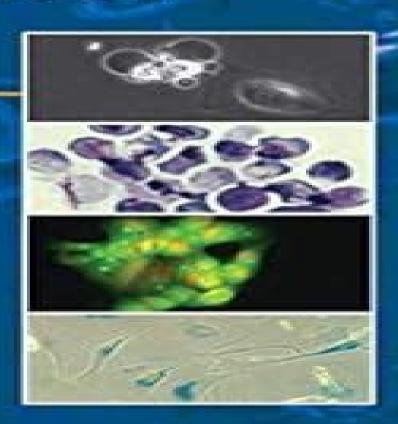
Apoptosis, Senescence, and Cancer

SECOND EDITION

Edited by

David A. Gewirtz Shawn E. Holt Steven Grant



Judit Hohmann, Anna Karolina Kiss, Judith Maria Rollinger, Atanas G. Atanasov

Apoptosis, Senescence and Cancer David A. Gewirtz, Shawn E. Holt, Steven Grant, 2007-12-17 Provides insight into established practices and research into apoptosis and senescence by examining techniques and research in the fields of cell death pathways senescence growth arrest drugs and resistance DNA damage response and other topics which still hold mysteries for researchers This book concludes with established cancer therapies Modeling and Parameter Estimation for Heterogeneous Cell Populations Jan Hasenauer, 2013 Most of the modeling performed in biology aims at achieving a quantitative description and understanding of the intracellular signaling pathways within a typical cell However in many biologically important situations even genetically identical cell populations show a heterogeneous response This means that individual members of the cell population behave differently Such situations require the study of cell to cell variability and the development of models for heterogeneous cell populations. The main contribution of this thesis is the development of unifying modeling frameworks for signal transduction and proliferation processes in heterogeneous cell populations These modeling frameworks allow for the detailed description of individual cells as well as differences between them In contrast to many existing modeling approaches the proposed frameworks allow for a direct comparison of model predictions with available data Beyond this the proposed population models can be simulated efficiently and by exploiting the model structures we are able to develop model tailored Bayesian parameter estimation methods. These methods enable the calculation of the optimal parameter estimates as well as the evaluation of the parameter and prediction uncertainties. The proposed tools allow for novel insights in population dynamics in particular the model based characterization of population heterogeneity and cellular subgroups This is illustrated for two different application examples pro and anti apoptotic signaling which is interesting in the context of cancer therapy and immune cell proliferation Transforming Growth Factor-Beta in Cancer Therapy, Volume I Sonia B. Jakowlew, 2010-01-01 Transforming Growth Factor il in Cancer Therapy Volume I Basicand Clinical Biology The present volume brings together a wealth of information that is fundamental to understanding the roleofTGF in the pathogenesis prevention and treatment of cancer It is not even 25 years sinceTGF was first isolated and characterized as a dimeric pep tide from both human and bovine sources 1 3 but the entire fieldofTGF research has grown and expanded so that it is now a central theme in all of cell biology There is almost no tissue or organ in the mammalian body in whichTGF does not playa central role in embryonic differentiation or in adult function and furthermore malfunction of the normal physiologyofTGF can have disastrous consequences in almost all ofthese sites Therefore the present comprehensive review of so many aspects of TGF function is a most welcome attempt to bring together a huge body of experimental data that is of the utmost importance in the field of oncology Frontiers in Anti-Cancer Drua Discovery Volume 10 Atta-ur-Rahman ,M. Igbal Choudhary, 2019-05-31 Frontiers in Anti Cancer Drug Discovery is a book series devoted to publishing the latest advances in anti cancer drug design and discovery In each volume eminent scientists

contribute reviews relevant to all areas of rational drug design and drug discovery including medicinal chemistry in silico drug design combinatorial chemistry high throughput screening drug targets recent important patents and structure activity relationships The book series should prove to be of interest to all pharmaceutical scientists involved in research in anti cancer drug design and discovery The book series is essential reading to all scientists involved in drug design and discovery who wish to keep abreast of rapid and important developments in the field The tenth volume of the series features chapters covering the following topics Challenges in the Management of Hepatoblastoma The Emerging Role of Monocarboxylate Transporter 1 in Cancer In vitro Anti Proliferative Assays and Techniques Used in Pre Clinical Anti Cancer Drug Discovery Recent Advances in the Development of Mesoporous Anti Cancer Drug Nanocarriers Polyphenols and Cancer Glioblastoma Multiforme Cutting Edge Targeting Strategies Utilizing Nanotechnology in Breast Cancer Therapy Discovery Kyu-Won Kim, Jae Kyung Roh, Hee-Jun Wee, Chan Kim, 2016-11-14 The reader will discover a comprehensive and multifaceted overview of the history of the development of anticancer drugs deeply influenced by the cell concept of cancer and future directions for the development of new anticancer drugs First this book documents the scientific progress in biological science over the last 70 years and the influence this progress had in cancer research Summaries and charts of important discoveries complete this overview Furthermore this book outlines the process of anticancer drug development with a focus on the characteristic drug groups of each era related to advancements of chemistry and biological sciences This book also provides brief mechanism of action of drugs illustrated by comprehensive timelines and conceptual cartoons This book finally sums up the limitations of the current anticancer drug development and seeks new directions for anticancer drug discovery considering under the systemic view of cancer Epigenetic Approaches in Drug Discovery, Development and <u>Treatment</u> Shibashish Giri, Chandravanu Dash, 2020-08-07 Establishment of a normal phenotype involves dynamic epigenetic regulation of gene expression that when affected contributes to human diseases On a molecular level epigenetic regulation is marked by specific covalent modifications acetylation methylation phosphorylation sumoylation PARylation and ubiquitylation of DNA and its associated histones Studies also suggest the influence of such epigenetic modifications on non coding RNA expression implicated in normal and diseased phenotypes Epigenetic control of genetic expression is a reversible process essential for normal development and function of an organism Alteration of epigenetic regulation leads to various disease forms such as cancer diabetes inflammation and neuropsychiatric disorders Assessing these alterations provides a deeper insight into the changes induced in the genome which is often informative for identifying disease subtypes or developing suitable treatments Therefore epigenetics proves to be a key area of clinical investigation in diagnosis prognosis and treatment of complex diseases Genetic mutations environmental stress pathogens and drugs of abuse are some of the predominant factors that induce and impact changes on chromatin which directly dictate a diseased phenotype It is essential to consider the interaction between genetic and epigenetic factors to understand the molecular mechanisms of complex

human diseases for safer and efficient drug development Furthermore genetic variation in absorption distribution metabolism and excretion ADME genes is insufficient to account for interindividual variability of drug response Therefore current efforts aim to identify epigenetic components of ADME gene regulation which include phase I and phase II enzymes uptake transporters efflux transporters and nuclear receptors involved in regulation of ADME genes Monitoring circulatory epigenetic biomarkers in liquid biopsies blood saliva urine cerebrospinal fluid of disease associated and drug associated epigenetic alterations may prove useful for decision support for routine clinical treatment and drug discovery Hence recent drug discovery efforts on targeting the epigenome has emerged an area of interest with several new drugs being developed tested and some already approved by the US Food and Drug Administration FDA These new insights into the complexities of epigenetic regulation are key contributors to our basic understanding of this process in human health and disease which will provide scope for innovative drug therapies It is of urgency to aid the present understanding of epigenomics driven diseased outcomes with the expectation that further studies will identify early markers of disease and targets for therapeutics

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