remove the entire *tie* coding region and no other gene

**C**

