



Volume 2 Virus Variability, Epidemiology, and Control

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Virus Variability, Epidemiology and Control Edouard Kurstak, R.G. Marusyk, F.A. Murphy, M.H.V. Van Regenmortel, 2013-11-11 Virus Variability and Impact on Epidemiology and Control of Diseases E Kurstak and A Hossain I INTRODUCTION An important number of virus infections and their epidemic developments demonstrate that ineffec tiveness of prevention measures is often due to the mutation rate and variability of viruses Kurstak et al 1984 1987 The new human immunodeficiency retroviruses and old influenza viruses are only one among several examples of virus variation that prevent or make very difficult the production of reliable vaccines It could be stated that the most important factor limiting the effectiveness of vaccines against virus infections is apparently virus variation. Not much is how ever known about the factors influencing and responsible for the dramatically diverse patterns of virus variability II MUTATION RATE AND VARIABILITY OF HUMAN AND ANIMAL VIRUSES Mutation is undoubtedly the primary source of variation and several reports in the literature suggest that extreme variability of some viruses may be a consequence of an unusually high mutation rate Holland et al 1982 Domingo et al 1985 Smith and Inglis 1987 The mutation rate of a virus is defined as the probability that during a single replication of the virus genome a particular nucleotide position is altered through substitution deletion insertion or recombination Different techniques have been utilized to measure virus mutation rates and these have been noted in the extent of application to different viruses Virus Variability, Epidemiology and Control Edouard Kurstak, R.G. Marusyk, F.A. Murphy, M.H.V. Van Regenmortel, 1990-09-30 Virus Variability and Impact on Epidemiology and Control of Diseases E Kurstak and A Hossain I INTRODUCTION An important number of virus infections and their epidemic developments demonstrate that ineffec tiveness of prevention measures is often due to the mutation rate and variability of viruses Kurstak et al 1984 1987 The new human immunodeficiency retroviruses and old influenza viruses are only one among several examples of virus variation that prevent or make very difficult the production of reliable vaccines It could be stated that the most important factor limiting the effectiveness of vaccines against virus infections is apparently virus variation Not much is how ever known about the factors influencing and responsible for the dramatically diverse patterns of virus variability II MUTATION RATE AND VARIABILITY OF HUMAN AND ANIMAL VIRUSES Mutation is undoubtedly the primary source of variation and several reports in the literature suggest that extreme variability of some viruses may be a consequence of an unusually high mutation rate Holland et al 1982 Domingo et al 1985 Smith and Inglis 1987 The mutation rate of a virus is defined as the probability that during a single replication of the virus genome a particular nucleotide position is altered through substitution deletion insertion or recombination Different techniques have been utilized to measure virus mutation rates and these have been noted in the extent of application to different viruses Genetic **Engineering** Jane K. Setlow, 2012-12-06 The current volume covers human gene therapy improving the nutritional value of maize restriction modification enzymes and eight other subjects Regulation of Gene Expression in Animal Viruses Luis

