Dansuke Kihara Editor

Protein Function Prediction for Omics Era



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Protein Function Prediction For Omics Era

Roman Wölfel

Protein Function Prediction For Omics Era:

Protein Function Prediction for Omics Era Daisuke Kihara, 2011-04-19 Gene function annotation has been a central question in molecular biology The importance of computational function prediction is increasing because more and more large scale biological data including genome sequences protein structures protein protein interaction data microarray expression data and mass spectrometry data are awaiting biological interpretation Traditionally when a genome is sequenced function annotation of genes is done by homology search methods such as BLAST or FASTA However since these methods are developed before the genomics era conventional use of them is not necessarily most suitable for analyzing a large scale data Therefore we observe emerging development of computational gene function prediction methods which are targeted to analyze large scale data and also those which use such omics data as additional source of function prediction In this book we overview this emerging exciting field The authors have been selected from 1 those who develop novel purely computational methods 2 those who develop function prediction methods which use omics data 3 those who maintain and update data base of function annotation of particular model organisms E coli which are frequently referred Analyzing Network Data in Biology and Medicine Nataša Pržuli, 2019-03-28 The increased and widespread availability of large network data resources in recent years has resulted in a growing need for effective methods for their analysis. The challenge is to detect patterns that provide a better understanding of the data However this is not a straightforward task because of the size of the data sets and the computer power required for the analysis The solution is to devise methods for approximately answering the questions posed and these methods will vary depending on the data sets under scrutiny This cutting edge text introduces biological concepts and biotechnologies producing the data graph and network theory cluster analysis and machine learning before discussing the thought processes and creativity involved in the analysis of large scale biological and medical data sets using a wide range of real life examples Bringing together leading experts this text provides an ideal introduction to and insight into the interdisciplinary field of network data analysis in biomedicine Biological Knowledge Discovery Handbook Mourad Elloumi, Albert Y. Zomaya, 2015-02-04 The first comprehensive overview of preprocessing mining and postprocessing of biological data Molecular biology is undergoing exponential growth in both the volume and complexity of biological data and knowledge discovery offers the capacity to automate complex search and data analysis tasks This book presents a vast overview of the most recent developments on techniques and approaches in the field of biological knowledge discovery and data mining KDD providing in depth fundamental and technical field information on the most important topics encountered Written by top experts Biological Knowledge Discovery Handbook Preprocessing Mining and Postprocessing of Biological Data covers the three main phases of knowledge discovery data preprocessing data processing also known as data mining and data postprocessing and analyzes both verification systems and discovery systems BIOLOGICAL DATA PREPROCESSING Part A Biological Data Management Part B Biological Data Modeling Part C Biological Feature Extraction Part D Biological

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Tuberculosis Johnjoe McFadden, Dany J.V. Beste, Andrzej M. Kierzek, 2012-12-09 The book starts with a general introduction into the relevance of systems biology for understanding tuberculosis This will be followed by several chapters which describe the application of systems biology to various aspects of the study of the pathogen Mycobacterium tuberculosis and its interaction with the host The book provides the reader with an account of how the new science of systems biology is providing novel insights into the ancient scourge of tuberculosis It will also describe how systems biology can be applied to the control of tuberculosis including the development of new treatments vaccines and diagnostics Biomedical Sciences and Environmental Sustainability Minu Kesheri, Swarna Kanchan, Donat-P. Häder, Rajeshwar P. Sinha, 2025-06-25 This book offers a comprehensive exploration of the cutting edge multi omics technologies that are revolutionizing research across biomedical sciences and environmental sustainability It addresses the urgent need for interdisciplinary research by integrating multi omics approaches with bioinformatics and artificial intelligence The book explores evolution of traditional omics technologies into comprehensive multi omics strategies that synergize data output through advanced computational tools It covers diverse topics such as health and disease mechanisms drug discovery innovations COVID 19 responses cancer treatment personalization neuroscience insights into brain disorders cyanobacterial natural compounds potential for biofuel production lichen symbiosis studies and more This volume integrates genomics proteomics metabolomics and more with bioinformatics machine learning and artificial intelligence to address complex challenges in health and the environment With contributions from renowned scholars worldwide this book illuminates recent advances through illustrative figures and case studies that enhance understanding of complex pathways while bioinformatics strategies streamline research outcomes This book is a must read for researchers academics and professionals in life sciences biomedical fields and environmental studies interested in advancing their knowledge of multi omics applications It is also beneficial for scientists involved in drug design or biotechnological innovations related to environmental sustainability

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physiomics signalomics thiolomics organelle omics micro morphomics microbiomics cryobionomics nanotechnology pharmacogenomics and computational systems biology for plants It provides up to date information technologies and their applications that can be adopted and applied easily for deeper understanding plant biology and therefore will be helpful in developing the strategy for generating cost effective superior plants for various purposes In the last chapter the editors have proposed several new areas in plant omics that may be explored in order to develop an integrated meta omics strategy to ensure the world and earth s health and related issues This book will be a valuable resource to students and researchers in Neglected and Underutilized Crops - Towards Nutritional Security and Sustainability the field of cutting edge plant omics Sajad Majeed Zargar, Antonio Masi, Romesh Kumar Salgotra, 2021-09-29 This book covers important topics on various neglected and underutilised crops vegetables cereals fruit crops It gives an overview of the potential availability of genetic and genomic resources and the future prospects of these food crops The book presents different chapters on the importance of underutilised crops with respect to sustainable agriculture and describes the approaches that must be followed for improving the yield and production of these crops It covers a wide range of food crops such as millet buckwheat underutilised spices underutilised vegetables and underutilised fruit crops It also provides insights on what smart foods are And whether these neglected crops qualify as smart foods This up to date and informative book is meant for food scientists geneticists breeders and biotechnologists It is of interest to students researchers and course instructors in these fields

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