

Cancer Drug Discovery and Development

Bioinformatics in Cancer and Cancer Therapy

Edited by
Gavin J. Gordon

 Humana Press

Bioinformatics In Cancer And Cancer Therapy Cancer Drug Discovery And Development

Gavin J. Gordon



Bioinformatics In Cancer And Cancer Therapy Cancer Drug Discovery And Development:

Bioinformatics in Cancer and Cancer Therapy Gavin J. Gordon, 2008-10-25 Bioinformatics can be loosely defined as the collection classification storage and analysis of biochemical and biological information using computers and mathematical algorithms Bioinformatics represents a marriage of biology medicine computer science physics and mathematics fields of study that have historically existed as mutually exclusive disciplines Edited by Gavin Gordon Bioinformatics in Cancer and Cancer Therapy the focus of this book is to provide a historical and technical perspective on the analytical techniques methodologies and platforms used in bioinformatics experiments to show how a bioinformatics approach has been used to characterize various cancer related processes and to demonstrate how a bioinformatics approach is being used to bridge basic science and the clinical arena to positively impact patient care and management **Systems**

Biology in Cancer Research and Drug Discovery Asfar S Azmi, 2012-09-29 Systems Biology in Cancer Research and Drug Discovery provides a unique collection of chapters by world class researchers describing the use of integrated systems biology and network modeling in the cancer field where traditional tools have failed to deliver expected promise This book touches four applications aspects of systems biology i in understanding aberrant signaling in cancer ii in identifying biomarkers and prognostic markers especially focused on angiogenesis pathways iii in unwinding microRNAs complexity and iv in anticancer drug discovery and in clinical trial design This book reviews the state of the art knowledge and touches upon cutting edge newer and improved applications especially in the area of network modeling It is aimed at an audience ranging from students academics basic researcher and clinicians in cancer research This book is expected to benefit the field of translational cancer medicine by bridging the gap between basic researchers computational biologists and clinicians who have one ultimate goal and that is to defeat cancer **Computational Drug Discovery of Medicinal Compounds for**

Cancer Management Sibhghatulla Shaikh ,Khurshid Ahmad ,Mohammad Ehtisham Khan ,Faez Iqbal Khan , Cancer is still a major public health concern and it is regarded as one of the leading causes of mortality globally Despite significant advances in biotechnologies developing practical and innovative small molecule drugs remains a difficult time consuming and expensive task that necessitates collaborations from many experts in multidisciplinary fields such as computational biology drug metabolism and clinical research among others Therefore new drug development procedures that save time and cost while increasing efficiency are in high demand In silico screening in combination with molecular simulations has become an increasingly important aspect of modern drug development processes Understanding ligand protein interaction is critical in all areas of drug design and discovery Computational approaches such as molecular docking molecular dynamics simulations pharmacophore modeling and QSAR etc are efficient tools for obtaining insights on structure function relationships for small molecules and or medicinal compounds with target proteins and are widely used in the identification and optimization of leads The goal of the drug discovery process is to predict a drug candidate s metabolic fate in order to build a link between

pharmacodynamics and pharmacokinetics and to identify the drug candidate's possible toxicity. The advancement of *in silico* techniques in recent years has enabled researchers to collect more trustworthy data. This Research Topic will focus on the use and application of computational methods that can aid in the drug design of medicinal compounds targeting various proteins for cancer management, which is a critical requirement in the pharmaceutical sectors.

Computational Methods in Drug Discovery and Repurposing for Cancer Therapy Ganji Purnachandra Nagaraju, Venkatesan Amouda, Ampasala Dinakara Rao, 2023-03-22. *Computational Methods in Drug Discovery and Repurposing for Cancer Therapy* provides knowledge about ongoing research as well as computational approaches for drug discovery and repurposing for cancer therapy. The book also provides detailed descriptions about target molecules, pathways, and their inhibitors for easy understanding and applicability. The book discusses tools and techniques such as integrated bioinformatics approaches, systems biology, tools, molecular docking, computational chemistry, artificial intelligence, machine learning, structure-based virtual screening, biomarkers, and transcriptome. These are discussed in the context of different cancer types such as colon, pancreatic, glioblastoma, endometrial, and retinoblastoma, among others. This book is a valuable resource for researchers, students, and members of the biomedical and medical fields who want to learn more about the use of computational modeling to better tailor the treatment for cancer patients. It discusses *in silico* remodeling of effective phytochemical compounds for discovering improved anticancer agents for substantial significant cancer therapy. It covers potential tools of bioinformatics that are applied toward discovering new targets by drug repurposing and strategies to cure different types of cancers. It demonstrates the significance of computational and artificial intelligence approaches in anticancer drug discovery. It explores how these various advances can be integrated into a precision and personalized medicine approach that can eventually enhance patient care.

