



# Molecular Microbial Ecology Manual

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# Molecular Microbial Ecology Manual

**Harinder P.S. Makkar, Gerrit J. Viljoen**



## **Molecular Microbial Ecology Manual:**

Molecular Microbial Ecology Manual A. D. L. Akkermans, J.D. van Elsas, Frans J. de Bruijn, 1996      **Molecular Microbial Ecology Manual** Antoon D. L. Akkermans, Jan Dirk Van Elsas, Frans J. De Bruijn, 2014-01-13 For a long time microbial ecology has been developed as a distinct field within Ecology In spite of the important role of microorganisms in the environment this group of invisible organisms remained unaccessible to other ecologists Detection and identification of microorganisms remain largely dependent on isolation techniques and characterisation of pure cultures We now realise that only a minor fraction of the microbial community can be cultivated As a result of the introduction of molecular methods microbes can now be detected and identified at the DNA RNA level in their natural environment This has opened a new field in ecology Molecular Microbial Ecology In the present manual we aim to introduce the microbial ecologist to a selected number of current molecular techniques that are relevant in microbial ecology The first edition of the manual contains 33 chapters and an equal number of additional chapters will be added this year Since the field of molecular ecology is in a continuous progress we aim to update and extend the Manual regularly and will invite anyone to deposit their new protocols in full detail in the next edition of this Manual We hope this book finds its place where it was born at the lab bench Antoon D L Akkermans Jan Dirk van Elsas and Frans J de Bruijn March 1995 Molecular Microbial Ecology Manual 1 3 6 1 8 1996 1996 Kluwer Academic Publishers      *Molecular microbial ecology manual*, 2004      **Molecular microbial ecology manual**, 2004      **Molecular Microbial Ecology Manual Including Supplement 3** Antoon D.L. Akkermans, Jan Dirk van Elsas, Frans J. de Bruijn, 1998 This manual provides general information about molecular techniques relevant to molecular ecological research Attention is paid to the use of molecular methods for studying the fate of genetically modified and native microorganisms in the environment      **Molecular Microbial Ecology Manual: folletos: Contents**, 2004 Section 1 Isolation of nucleic acids Section 2 Detection of microbial nucleic acid sequences Section 3 Identification and classification of microbes using DNA and RNA sequences Section 4 Detection identification and classification of microbes using other methods Section 5 Detection of gene transfer in the environment Section 6 Tracking of specific microbes in the environment Section 7 Statistical computer assisted and other analyses Section 8 Molecular tools to assess microbial activities

*Handbook of Molecular Microbial Ecology I* Frans J. de Bruijn, 2011-10-14 The premiere two volume reference on revelations from studying complex microbial communities in many distinct habitats Metagenomics is an emerging field that has changed the way microbiologists study microorganisms It involves the genomic analysis of microorganisms by extraction and cloning of DNA from a group of microorganisms or the direct use of the purified DNA or RNA for sequencing which allows scientists to bypass the usual protocol of isolating and culturing individual microbial species This method is now used in laboratories across the globe to study microorganism diversity and for isolating novel medical and industrial compounds Handbook of Molecular Microbial Ecology is the first comprehensive two volume reference to cover unculturable

microorganisms in a large variety of habitats which could not previously have been analyzed without metagenomic methodology. It features review articles as well as a large number of case studies based largely on original publications and written by international experts. This first volume, *Metagenomics and Complementary Approaches*, covers such topics as background information on DNA reassociation and use of 16 rRNA and other DNA fingerprinting approaches, species designation in microbiology, metagenomics, introduction to the basic tools with examples, consortia and databases, bioinformatics, computer-assisted analysis, complementary approaches, microarrays, metatranscriptomics, metaproteomics, metabolomics, and single cell analysis. A special feature of this volume is the highlighting of the databases and computer programs used in each study; they are listed along with their sites in order to facilitate the computer-assisted analysis of the vast amount of data generated by metagenomic studies.

*Handbook of Molecular Microbial Ecology I* is an invaluable reference for researchers in metagenomics, microbiology, and environmental microbiology, those working on the Human Microbiome Project, microbial geneticists, molecular microbial ecologists, and professionals in molecular microbiology and bioinformatics. *Molecular Microbial Ecology Manual*, 2004. Methods in Gut Microbial Ecology for Ruminants, Harinder P.S. Makkar, Christopher S. McSweeney, 2006-02-23.

As a result of various human activities such as increase in human population, decrease in arable land due to soil degradation, urbanization, industrialization, and associated increase in the demand for livestock products, dramatic changes are occurring in the global ruminant livestock sector. These changes include shift in the size of regional livestock populations and in the types of management and feeding systems under which ruminant livestock are held, and increased demand of a wider range of quality attributes from animal agriculture, not just of the products themselves but also of the methods used in their production. The livestock sector will need to respond to new challenges of increasing livestock productivity while protecting environment and human health and conserving biodiversity and natural resources. The microorganisms in the digestive tracts of ruminant livestock have a profound influence on the conversion of feed into end products which can impact on the animal and the environment. As the livestock sector grows, particularly in developing countries, there will be an increasing need to understand these processes for better management and use of both feed and other natural resources that underpin the development of sustainable feeding systems. *Molecular Microbial Ecology Manual*, Antoon D. L. Akkermans, Frans J. de Bruijn, Jan Dirk van Elsas, 1996. **Handbook of Molecular Microbial Ecology II**, Frans J. de Bruijn, 2011-10-14.

