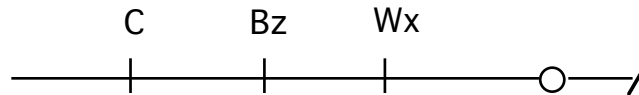


#### Problem set 4: Transposable element answers

1. The map of the left arm of maize chromosome 9 is shown below. C is the colorless gene, which is required for pigmentation of the kernel's aleurone layer, Bz is the bronze gene and is also required for pigmentation of the aleurone, and Wx is the waxy gene, which is required for production of starch in the endosperm of the kernel. The telomere is at the left end of the map and the circle represents the centromere.



In a cross the male parent was  $C^+ Wx^+ / C^+ Wx^+; Ac / Ac$  and the female parent  $c wx / c wx; Ac^+ / Ac^+$ . Remember the nomenclature: Ds and Ac are active dissociation and activator elements and  $Ds^+$  and  $Ac^+$  lack these activities.

a) 50% of the kernels from this cross are variegated for C but not Wx (the aleurone layer contains unpigmented spots and the endosperm is uniformly  $Wx^+$ ), and 50% of the kernels are variegated for both C and Wx (the aleurone layer contains unpigmented spots and the endosperm contains sections that are mutant for Wx). Which chromosomes contain Ds and where do(es) the Ds element(s) map?

b) You find a kernel among the progeny of this cross that you believe contains an unstable allele of C ( $c^u$ ). What phenotype suggests that the kernel contains an unstable allele of C? Can you determine from the phenotype of the kernel containing  $c^u$  if chromosome breakage occurred at this unstable allele? Explain.

2. The following data are from McClintock's maize experiments. She scored the kernel phenotypes that resulted from two crosses. In both crosses the female parent was  $C^+ bz wx / C^+ bz wx$  and the male parent was  $C-I Bz^+ Wx^+ Ds / C^+ Bz^+ wx Ds$ . The male parents in the two crosses were different plants. She scored the phenotype from the two crosses. Below are her results.

### Cross I

| <u>Kernel phenotype<br/>at Wx and C loci</u> | <u>Variegated</u> | <u>Not variegated</u> |
|--|-------------------|-----------------------|
| C-I W <sub>x</sub> <sup>+</sup>              | 268               | 255                   |
| C <sup>+</sup> w <sub>x</sub>                | 248               | 242                   |
| C-I w <sub>x</sub>                           | 88                | 91                    |
| C <sup>+</sup> W <sub>x</sub> <sup>+</sup>   | 84                | 83                    |

### Cross II

| <u>Kernel phenotype<br/>at Wx and C loci</u> | <u>Variegated</u> | <u>Not variegated</u> |
|--|-------------------|-----------------------|
| C-I W <sub>x</sub> <sup>+</sup>              | 928               | 246                   |
| C <sup>+</sup> w <sub>x</sub>                | 164               | 893                   |
| C-I w <sub>x</sub>                           | 62                | 387                   |
| C <sup>+</sup> W <sub>x</sub> <sup>+</sup>   | 344               | 100                   |

- Describe the phenotypes for each marker of each class of kernel.
- In cross I, how do kernels exhibiting the C-I w<sub>x</sub> and C<sup>+</sup> W<sub>x</sub><sup>+</sup> phenotypes arise? What do their frequencies tell you?
- In cross I, why are the frequencies of variegated and not variegated kernels within a phenotypic class approximately equivalent?
- In cross II, why are the frequencies of variegated and not variegated kernels within a phenotypic class not equal. What does the frequency of variegated and not variegated in a phenotypic class tell you?

3. A wild-type *ry*<sup>+</sup> (*rosy*) gene was introduced into a *ry* mutant using P element-mediated gene transformation, and after several additional crosses a strain containing a stable *ry*<sup>+</sup> gene was established.

If a transformed male is mated to an M strain female, would you expect the P element construct used in this experiment to transpose

to another site in the genome of the F1?

If a transformed male is mated to an P strain female, would you expect the P element construct used in this experiment to transpose to another site in the genome of the F1?

4. In a screen for P element mutations in the miniature wings gene on X, you cross a male to a female that has a recessive lethal mutation on one X chromosome. The other chromosome is a balancer X chromosome containing an inversion, a recessive miniature wings mutation, a different recessive lethal mutation and the dominant Bar mutation. How would you have generated the male?

What will be the phenotypes most of progeny from the cross to generate new alleles of miniature wings?

What is the phenotype and genotype of the animals that you will be screening for?

5. You cross two *Drosophila* strains of unknown origin. Strain A females are crossed to Strain B males. The progeny are partially sterile. Two ideas spring to mind—hybrid dysgenesis or a mitochondrial mutation that leads to the gonadal defects.

a. If the cause is hybrid dysgenesis, what type of strains did you cross and what is causing the sterility?

b. If the cause is a mitochondrial mutation, which strain contained the mutation? Why?