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
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Systems biology is the study of the dynamic interactions of more than one component in a biological system in order to understand and predict the behavior of the system as a whole. Systems biology is a rapidly expanding discipline fuelled by the 'omics' era and new technological advances that have increased the precision of data. A focus on simple single cell organisms, such as bacteria, aids tractability and means that systems microbiology is a rapidly maturing science.

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Recommended for all microbiology laboratories, this book contains cutting-edge reviews by world-leading experts on the systems biology of microorganisms. As well as covering theoretical approaches and mathematical modeling, the book includes case studies on single microbial species of bacteria and archaea, and it explores the systems analysis of microbial phenomena, such as chemotaxis and phagocytosis. The topics covered include: the mathematical models for systems biology * systems biology of *Escherichia coli* metabolism * bacterial

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Deep subsurface microbiology is a highly active and rapidly advancing research field at the interface of

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Microbiology and the geosciences; it focuses on the detection, identification, quantification, cultivation and activity measurements of bacteria, archaea and eukaryotes that permeate the subsurface biosphere of deep marine sediments and the basaltic ocean and continental crust. The deep subsurface biosphere abounds with uncultured, only recently discovered and - at best - incompletely understood microbial populations. In spatial extent and volume, Earth's subsurface biosphere is only rivaled by the deep sea water column. So far, no

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Paper deep subsurface sediment has been found that is entirely devoid of microbial life; microbial cells and DNA remain detectable at sediment depths of more than 1 km; microbial life permeates deeply buried hydrocarbon reservoirs, and is also found several kilometers down in continental crust aquifers. Severe energy limitation, either as electron acceptor or donor shortage, and scarcity of microbially degradable organic carbon sources are among the evolutionary pressures that have shaped the genomic and physiological repertoire of the deep subsurface

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Biosphere. Its biogeochemical role as long-term organic carbon repository, inorganic electron and energy source, and subduction recycling engine continues to be explored by current research at the interface of microbiology, geochemistry and biosphere/geosphere evolution. This Research Topic addresses some of the central research questions about deep subsurface microbiology and biogeochemistry: phylogenetic and physiological microbial diversity in the deep subsurface; microbial activity and survival

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Paper strategies in severely energy-limited subsurface habitats; microbial activity as reflected in process rates and gene expression patterns; biogeographic isolation and connectivity in deep subsurface microbial communities; the ecological standing of subsurface biospheres in comparison to the surface biosphere - an independently flourishing biosphere, or mere survivors that tolerate burial (along with organic carbon compounds), or a combination of both? Advancing these questions on Earth's deep subsurface biosphere redefines the habitat range, environmental tolerance,

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Paper activity and diversity of microbial life.

Forest Microbiology, Volume One: Tree Microbiome: Phyllosphere, Endosphere and Rhizosphere places an emphasis on the microbiology of leaves, needles, stems, roots, litter and soil. This comprehensive title is split into five sections, including the phyllosphere microbiome, endosphere, rhizosphere, archaea, viruses in forest ecosystem and microbiota of forest nurseries and tree pests, challenges and potentials. Microbial communities associated with various host trees and different tree

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Paper tissues are compared, and generalists and specialists among tree-associated microbes are identified. In addition, biotic and abiotic factors determining the composition and the structure of forest tree microbial communities are presented, along with the concept of microbial 'hubs.' Together, the book's editors have 25 years' worth of experience teaching and conducting research on forest microbiology, making this an essential read for any scientist interested in the forest microbiome. Addresses the microbiology of living organs of forest trees including needles,

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Paper, stems and roots
Highlights the potential
impact of microbiota
inhabiting forest trees on
the health and fitness of,
and disease progression in,
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phyllosphere, endosphere and
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microbiome

Marine and freshwater polar
environments are
characterized by intense
physical forces and strong
seasonal variations. The
persistent cold and
sometimes inhospitable
conditions create unique
ecosystems and habitats for
microbial life. Polar
microbial communities are

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Paper diverse productive assemblages, which drive biogeochemical cycles and support higher food-webs across the Arctic and over much of the Antarctic. Recent studies on the biogeography of microbial species have revealed phylogenetically diverse polar ecotypes, suggesting adaptation to seasonal darkness, sea-ice coverage and high summer irradiance. Because of the diversity of habitats related to atmospheric and oceanic circulation, and the formation and melting of ice, high latitude oceans and lakes are ideal environments to investigate

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Paper composition and functionality of microbial communities. In addition, polar regions are responding more dramatically to climate change compared to temperate environments and there is an urgent need to identify sensitive indicators of ecosystem history, that may be sentinels for change or adaptation. For instance, Antarctic lakes provide useful model systems to study microbial evolution and climate history. Hence, it becomes essential and timely to better understand factors controlling the microbes, and how, in turn, they may affect the functioning of these fragile

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Paper ecosystems. Polar microbiology is an expanding field of research with exciting possibilities to provide new insights into microbial ecology and evolution. With this Research Topic we seek to bring together polar microbiologists studying different aquatic systems and components of the microbial food web, to stimulate discussion and reflect on these sensitive environments in a changing world perspective.

