



**2.** You have isolated two different yeast mutants that will not grow on medium that lacks arginine. You call these mutants  $arg1^-$  and  $arg2^-$ .

**a)** Mating of either  $arg1^-$  or  $arg2^-$  to wild type produces diploids that can grow without arginine. Mating of  $arg1^-$  to  $arg2^-$  produces a diploid that also can grow on medium without arginine. What do these results tell you about the  $arg1^-$  and  $arg2^-$  mutations?

**b)** Sporulation of the diploids produced by the mating of  $arg1^-$  to  $arg2^-$  yields tetrads of the following types:

| <u>type 1</u>                           | <u>type 2</u>      |
|---|--------------------|
| 1 Arg <sup>+</sup> , 3 Arg <sup>-</sup> | 4 Arg <sup>-</sup> |

Out of 20 tetrads, 2 are type 1, and 18 are type 2. Categorize each of the tetrad types as parental ditype (P), tetratype (T), or nonparental ditype (N). If the markers are linked give the distance between them in cM. Does this result tell you anything new about the gene(s) mutated in  $arg1^-$  and  $arg2^-$ ?

**c)** Now you isolate a third mutant called  $arg3^-$ . When this mutant is mated to wild type the resulting diploids cannot grow without arginine. When  $arg3^-$  is mated to  $arg1^-$  the resulting diploid cannot grow without arginine. What do these results tell you about  $arg3^-$  and its relationship to  $arg1^-$ ?

**d)** When the diploid produced by mating  $arg3^-$  to  $arg1^-$  is sporulated tetrads of three types are produced.

| <u>type 1</u>                           | <u>type 2</u>     | <u>type 3</u>                           |
|---|-------------------|---|
| 1 Arg <sup>+</sup> , 3 Arg <sup>-</sup> | 4Arg <sup>-</sup> | 2 Arg <sup>+</sup> , 2 Arg <sup>-</sup> |

Out of 20 tetrads, 12 are type 1, 5 are type 2, and 3 are type 3. Categorize each of the tetrad types as parental ditype (P), tetratype (T), or nonparental ditype (N). What does this result tell you about the relationship between  $arg3^-$  and  $arg1^-$ ? Give any relevant distances in cM.

**e)** Based on these results deduce the relationship between  $arg3^-$  and  $arg2^-$ . In a mating of  $arg3^-$  to  $arg2^-$  what would the phenotype of the resulting diploid be? If this diploid were sporulated and 60 tetrads were examined how many tetrads of each type would you expect?

**3.** You have isolated two temperature-sensitive mutations in phage  $\phi$ . These phage mutants are called **ts-1** and **ts-2**. Each mutant will form plaques at 30°C but not at 42°C. You cross **ts-1** to **ts-2** phage by coinfecting *E. coli* at the permissive temperature of 30°C. When the resulting phage lysate is plated at 30°C you count  $10^5$  plaques per ml of phage lysate, but when the same phage lysate is plated at 42°C, there are 300 plaques per ml.

(a) What is the distance between the **ts-1** and **ts-2** mutations in m.u.?

You next cross a **ts-1** phage strain to a **ts-2** phage strain that also carries a **cl<sup>-</sup>** mutation, which gives plaques with clear centers. When the resulting lysate is plated out at 42°C and 100 plaques are examined, 85 are clear and 15 have normal turbid centers.

(b) If the phage produced from this cross were plated at 30°C, what fraction of the plaques would you expect to have clear centers?

(c) Draw a map showing the relative order of **cl<sup>-</sup>**, **ts-1** and **ts-2** as well as all of the distances you can calculate in m.u.