

Lecture Notes in Biomathematics

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15

Thomas Nagylaki

Selection in One-
and Two-Locus Systems

With a Foreword by R. M. May



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Selection In One And Two Locus Systems Lecture Notes In Biomathematics Volume 15

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Selection In One And Two Locus Systems Lecture Notes In Biomathematics Volume 15:

Vito Volterra Symposium on Mathematical Models in Biology Claudio Barigozzi, 2013-03-13 The idea of organizing a symposium on mathematical models in biology came to some colleagues members of the Accademia dei Lincei in order to point out the importance of mathematics not only for supplying instruments for the elaboration and the evaluation of experimental data but also for discussing the possibility of developing mathematical formulations of biological problems This appeared particularly appropriate for genetics where mathematical models have been of historical importance When the organizing work had started it became clear to us that the classic studies of Vito Volterra who was also a Member of the Academy and its President from 1923 to 1926 might be considered a further reason to have the meeting in Rome at the Accademia dei Lincei thus the meeting is dedicated to his memory Biology in its manifold aspects proved to be a difficult object for an exhaustive approach thus it became necessary for practical reasons to make a choice of problems Therefore not all branches of biology have been represented The proceedings of the symposium as a whole assume a knowledge of mathematics on the part of the reader however the problem of teaching mathematics to biologists was the subject of a round table discussion not recorded in these proceedings On this were brought up some basic points to be recommended to teachers on an international basis and a statement was prepared for circulation The Organizing Committee

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TOPIC I MODELS OF NATURAL SELECTION

Analysis of Neural Networks U. van der Heiden, 2013-03-08 The purpose of this work is a unified and general treatment of activity in neural networks from a mathematical point of view Possible applications of the theory presented are indicated throughout the text However they are not explored in detail for two reasons first the universal character of neural activity in nearly all animals requires some type of a general approach secondly the mathematical perspicuity would suffer if too many experimental details and empirical peculiarities were interspersed among the mathematical investigation A guide to many applications is supplied by the references concerning a variety of specific issues Of course the theory does not aim at covering all individual problems Moreover there are other approaches to neural network theory see e.g. Poggio and Torre 1978 based on the different levels at which the nervous system may be viewed The theory is a deterministic one reflecting the average behavior of neurons or neuron pools In this respect the essay is written in the spirit of the work of Cowan, Feldman and Wilson see sect 2.2 The networks are described by systems of nonlinear integral equations Therefore the paper can also be read as a course in nonlinear system theory The interpretation of the elements as neurons is not a necessary one However for vividness the mathematical results are often expressed in neurophysiological terms such as excitation, inhibition, membrane potentials and impulse frequencies The nonlinearities are essential constituents of the theory

Population Biology and Evolution K. Wöhrmann, V. Lösckke, 2012-12-06 This volume contains the papers presented at a symposium on population biology sponsored by the Deutsche Forschungsgemeinschaft It was held at the guest house of the University of Tübingen at Oberjoch on May 15-19

1983 Prior to this conference a small group of European biologists had met in Berlin June 1981 and Pavia September 1982 to discuss research problems on the borderline between population genetics and evolutionary ecology From the contributions and discussions at these meetings it became evident that the unification of approaches to evolutionary problems in population genetics and evolutionary ecology has not yet been successful and requires further efforts It was the consensus that a larger symposium with international participation would be helpful to confront and discuss the different approaches to population biology in order to assess where we are now and where we should be going As a result an organizational committee was formed F Christiansen S Jayakar V Loeschke W Scharloo and K W6hrmann to identify topics that seemed at least to them to be fruitful in tackling problems in population biology Consequently a number of colleagues were asked to participate in the meeting We have divided this book into chapters corresponding to the eight topics chosen The volume begins with the relation between genotype and phenotype and is followed by a chapter on quantitative genetics and selection in natural populations

Geometrical Probability and Biological Structures: Buffon's 200th Anniversary R. E.

Miles, J. Serra, 2013-03-08

Stochastic Transport Processes in Discrete Biological Systems Eckart Frehland, 2013-03-13

These notes are in part based on a course for advanced students in the applications of stochastic processes held in 1978 at the University of Konstanz These notes contain the results of recent studies on the stochastic description of ion transport through biological membranes In particular they serve as an introduction to an unified theory of fluctuations in complex biological transport systems We emphasize that the subject of this volume is not to introduce the mathematics of stochastic processes but to present a field of theoretical biophysics in which stochastic methods are important In the last years the study of membrane noise has become an important method in biophysics Valuable information on the ion transport mechanisms in membranes can be obtained from noise analysis A number of different processes such as the opening and closing of ion channels have been shown to be sources of the measured current or voltage fluctuations Biological transport systems can be complex For example the transport process can be coupled to other processes such as chemical reactions and take place in discontinuous structures of molecular dimensions Furthermore since there are strong electric fields or high concentration gradients across biological membranes ion transport processes of biological relevance are mostly processes far from equilibrium For these reasons the development of new theoretical concepts has been necessary The concept of transport in discrete systems has turned out to be more appropriate than continuum models

Stochastic Methods in Biology Motoo Kimura, Gopinath Kallianpur, Takeyuki Hida, 2013-03-13

