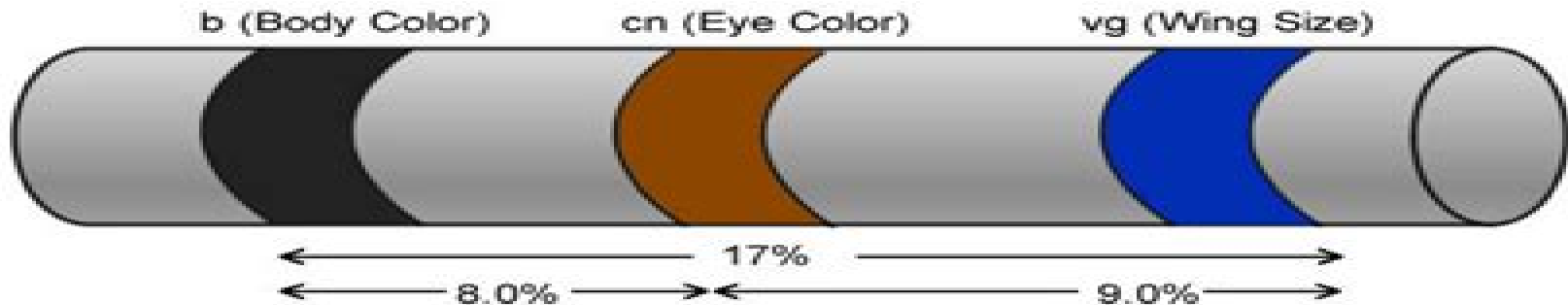


How to Create a Chromosome Map from Crossover Frequencies

Recombination: During crossing-over (prophase I of Meiosis), genes on chromosomes switch places. Crossover is random, but the likelihood that 2 genes crossover will increase if those genes are farther apart. Genes closer together are more likely to "stick together" and not switch places.

Gene Linkage Maps: Using the crossover frequencies, you can construct a map to represent the distances between genes.

This map shows chromosome #2 of *Drosophila melanogaster*. The distance between the genes can be written as a percentage or as a MAP UNIT. The gene for body color and wing size are 17 map units apart.



Sample Problem:

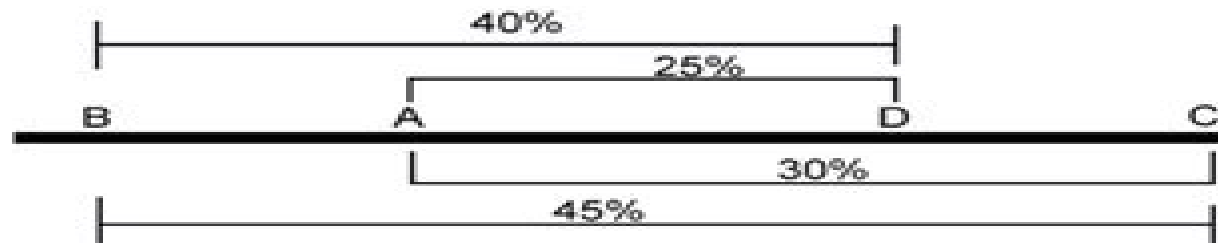
Given the crossover frequency of each of the genes on the chart, construct a chromosome map.

Gene	Frequency of Crossover
A-C	30%
B-C	45%
B-D	40%
A-D	25%

Step 1: Start with the genes that are the farthest apart first: B and C are 45 map units apart and would be placed far apart.

B 45% C

Step 2: Solve it like a puzzle, using a pencil to determine the positions of the other genes.



Step 3: Subtraction will be necessary to determine the final distances between each gene.

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Harcourt Science: Teacher's ed., life science units A and B ,2005 *Automated Design and Optimization of Wire Antennas Using Genetic Algorithms* Derek Scott Linden,1997 A Genetic Algorithm GA has been used in conjunction with the Numerical Electromagnetics Code Version 2 NEC2 to create and optimize atypical wire antenna designs with impressive characteristics Antenna design parameters are encoded into an ordered series of numbers and or symbols analogous to a biological chromosome A cost function that quantifies how well a design meets the engineer s specifications is created The GA uses these to generate and evaluate a population of designs The most successful designs are then promoted and mixed through mating and mutation while poor designs are removed This process difficult to trap in local minima continues until convergence criteria are met generally yielding excellent designs with no user intervention or initial guesses Three antennas have been optimized a monopole loaded with a modified folded dipole the Yagi antenna and the crooked wire genetic antenna Prior study of the loaded monopole had shown hemispherical coverage was possible The GA found an asymmetric loaded monopole with an average variation in gain over the hemisphere of only 0.4dB confirmed by measurement GA optimized Yagi antennas surpassed the gain of conventional Yagis by about 1dB improvement also confirmed by measurement The GA designed a Yagi with a beamwidth of 50 deg 60 deg sidelobes nearly 25dB down and a 14% bandwidth specifications difficult to achieve using conventional techniques The crooked wire genetic antenna is several wires joined in series locations and lengths are determined by the GA Optimization for hemispherical coverage with right hand circular polarization RHCP produced highly unusual shapes unrealizable using a conventional approach RHCP hemispherical coverage was achieved with less than 4dB variation Measurements verify the results

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