Advances in Targeted Cancer Therapy Richard M. Schultz, 2006-01-24. There have been tremendous advances in our understanding of molecular and tumor biology during the past few years. In the field of cancer therapeutics, it is expected that cytotoxic drug approaches will be gradually replaced with treatments based on biological targeted approaches. Hopefully, these new targeted therapies will significantly increase efficacy and lack the devastating and troublesome side effects elicited by cytotoxic chemotherapy. This volume is the first book to cover the general topic of targeted cancer therapy. It presents a range of targets such as tumor angiogenesis, cell cycle control, and cell signalling, COX 2, apoptosis, cell survival, invasion, and metastasis, and approaches like kinase inhibitors, antisense, and antibody-based therapeutics. The emphasis is on preclinical development, including target validation, development of biomarkers, strategies for combination approaches, and development of resistance. The particular challenges involved in translating these data to clinical application are discussed. This volume should be of broad general interest to researchers and clinicians involved in cancer therapy as well as other scientists interested in current strategies for cancer treatment.

The Oncogenomics Handbook William J. LaRochelle, Richard A. Shimkets, 2007-11-09. An integrated overview of cancer drug discovery and development from the bench to the clinic, showing

with broad strokes and representative examples the drug development process as a network of linked components leading from the discovered target to the ultimate therapeutic product Following a systems biology approach the authors explain genomic databases and how to discover oncological targets from them how then to advance from the gene and transcript to the level of protein biochemistry how next to move from the chemical realm to that of the living cell and ultimately pursue animal modeling and clinical development Emerging cancer therapeutics including Ritux an Erbitux Gleevec Herceptin Avastin ABX EGF Velcade Kepivance Iressa Tarceva and Zevalin are addressed Highlights include cancer genomics pharmacogenomics transcriptomics gene expression analysis proteomic and enzymatic cancer profiling technologies and cellular and animal approaches to cancer target validation

Targeting Autophagy in Cancer Therapy Jin-Ming Yang, 2016-09-02 This volume will detail the current state and perspectives of autophagy based cancer therapy Covering a wide range of topics it will include an overview of autophagy as a therapeutic target in cancer autophagy modulators as cancer therapeutic agents implications of micro RNA regulated autophagy in cancer therapy modulation of autophagy through targeting PI3 kinase in cancer therapy targeting autophagy in cancer stem cells and roles of autophagy in cancer immunotherapy In addition the volume will review applications of system biology and bioinformatics approaches to discovering cancer therapeutic targets in the autophagy regulatory network The volume will be beneficial for a variety of basic and clinical scientists including cancer biologists autophagy researchers pharmacologists and clinical oncologists who wish to delve more deeply into this field of cancer research This volume will be the first book to focus solely on autophagy as a target in cancer therapy As well it will comprehensively discuss the roles of autophagy in most currently available cancer treatments

Cancer Bioinformatics Sylvia Nagl, 2006-03-06 The development and application of bioinformatics tools to basic and translational cancer research is in fact a rapidly expanding field that deserves a timely review Therefore a publication of this type is needed The editors have done an excellent job in recruiting well established scientists to author the various chapters of the book Dieter Naf Jackson Laboratory USA Cancer bioinformatics is now emerging as a new interdisciplinary field which is facilitating an unprecedented synthesis of knowledge arising from the life and clinical sciences This groundbreaking title provides a comprehensive and up to date account of the enormous range of bioinformatics for cancer therapy development from the laboratory to clinical trials It functions as a guide to integrated data exploitation and synergistic knowledge discovery and support the consolidation of the interdisciplinary research community involved

Pharmacoinformatics: New developments and challenges in drug design José L. Medina-Franco, David Ramírez, Rafael Peláez, Carlos F. Lagos, 2024-08-08 Pharmacoinformatics combines bio and chemoinformatics approaches as well as artificial intelligence to support drug design and development at various stages starting from preclinical research support to clinical trial design and execution support stages I II and III as well as pharmacovigilance pharmacoeconomics and personalized medicine The rational use of these methods combined with the relevant experimental approaches is crucial

to face the new challenges in drug design and development The Biotech Investor's Bible George Wolff, 2001-06-13 A one stop source for investing in biotech with detailed coverage of the science the business the players and the strategies for one of today's most promising and volatile industries To invest in biotech is to invest in the future and as such investors need to learn the nuances of the science they're putting their money on The core asset of biotech companies is knowledge and sound investment decisions are impossible without an understanding of this complex science That's where The Biotech Investor's Bible fits in This much needed one of a kind resource simplifies the complex science surrounding the business of biotech and clarifies subtle distinctions within the context of their financial repercussions The book explains the basics of genetics patents and therapies and teaches investors how to value biotech companies and their state of the art products and technology The Biotech Investor's Bible offers an informative summary of the relatively short history of the industry and provides a comprehensive review of various industry sectors

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