Carrasco, Nahum Sonenberg, Eckard Wimmer, 2012-12-06 Viruses being obligatory parasites of their host cells rely on a vast supply of cellular components for their replication regardless of whether infection leads to cell death or to the state of persistence Animal viruses are providing scientists with relatively simple models to study the molecular biology of genome replication and gene expression Whereas viruses use in general pathways of macromolecular biosynthesis common to the host cell they have a cunning ability to adopt unusual mechanisms of gene expression and gene replication provided these special pathways offer an advantage in competition for cellular resources Any study of viral gene expression and replication is likely to lead also to new insights in cellular metabolism. The discoveries of cis acting regulatory elements in transcription the phenomenon of splicing of pre mRNA and cap dependent and cap independent initiation of translation may be cited as examples In addition animal virus genomes contain elements and encode proteins that are very useful for the design of vectors for gene cloning and expression in mammalian cells Apart from the basic interest in their biology viruses have gained notoriety of course because they are pathogens Human animal viruses may cause diseases ranging from the deadly AIDS to the benign common cold All studies on animal viruses potentially lead to the development of tools for their control be it through prevention by immunization or treatment with antiviral drugs Finally viruses have yielded invaluable reagents in molecular biology as for example the vaccinia virus vector for the expression of foreign genes **Genetic Diversity of RNA** Viruses John J. Holland, 2012-12-06 Many RNA viruses have been known for decades to be genetically and biologically quite variable Some well known examples are influenza viruses foot and mouth disease viruses and Newcastle disease virus During the past decade it has become clear that most it not all RNA viruses riboviruses and retroviruses are much more mutable than was recognized previously and that this great mutability generates extremely complex populations consisting of indeterminate mixtures of related variants Le mutant swarms or guasispecies populations This is also true of DNA viruses such as hepatitis DNA genomes via RNA transcripts B virus which replicate their that are reverse transcribed back to DNA This hypermutability of RNA replicons provides great biological adaptability for RNA virus genomes It also allows but does not necessitate RNA viruses so that they can extremely rapid evolution of evolve over a million times more quickly than their eukaryotic DNA based hosts The genetics of RNA replicons is so unusual and often counterintuitive that it has many important biological conse quences which are neither readily apparent nor widely under stood Failure to understand the distinctive aspects of RNA genetics frequently generates confusion and controversy and can adversely impact vaccine and antiviral drug programs and other applications of medical virology The 14 chapters in this volume describe advances in a number of significant areas of RNA virus genetics and evolution Virus Dynamics Martin A. Nowak, Robert M. May, 2000-11-23 This text describes the emerging field of theoretical immunology in particular the use of mathematical models to describe the spread of infectious diseases within patients It reveals insights into the dynamics of viral other infections The Bunyaviridae Richard M. Elliott, 2013-06-29 In this comprehensive reference leading researchers examine

the biology molecular biology and diseases of the Bunyaviridae and provide up to date information on the genetic characterization and variations of this virus group Chapters deal with the molecular biology of five genera Bunyavirus Hantavirus Nairovirus Phlebovirus and Tospovirus Chapters examine Bunyaviridae assembly and intracelluar protein transport as well as Bunyaviridae genetics Contributors discuss the Bunyaviridae diseases including the hantavirus pulmonary syndrome Methods in Muscle Biology, 1997-11-14 Methods in Muscle Biology is a comprehensive laboratory guide that details the methods used in the study of muscle biology. The techniques included embrace cell developmental and **Evolution of Infectious Disease** Paul W. molecular biology as well asphysiology neurobiology and medical research Ewald, 1994-01-06 Findings from the field of evolutionary biology are yielding dramatic insights for health scientists especially those involved in the fight against infectious diseases. This book is the first in depth presentation of these insights In detailing why the pathogens that cause malaria smallpox tuberculosis and AIDS have their special kinds of deadliness the book shows how efforts to control virtually all diseases would benefit from a more thorough application of evolutionary principles When viewed from a Darwinian perspective a pathogen is not simply a disease causing agent it is a self replicating organism driven by evolutionary pressures to pass on as many copies of itself as possible In this context so called cultural vectors those aspects of human behavior and the human environment that allow spread of disease from immobilized people become more important than ever Interventions to control diseases don't simply hinder their spread but can cause pathogens and the diseases they engender to evolve into more benign forms In fact the union of health science with evolutionary biology offers an entirely new dimension to policy making as the possibility of determining the future course of many diseases becomes a reality By presenting the first detailed explanation of an evolutionary perspective on infectious disease the author has achieved a genuine milestone in the synthesis of health science epidemiology and evolutionary biology Written in a clear accessible style it is intended for a wide readership among professionals in these fields and general readers interested in Emerging Viruses Stephen S. Morse, 1996-08-01 New epidemics such as AIDS and mad cow disease science and health have dramatized the need to explore the factors underlying rapid viral evolution and emerging viruses This comprehensive volume is the first to describe this multifaceted new field It places viral evolution and emergence in a historical context describes the interaction of viruses with hosts and details the advances in molecular biology and epidemiology that have provided the tools necessary to track developing viral epidemics and to detect new viruses far more successfully than could be done in the recent past This unique book also lucidly details case histories and offers practical suggestions for the prevention of future epidemics The contributors are leading authorities in their disciplines and were selected both for their expert knowledge and for their ability to define and elucidate the fundamental issues The book is highly accessible and has been written for a wide audience that includes virologists public health authorities medical anthropologists evolutionary biologists geneticists infectious disease specialists and social scientists interested in medical and health issues

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