The premiere two-volume reference on revelations from studying complex microbial communities in many distinct habitats. Metagenomics is an emerging field that has changed the way microbiologists study microorganisms. It involves the genomic analysis of microorganisms by extraction and cloning of DNA from a group of microorganisms or the direct use of the purified DNA or RNA for sequencing, which allows scientists to bypass the usual protocol of isolating and culturing individual microbial species. This method is now used in laboratories across the globe to study microorganism diversity and for isolating novel medical and industrial compounds. *Handbook of*

Molecular Microbial Ecology is the first comprehensive two volume reference to cover unculturable microorganisms in a large variety of habitats which could not previously have been analyzed without metagenomic methodology It features review articles as well as a large number of case studies based largely on original publications and written by international experts This second volume Metagenomics in Different Habitats covers such topics as Viral genomes Metagenomics studies in a variety of habitats including marine environments and lakes soil and human and animal digestive tracts Other habitats including those involving microbiome diversity in human saliva and functional intestinal metagenomics diversity of archaea in terrestrial hot springs and microbial communities living at the surface of building stones Biodegradation Biocatalysts and natural products A special feature of this book is the highlighting of the databases and computer programs used in each study they are listed along with their sites in order to facilitate the computer assisted analysis of the vast amount of data generated by metagenomic studies Such studies in a variety of habitats are described here which present a large number of different system dependent approaches in greatly differing habitats Handbook of Molecular Microbial Ecology II is an invaluable reference for researchers in metagenomics microbial ecology microbiology and environmental microbiology those working on the Human Microbiome Project microbial geneticists and professionals in molecular microbiology and bioinformatics Handbook of Molecular Microbial Ecology I Frans J. De Bruijn, 2011 Handbook of Molecular Microbial Ecology, 2 Volume Set Frans J. de Bruijn, 2011-06-07 Metagenomics has revolutionized microbiology and many associated health and environmental fields This is the first comprehensive treatise covering the various omics in a large variety of habitats which could not previously have been analysed without metagenomic methodology This is a reference work both for people in the field and those who would like to enter it Two volume set covers topics including Volume One Species designations in microbiology Metagenomics Consortia and databases Bioinformatics Microarrays Other omics applications Volume Two Viral genomes Metagenomics studies in a variety of habitats including marine soil and the digestive tract Biodegradation Biocatalysts and natural products Microbial Ecology of Activated Sludge Robert Seviour, Per Halkjaer Nielsen, 2010-01-15 Microbial Ecology of Activated Sludge written for both microbiologists and engineers critically reviews our current understanding of the microbiology of activated sludge the most commonly used process for treating both domestic and industrial wastes The contributors are all internationally recognized as leading research workers in activated sludge microbiology and all have made valuable contributions to our present understanding of the process The book pays particular attention to how the application of molecular methods has changed our perceptions of the identity of the filamentous bacteria causing the operational disorders of bulking and foaming and the bacteria responsible for nitrification and denitrification and phosphorus accumulation in nutrient removal processes Special attention is given to how it is now becoming possible to relate the composition of the community of microbes present in activated sludge and the in situ function of individual populations there and how such information might be used to manage and control these systems better

Detailed descriptions of some of these molecular methods are provided to allow newcomers to this field of study an opportunity to apply them in their research Comprehensive descriptions of organisms of interest and importance are also given together with high quality photos of activated sludge microbes Activated sludge processes have been used globally for nearly 100 years and yet we still know very little of how they work In the past 15 years the advent of molecular culture independent methods of study have provided tools enabling microbiologists to understand which organisms are present in activated sludge and critically what they might be doing there Microbial Ecology of Activated Sludge will be the first book available to deal comprehensively with the very exciting new information from applying these methods and their impact on how we now view microbiologically mediated processes taking place there As such it will be essential reading for microbial ecologists environmental biotechnologists and engineers involved in designing and managing these plants It will also be suitable for postgraduate students working in this field

**Marine Microbiology** John Paul, 2001-05-08 Written by experts in the field Marine Microbiology presents the latest experimental techniques in the detail required for modern environmental microbiological research Chapters start with the introduction and background of a particular method followed by a concise description of the procedures involved There is also a list of vendors who supply critical components which includes names addresses and websites at the end of each chapter Covers both established and novel current methods Historical perspective Enumeration of autotrophic picoplankton bacteria and viruses Fingerprinting Viral Assemblages by Pulsed Field Gel electrophoresis PFGE Fluorescence in situ hybridization with rRNA targeted oligonucleotide probes Detection of phytoplankton by remote sensing Marine pollution microbiology Microbes in extreme environments List of suppliers provided at end of each chapter Colour plate section

