

Name:

## Introduction to Genetic Analysis

### Problem Set #1-2

Please answer the following questions on a **SEPARATE SHEET OF PAPER**.

### Problem 1: Complementation Test

You have isolated a collection of yeast mutants that form small colonies when plated on a concentration of the drug geldanamycin that does not affect the growth of wild type cells. Geldanamycin sensitive mutants 1–10 are MAT $\alpha$  and mutants 11–21 are MAT $a$ . As mentioned in lecture, MAT $\alpha$  can only mate with MAT $a$ . Your analysis begins by pairwise mating of each mutant to a wild-type strain and to the mutants of the opposite mating type.

The size of the colonies of the resulting diploids is shown in the table below (“wt” indicates wild-type colony size following geldanamycin treatment, “-” indicates small colony size following treatment):

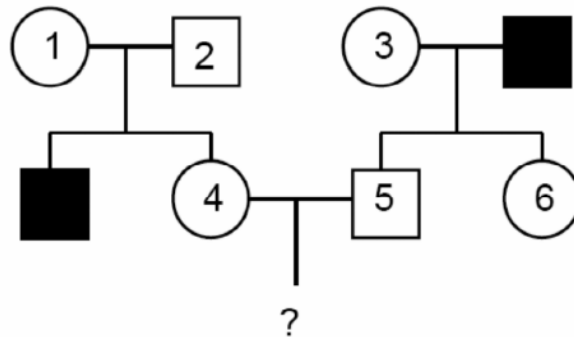
	Wild-type	11	12	13	14	15	16	17	18	19	20	21
Wild-type	wt	-	wt	wt	wt	wt	-	wt	-	-	wt	wt
1	wt	-	wt	wt	wt	wt	-	wt	-	-	wt	-
2	wt	-	wt	-	wt	wt	-	-	-	-	-	wt
3	wt	-	wt	-	wt	wt	-	-	-	-	-	-
4	wt	-	-	wt	-	-	-	wt	-	-	wt	wt
5	wt	-	wt	-	wt	wt	-	-	-	-	-	wt
6	wt	-	wt	-	wt	wt	-	-	-	-	-	wt
7	-	-	-	-	-	-	-	-	-	-	-	-
8	-	-	-	-	-	-	-	-	-	-	-	-
9	wt	-	-	wt	-	-	-	wt	-	-	wt	wt
10	wt	-	-	wt	-	-	-	wt	-	-	wt	wt

- Which of the mutants are dominant and which are recessive?
- What is anomalous about the behavior of Mutant 3? Provide a simple genetic explanation.
- Organize the 21 mutations into complementation groups (genes). Please indicate any remaining ambiguities.

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## **Problem 2: Probability & Bayes' Theorem**

In the following pedigree assume that no new mutations arise. You may further assume that each possible mode of inheritance (autosomal recessive, autosomal dominant, X-linked recessive, X-linked dominant) has equal a prior probability. (Show your work).



- What is the probability that ? is an affected male, assuming an X-linked recessive trait.
- What is the probability that ? is an affected male, assuming an autosomal recessive trait?
- What is the probability that ? is an affected female, assuming an X-linked recessive trait?
- Given that ? is an affected male, what is the probability that this pedigree describes an autosomal recessive trait?
- What is the probability ? will be affected without knowing the sex of the individual?