Exponentially increasing information on biological organisms coupled with increasing computational

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Paper in the past decade have broadened the perspective of fundamental biological research, bringing about considerable promise and unprecedented potential for practical applications in biotechnology. As one emergent discipline, synthetic biology aims to design and engineer novel biologically-based parts, devices, and systems, in addition to redesigning existing, natural biological systems. Although previously relegated to demonstration studies, more recent research in synthetic biology has focused on the rational engineering of

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Industrial microorganisms with the potential to address many of society's critical challenges. Within the realm of industrial microbiology, progress in the field of synthetic biology has enabled the development of, for example, new biosynthetic pathways for the production of renewable fuels and chemicals, programmable logic controls to regulate and optimize cell function, and robust microbes for the destruction of harmful environmental contaminants. Some of the exciting examples included producing anti-malarial drug, anti-cancer taxol precursor and

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Various biofuel molecules in E. coli and yeast. In addition, these researches have also greatly enhanced our understanding of the cellular machinery and its regulation in some of the industry important microbes, laying an important foundation for further design and engineering of biological function for even greater application. For these reasons, we present here a collection of articles from the leading edge of the field of synthetic biology, with a specific focus on the development in industrial microorganisms. It is the intent of this collection to

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Paper reach a wide audience whose interests and expertise spans from development of novel synthetic biology methodologies and theories (both experimental and computational) to practical applications seeking to address issues facing the world today.

Advances in Applied Microbiology, Volume 115 continues the comprehensive reach of this widely read and authoritative review source in microbiology. Users will find invaluable references and information on a variety of areas relating to the topic of microbiology, with this

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Paper focusing on the preparation of functional oligosaccharides using microbial coupling fermentation. Contains contributions from leading authorities in the field
Informs and updates on the latest developments in the field of microbiology
Includes discussions on the role of specific molecules in pathogen life stages, interactions, and much more

This book ventures into a new and exciting area of discovery that directly ties our current knowledge of cancer to the discovery of microorganisms associated with different types of

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Paper. Recent studies demonstrate that microorganisms are directly linked to the establishment of cancers and that they can also contribute to the initiation, as well as persistence of, the cancers. *Microbiome and Cancer* covers the current knowledge of microbiome and its association with human cancers. It provides important reading for novices, senior undergraduates in cancer and microbiology, graduate students, junior investigators, residents, fellows and established investigators in the fields of cancer and microbiology.

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Paper We cover areas related to known, broad concepts in microbiology and how they can relate to the ongoing discoveries of the micro-environment and the changes in the metabolic and physiologic states in that micro-environment, which are important for the ongoing nurturing and survival of the poly-microbial content that dictates activities in that micro-environment. We cover the interactions of microorganisms associated with gastric carcinomas, which are important for driving this particular cancer. Additional areas include oral cancers, skin cancers, ovarian cancers,

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Paper breast cancers, nasopharyngeal cancers, lung cancers, mesotheliomas, Hodgkin's and non-Hodgkin's lymphomas, glioblastoma multiforme, hepatocellular carcinomas, as well as the inflammatory response related to the infectious agents in cancers. This book covers the metabolic changes that occur because of infection and their support for development of cancers, chronic infection and development of therapeutic strategies for detection and control of the infection. The field of microbiome research has exploded over the last five years, and we are now understanding more

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Paper and more about the context in which microorganisms can contribute to the onset of cancers in humans. The field of microbiome research has demonstrated that the human body has specific biomes for tissues and that changes in these biomes at the specific organ sites can result in disease. These changes can result in dramatic differences in metabolic shifts that, together with genetic mutations, will produce the perfect niche for establishment of the particular infection programmes in that organ site. We are just beginning to understand what those changes are and how they

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Paper influence the disease state. Overall, we hope to bring together the varying degrees of fluctuations in the microbiome at the major organ sites and how these changes affect the normal cellular processes because of dysregulation, leading to proliferation of the associated tissues.

The Microbiology of Nuclear Waste Disposal is a state-of-the-art reference featuring contributions focusing on the impact of microbes on the safe long-term disposal of nuclear waste. This book is the first to cover this important emerging topic, and is written for a wide

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Paper encompassing regulators, implementers, academics, and other stakeholders. The book is also of interest to those working on the wider exploitation of the subsurface, such as bioremediation, carbon capture and storage, geothermal energy, and water quality. Planning for suitable facilities in the U.S., Europe, and Asia has been based mainly on knowledge from the geological and physical sciences. However, recent studies have shown that microbial life can proliferate in the inhospitable environments

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associated with radioactive waste disposal, and can control the long-term fate of nuclear materials. This can have beneficial and damaging impacts, which need to be quantified.

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