*Manual of Environmental Microbiology* Christon J. Hurst, Ronald L. Crawford, Jay L. Garland, David A. Lipson, 2007-05-14 The most definitive manual of microbes in air water and soil and their impact on human health and welfare Incorporates a summary of the latest methodology used to study the activity and fate of microorganisms in various environments Synthesizes the latest information on the assessment of microbial presence and microbial activity in natural and artificial environments Features a section on biotransformation and biodegradation Serves as an indispensable reference for environmental microbiologists microbial ecologists and environmental engineers as well as those interested in human diseases water and wastewater treatment and biotechnology

**New Approaches for the Generation and Analysis of Microbial Typing Data** L. Dijkshoorn, K.J. Towner, Mark J Struelens, 2001-07-10 Rapid molecular identification and typing of micro organisms is extremely important in efforts to monitor the geographical spread of virulent epidemic or antibiotic resistant pathogens It has become a mainstay of integrated hospital infection control service In addition numerous industrial and biotechnological applications require the study of the diversity of organisms Conventional phenotypic identification and typing methods have long been the mainstay of microbial population and epidemiological studies but such methods often lack adequate discrimination and their use is normally confined to the group

of organisms for which they were originally devised Molecular fingerprinting methods have flourished in recent years and many of these new methods can be applied to numerous different organisms for a variety of purposes Standardisation of these methods is vitally important In addition the generation of large numbers of complex fingerprint profiles requires that a computer assisted strategy is used for the formation and analysis of databases The purpose of this book is to describe the best fingerprinting methods that are currently available and the computer assisted strategies that can be used for analysis and exchange of data between laboratories This book is dedicated to the memory of Jan Ursing 1926 2000 Swedish microbiologist taxonomist and philosopher taxonomy is on the borders of philosophy because we do not know the natural continuities and discontinuities

**Modern Soil Microbiology, Second Edition** Jan Dirk van Elsas, Janet K. Jansson, Jack T. Trevors, 2006-12-21 In the ten years since the publication of Modern Soil Microbiology the study of soil microbiology has significantly changed both in the understanding of the diversity and function of soil microbial communities and in research methods Ideal for students in a variety of disciplines this second edition provides a cutting edge examination of a fascinating discipline that encompasses ecology physiology genetics molecular biology and biotechnology and makes use of biochemical and biophysical approaches The chapters cover topics ranging from the fundamental to the applied and describe the use of advanced methods that have provided a great thrust to the discipline of soil microbiology Using the latest molecular analyses they integrate principles of soil microbiology with novel insights into the physiology of soil microorganisms The authors discuss the soil and rhizosphere as habitats for microorganisms then go on to describe the different microbial groups their adaptive responses and their respective processes in interactive and functional terms The book highlights a range of applied aspects of soil microbiology including the nature of disease suppressive soils the use of biological control agents biopesticides and bioremediation agents and the need for correct statistics and experimentation in the analyses of the data obtained from soil systems

Biological Nitrogen Fixation for the 21st Century Claudine Elmerich, Adam Kondorosi, William E. Newton, 2013-12-01 Nitrogen availability is one of the most critical factors that limits plant productivity The largest reservoir of nitrogen is the atmosphere but this gaseous molecular nitrogen only becomes available to plants through the biological nitrogen fixation process which only prokaryotic cells have developed The discovery that microbes were providing fixed nitrogen to legumes and the isolation of the first nitrogen fixing bacteria occurred at the end the 19th Century in Louis Pasteur's time We are now building on more than 100 years of research in this field and looking towards the 21st Century The International Nitrogen Fixation Congress series Started more than 20 years ago The format of this Congress is designed to gather scientists from very diverse origins backgrounds interests and scientific approaches and is a forum where fundamental knowledge is discussed alongside applied research This confluence of perspectives is we believe extremely beneficial in raising new ideas questions and concepts

Applications of Gene-Based Technologies for Improving Animal Production and Health in Developing Countries Harinder P.S. Makkar, Gerrit J. Viljoen, 2005-08-12 Modern Biotechnology has

potential for solving many problems associated with animal productivity and health and offers exciting opportunities for enhancing agricultural productivity At present the focus is however on the issues and problems of significance for livestock producers in the developed world In order to fully realize the benefits of this technology in developing countries there is a need to identify characterize and apply appropriate gene based technologies for these regions These proceedings present peer reviewed state of the art papers describing the achievements in the areas of animal breeding and genetics animal nutrition animal health and environment ethics safety and regulatory aspects of gene based technologies achievements which could be realized using these modern scientific tools to maximise the benefits from the livestock revolution that is taking place and the constraints in the use of gene based technologies and their specific research needs This book will help in bridging the wide gap between developed and developing countries in the development and use of gene based technologies and to elucidate the current and future roles of such technologies in the developing world It is a good reference source for researchers students and policy makers alike



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