

Chromosome mapping

Your goal is to map the position of several genes A-K on their respective chromosomes.

Information you are given:

The ratios of phenotypes obtained when crossing heterozygotic individuals (obtained from crossing homozygotic pure breeding recessive or dominant for both genes analyzed).

Take in account that:

The expected ratios for independent heredity are 9:3:3:1

The expected ratio for linked heredity (virtually, at the same place) are: 3:1, it is 12:4

And rates can range from these two edges. For example:

When genes are in the same chromosome, but separated, rates about 10:2:2:2 could be expected

When genes are near in the same chromosome, you could expect rates about 11:1:1:2

Information you are not given:

The total number of chromosomes (but you can discover it from your data).

Process:

Every team can ask for a crossing in each round.

Each team will be the expert on one gene, and in this first step will analyze the data from this gene. When ready, you can choose one of the other teams (you will be able to know which are the speciality of the other teams). And work together to complete your work.

Answer:

Chromosome 1: A.....DB

Chromosome 2: JC

Chromosome 3: E...I...F...K

Chromosome 4: H.....G