

Probability Models and
Statistical Methods in
Genetics (Wiley series
in probability &
mathematical statistics)

Johnson, Regina C. Elandt-

Probability Models And Statistical Methods In Genetics

Didier Musso



Probability Models And Statistical Methods In Genetics:

Probability Models and Statistical Methods in Genetics Regina C. Elandt-Johnson,1971 Probability Models and Statistical Methods in Genetics Regina C. Elandt-Johnson,1971 Basic terms and definitions in genetics Probability models general concepts and definitions Genotypes and phenotypes in experimental populations Equilibrium laws in panmictic populations Random variables and their distributions Moments of random variables Genotype distributions for relatives in Randomly mating populations Some standard distributions and their properties Inbreeding and nonrandom mating Natural selection and mutation Estimation of genetic parameters general theory Estimation of parameters associated with multinomial distributions Testing genetic hypotheses Human blood groups Autosomal linkage in experimental populations Statistically equivalent models of Inheritance Segregation ratios in families Simple modes of inheritance Complex segregation analysis Histocompatibility testing *Statistics in Genetics* M.Elizabeth Halloran,Seymour Geisser,1999-06-04 Though the Genome Project will eventually result in the sequencing of the human genome as well as the genomes of several other organisms there will still be a need for good statistics for family studies of complex diseases The papers in this volume are contributions by some of the leading researchers in the field to the current topics in statistical genetics One section deals with DNA sequence matching and issues related to forensics while another deals with statistical problems of modeling phylogenies and inferential difficulties related to the complex tree structures produced as well as the method of coalescence

Genetic Variation and Human Disease Kenneth M. Weiss,1993 Recent developments in molecular and computational methods have made it possible to identify the genetic basis of any biological trait and have led to spectacular advances in the study of human disease This book provides an overview of the concepts and methods needed to understand the genetic basis of biological traits including disease in humans Using examples of qualitative and quantitative phenotypes Professor Weiss shows how genetic variation may be quantified and how relationships between genotype and phenotype may be inferred This book will appeal to many biologists and biological anthropologists interested in the genetic basis of biological traits as well as to epidemiologists biomedical scientists human geneticists and molecular biologists **Probability Models for DNA Sequence Evolution** Richard Durrett,2008-12-15 Our basic question is Given a collection of DNA sequences what underlying forces are responsible for the observed patterns of variability To approach this question we introduce and analyze a number of probability models the Wright Fisher model the coalescent the infinite alleles model and the infinite sites model We study the complications that come from nonconstant population size recombination population subdivision and three forms of natural selection directional selection balancing selection and background selection These theoretical results set the stage for the investigation of various statistical tests to detect departures from neutral evolution The final chapter studies the evolution of whole genomes by chromosomal inversions reciprocal translocations and genome duplication Throughout the book the theory is developed in close connection with data from more than 60 experimental studies from the biology

literature that illustrate the use of these results This book is written for mathematicians and for biologists alike We assume no previous knowledge of concepts from biology and only a basic knowledge of probability a one semester undergraduate course and some familiarity with Markov chains and Poisson processes *Journal of the American Statistical Association*, 1972 Genes in Populations Eliot B. Spiess, 1977 *Likelihood, Bayesian, and MCMC Methods in Quantitative Genetics* Daniel Sorensen, Daniel Gianola, 2007-03-22 This book suitable for numerate biologists and for applied statisticians provides the foundations of likelihood Bayesian and MCMC methods in the context of genetic analysis of quantitative traits Although a number of excellent texts in these areas have become available in recent years the basic ideas and tools are typically described in a technically demanding style and contain much more detail than necessary Here an effort has been made to relate biological to statistical parameters throughout and the book includes extensive examples that illustrate the developing argument *Genetika*, 1992 Mathematical and Statistical Methods for Genetic Analysis Kenneth Lange, 2012-12-06 During the past decade geneticists have cloned scores of Mendelian disease genes and constructed a rough draft of the entire human genome The unprecedented insights into human disease and evolution offered by mapping cloning and sequencing will transform medicine and agriculture This revolution depends vitally on the contributions of applied mathematicians statisticians and computer scientists *Mathematical and Statistical Methods for Genetic Analysis* is written to equip students in the mathematical sciences to understand and model the epidemiological and experimental data encountered in genetics research Mathematical statistical and computational principles relevant to this task are developed hand in hand with applications to population genetics gene mapping risk prediction testing of epidemiological hypotheses molecular evolution and DNA sequence analysis Many specialized topics are covered that are currently accessible only in journal articles This second edition expands the original edition by over 100 pages and includes new material on DNA sequence analysis diffusion processes binding domain identification Bayesian estimation of haplotype frequencies case control association studies the gamete competition model QTL mapping and factor analysis the Lander Green Kruglyak algorithm of pedigree analysis and codon and rate variation models in molecular phylogeny Sprinkled throughout the chapters are many new problems **Simulation and the Monte Carlo Method** Reuven Y. Rubinstein, 1981-05-14 This book provides the first simultaneous coverage of the statistical aspects of simulation and Monte Carlo methods their commonalities and their differences for the solution of a wide spectrum of engineering and scientific problems It contains standard material usually considered in Monte Carlo simulation as well as new material such as variance reduction techniques regenerative simulation and Monte Carlo optimization *Manual of Quantitative Genetics* Walter Alvin Becker, 1992 **Genetic Basis of the Epilepsies** Victor Elving Anderson, 1982 Analysis of Human Genetic Linkage Jurg Ott, 1991 For the last decade this book has been the unrivaled standard text on human genetic linkage analysis Biometrics One of the best books that cover the theory behind linkage analysis Scientists with an interest in genetic linkage

analysis should definitely consider allocating space on their bookshelves for this book Biotech Software and Internet Report A valuable resource It is the most accessible work on the subject and should be in the collection of everyone interested in mapping human genes Doody s Book Review Service

The Genetic Basis of Common Diseases Richard Allen King, Jerome I. Rotter, 1992 Here is a unique and comprehensive resource for human geneticists and physicians concerned with the genetic causes of diseases routinely seen in research clinical practice and genetic counseling Each of the chapters covers a particular disease describes the genetic factors involved and define the relevant biochemical immunological and physiological markers The expert widely known contributors also detail clinical applications with advice on how the genetic data can be used to evaluate individuals and families interpret diagnostic texts and manage the disease

Animal Breeding Abstracts ,1984 *General Catalogue* University of California, Los Angeles,1975 **Prior Information in Linear Models** Helge Toutenburg,1982 Cumulated Index Medicus ,1989 **Multilocus Genetic Structure of Natural Populations** **Lodgepole Pine** Bryan Keith Epperson,1983

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