The use of probabilistic methods in the biological sciences has been so well established by now that mathematical biology is regarded by many as a distinct discipline with its own repertoire of techniques The purpose of the Workshop on stochastic methods in biology held at Nagoya University during the week of July 8-12 1985 was to enable biologists and probabilists from Japan and the U S to discuss the latest developments in their respective fields and to exchange ideas on the applicability of the more recent developments in

stochastic process theory to problems in biology Eighteen papers were presented at the Workshop and have been grouped under the following headings I Population genetics five papers II Measure valued diffusion processes related to population genetics three papers III Neurophysiology two papers IV Fluctuation in living cells two papers V Mathematical methods related to other problems in biology epidemiology population dynamics etc six papers An important feature of the Workshop and one of the reasons for organizing it has been the fact that the theory of stochastic differential equations SDE s has found a rich source of new problems in the fields of population genetics and neuro biology This is especially so for the relatively new and growing area of infinite dimensional i e measure valued or distribution valued SDE s The papers in II and III and some of the papers in the remaining categories represent these areas

Recognition of Pattern and Form Duane G. Albrecht, 2013-03-08 Sitzungsberichte der Mathematisch-Naturwissenschaftliche Classe ,1984 **Trees and Hierarchical Structures** Andreas Dress, Arndt van Haeseler, 2013-03-09 The raison d'être of hierarchical clustering theory stems from one basic phenomenon This is the notorious non transitivity of similarity relations In spite of the fact that very often two objects may be quite similar to a third without being that similar to each other one still wants to classify objects according to their similarity This should be achieved by grouping them into a hierarchy of non overlapping clusters such that any two objects in one cluster appear to be more related to each other than they are to objects outside this cluster In everyday life as well as in essentially every field of scientific investigation there is an urge to reduce complexity by recognizing and establishing reasonable classification schemes Unfortunately this is counterbalanced by the experience of seemingly unavoidable deadlocks caused by the existence of sequences of objects each comparatively similar to the next but the last rather different from the first *Berichte über die wissenschaftliche Biologie* ,1978 **Das Öffentliche Gesundheitswesen** ,1978 **Scientific and Technical Books and Serials in Print** ,1989 Journal of Mathematical Biology ,1984 **Zeitschrift für angewandte Mathematik und Mechanik** ,1989 Subject Guide to Children's Books in Print 1997 Bowker Editorial Staff, R R Bowker Publishing, 1996-09 **Verzeichnis lieferbarer Bücher** ,1983 *Subject Guide to Books in Print* ,2001 **Population Genetics of Multiple Loci** Freddy B. Christiansen, 2000-01-10 Population Genetics of Multiple Loci F B Christiansen University of Aarhus Denmark This is a very beautiful and powerful study of an area that Christiansen has dominated for many years Marcus Feldman Stanford University USA Population genetics thrives on the constant interaction between theoretical and empirical knowledge In the first instance population genetics was developed using one locus two allele models for genetic variation The simplicity of these models opened up theoretical developments in population and evolutionary genetics to biologists without specialist training in mathematics Population genetics of multi allelic loci is more complex and requires more mathematical insight and its study is predominantly undertaken by mathematical biologists Traditional formulations of multi locus theory do not simplify by assuming two alleles per locus In this elegant presentation the author provides a formulation of multi locus population genetics that retains the

simplicity of two allele models Provides an accessible and natural extension of classical population genetics to multiple loci
 Exposes the population genetic aspects of sexual reproduction Describes the complexity of evolutionary interactions among
 genes Provides the background for insight into the functioning of genetic algorithms applied in computer science Written by
 a world leader in the field The book is divided into two main sections Part I Recombination and Segregation includes
 coverage of random mating inbreeding migration and mixing Part II Selection covers numerous phenomena involving natural
 selection including viability fertility mutation and migration The author has successfully presented the theory in a way that is
 intelligible to anyone with a reasonably good background in basic mathematics and is devoted to learning multiple loci
 population genetics The text is primarily aimed at advanced undergraduate and postgraduate students and researchers
 interested in genetics and population biology It is also essential reading for those working or researching in biomathematics
 and adaptive computing 'American Book Publishing Record' Cumulative R. R. Bowker LLC, 1978 Rhythms in Biology
and Other Fields of Application Société mathématique de France, Société mathématique de France. Journées, 1983-06 This
 volume contains most of the talks presented at the Journées de la Société Mathématique de France entitled Rhythms in Biology
 and other fields of application Deterministic and Stochastic Approaches held in Luminy from the 14th to the 18 of
 September 1981 The aim of our meeting was to bring together scientists from different disciplines to discuss a common
 topic and to stimulate exchanges between participants We hope that this goal was reached This volume is divided into four
 chapters In each one the papers are arranged in alphabetical order by first author Chapters one and two contain papers
 devoted to description or modelling of rhythmic biological phenomena Chapters three and four deal with models for the
 study of rhythms involving the use of deterministic or stochastic tools capable of fruitful transfer to Biology We are pleased
 that these Proceedings appear in a series which constitutes an interface between Biologists and Mathematicians We are
 indebted to all who provided us with their help particularly the Centre International de Rencontres Mathématiques C I R M at
 Luminy the Société Mathématique de France S M F the Delegation aux Relations Universitaires Internationales D R U I and
 the Laboratoire d'Informatique et de Mathématiques Appliquées de Grenoble I M A G Special thanks are due to Mrs A Litman
 for her dedication and her efficiency throughout the organization of this meeting Grenoble December 1982

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