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Second Edition



Rna Bioinformatics Methods In Molecular Biology

Peter G. Clote

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RNA Bioinformatics Ernesto Picardi, 2021-05-22 This detailed book aims to provide an overview of novel bioinformatics resources for exploring diverse aspects of RNA biology This edition focuses on methods dealing with non coding RNA miRNAs circRNAs or lncRNAs RNA modifications m6A or RNA editing single cell RNA seg and statistical models to handle count data from RNA seg experiments The book also includes chapters based on the classical RNA bioinformatics methods such as those for deciphering secondary and tertiary RNA structures however they are revised to take into account deep sequencing data Finally chapters describing methods to analyze RNA sequencing data from emerging third generation sequencing technologies that could provide interesting insights into the transcriptional process and its regulation are also included Written for the highly successful Methods in Molecular Biology series chapters include the kind of expert implementation advice that encourages quality results Comprehensive and up to date RNA Bioinformatics Second Edition serves as an ideal guide for researchers digging ever deeper into the depths of the study of RNAs The chapter RNA Seg Data Analysis in Galaxy is open access under a CC BY 4 0 license RNA Sequence, Structure, and Function Jan Gorodkin, Walter L. Ruzzo, 2014 The existence of genes for RNA molecules not coding for proteins ncRNAs has been recognized since the 1950 s but until recently aside from the critically important ribosomal and transfer RNA genes most focus has been on protein coding genes However a long series of striking discoveries from RNA's ability to carry out catalytic function to discovery of riboswitches microRNAs and other ribo regulators performing critical tasks in essentially all living organisms has created a burgeoning interest in this primordial component of the biosphere However the structural characteristics and evolutionary constraints on RNA molecules are very different from those of proteins necessitating development of a completely new suite of informatic tools to address these challenges In RNA Sequence Structure Function Computational and Bioinformatic Methods expert researchers in the field describe a substantial and relevant fraction of these methodologies from both practical and computational algorithmic perspectives Focusing on both of these directions addresses both the biologist interested in knowing more about RNA bioinformatics as well as the bioinformaticist interested in more detailed aspects of the algorithms Written in the highly successful Methods in Molecular Biology series format the chapters include the kind of detailed description and implementation advice that is crucial for getting optimal results Thorough and intuitive RNA Sequence Structure Function Computational and Bioinformatic Methods aids scientists in continuing to study key methods and principles of RNA bioinformatics Computational Biology of Non-Coding RNA Xin Lai, Shailendra Gupta, Julio Vera Gonzalez, 2024-12-19 This second edition details a collection of state of art methods including identification of novel ncRNAs and their targets functional annotation and disease association in different biological contexts Chapters guide readers through an overview of disease specific ncRNAs computational methods and workflows for ncRNA discovery annotation based on high throughput sequencing data bioinformatics tools and databases for ncRNA analyses

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