

GENE MAPPING IN LABORATORY MAMMALS PART B%0A

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Gene mapping in laboratory mammals, Part A, Part B, Part A is a comprehensive account of statistical methods for detecting and estimating linkage, and for devising linkage maps. It covers not only all types of segregation, from the various two-point crosses to the multi-locus testcross, but also segregations with deviations from the classic Mendel-based expectations - those disturbed by partial

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Part A presents a general account of statistical methods which can be brought to bear on the problem. The primary emphasis is on the practical aspects of estimation. A large proportion, if not The primary emphasis is on the practical aspects of estimation.

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