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Pediatric Inflammatory Bowel Disease Petar Mamula,Jonathan E. Markowitz,Robert N. Baldassano,2012-12-14 Pediatric Inflammatory Bowel Disease Second Edition provides an essential reference with an emphasis on the unique pediatric issues of IBD Chapters focus on complications of IBD specific to children and adolescents Treatment recommendations are based on the latest clinical research available The textbook also presents sections dedicated to the aspects of participation in clinical research unique to children and adolescents and the complicated yet vital process of successfully transitioning a patient from a pediatric to adult specialist Controversies in pediatric IBD care such as the off label use of medications are also covered The format incorporates multiple tables graphs and figures to improve readability and make for an efficient reference for clinicians to use Thoroughly revised and updated from the first edition the volumes includes new therapies that are currently being used or tested for treatment of IBD important areas regarding incidence and prevalence immunization and response to vaccine administration as well as advancements in our understanding of growth and development with particular to the use of growth hormone therapy Other new areas covered include important topics of complementary and alternative medicine use in IBD immunization and liver disease in IBD Pediatric Inflammatory Bowel Disease Second Edition is a valuable resource for pediatric gastroenterologists as well as adult gastroenterologists

Evidence-Based Gastroenterology and Hepatology John W. D. McDonald,Andrew K. Burroughs,Brian G. Feagan,2008-04-15 Traditional textbooks in this field have emphasized the basic sciences of pathology biochemistry and physiology Evidence based Gastroenterology and Hepatology covers all the major diseases of the gastrointestinal tract and liver utilizing clinical epidemiology to present the strongest and most current evidence for interventions This second edition is edited and written by leading gastroenterologists from around the world each chapter summarizes the evidence so that better informed decisions will be made about which treatments to offer to patients It provides practising Gastroenterologists and Surgeons with clear information regarding the diagnosis and treatment of pancreatic diseases giving clear evidence and experience based material that is immediately relevant to clinical practice Also contains a list of recommended reading at the end of each chapter Take a look at up to date information at www evidbasedgastro com **The science and art of value in healthcare: Measures, voices and methods** Jon Arsen Chilingerian,Laura Lorenz,Gala True,Stephen E. Chick,Christopher Tompkins,2022-12-29 **Pediatric Endoscopy, An Issue of Gastrointestinal Endoscopy Clinics, E-Book** Catharine M. Walsh,2023-03-26 In this issue of Gastrointestinal Endoscopy Clinics of North America guest editor Dr Catharine M Walsh brings her considerable expertise to the topic of Pediatric Endoscopy Since its inception in the early 1980s pediatric endoscopy has seen dramatic innovations in both diagnostic procedures and advanced procedures that are increasingly being performed by pediatric trained endoscopists In this issue top experts bring you fully up to date with recent advances in this fast changing field Contains 14 practice oriented topics including pediatric unsedated transnasal endoscopy tools for

improving quality in pediatric endoscopy artificial intelligence in pediatric endoscopy advances in endoscopy for pediatric inflammatory bowel disease and more Provides in depth clinical reviews on pediatric endoscopy offering actionable insights for clinical practice Presents the latest information on this timely focused topic under the leadership of experienced editors in the field Authors synthesize and distill the latest research and practice guidelines to create clinically significant topic based reviews **The Software Encyclopedia 2001** ,2001 *Adweek* ,1999 National Library of Medicine Audiovisuals Catalog National Library of Medicine (U.S.), *Drug Intelligence & Clinical Pharmacy* ,1987 **Logical Modeling of Cellular Processes: From Software Development to Network Dynamics** Matteo Barberis,Tomáš Helikar,2019-08-16 Mathematical models have become invaluable tools for understanding the intricate dynamic behavior of complex biochemical and biological systems Among computational strategies logical modeling has been recently gaining interest as an alternative approach to address network dynamics Due to its advantages including scalability and independence of kinetic parameters the logical modeling framework is becoming increasingly popular to study the dynamics of highly interconnected systems such as cell cycle progression T cell differentiation and gene regulation Novel tools and standards have been developed to increase the interoperability of logical models which can now be employ to respond a variety of biological questions This Research Topic brings together the most recent and cutting edge approaches in the area of logical modeling including among others novel biological applications software development and model analysis techniques Revolutionizing Immunological Disease Understanding Through Single Cell Multi-Omics Technologies Jennie R. Lill,Veronica Anania,Ying Zhu,Zora Modrusan,2025-06-17 Single cell technologies encompassing genomics and proteomics are at the forefront of revolutionizing our understanding of biological mechanisms at the cellular level particularly in the context of immunological diseases These technologies have enabled researchers to dissect complex cellular processes with unprecedented detail providing crucial spatial context that was previously unattainable Recent advancements have led to the formulation of new signaling hypotheses and opened novel pathways for therapeutic interventions The integration of multi omics approaches which combine various single cell technologies has further enhanced our ability to explore intracellular signaling and disease pathology Despite these advancements challenges remain in optimizing data interpretation and understanding the full scope of these technologies impact on immunological and cancer immunological disease pathogenesis There is a pressing need for continued exploration and refinement of these methodologies to fully harness their potential in elucidating disease mechanisms Computer Aid in Gastroenterology P. Rozen,F. T. de Dombal,1984-04-26 **Construction and utilization of digital twins for personalized therapeutic predictions** Samuel Schäfer,2024-10-18 Modern healthcare faces a significant challenge namely that 25 70% of patients with common diseases do not benefit from standard treatments despite the availability of over 13 000 drugs registered in DrugBank This discrepancy is likely due to these diseases complex and heterogeneous molecular nature rather than a lack of therapeutic options Emerging technologies have revealed the immense

molecular complexity underlying common diseases For instance singlecell RNA sequencing scRNA seq has demonstrated altered gene interactions in and across multiple cell types in numerous tissues Furthermore these technologies have revealed vast molecular differences between patients with the same diagnosis There is a wide gap between this complexity and the current diagnostic and therapeutic approaches Aim To bring personalized medicine one step closer to the clinic this thesis focuses on developing digital disease models that can capture the molecular biological complexity of disease in individual patients We aim to harness these disease models to identify optimal treatments for each individual patient Paper I We started by exploring the usefulness of OMIC based approaches for diagnostic and therapeutic predictions Utilizing a single cell RNA sequenced mouse model of antigen induced arthritis we aimed to prioritize cell types and therapeutic targets Initial pathway enrichment analyses did not yield relevant prioritization prompting an investigation into network based approaches Multi cellular disease models MCDMs for AIA and human rheumatoid arthritis were constructed incorporating predicted cell type interactions Centrality analysis indicated that these interactions could quantify a cell type s relative importance in disease pathogenesis We hypothesized that transcriptomic alterations in central cell types might reflect the MCDM serving as potential diagnostic markers An analysis of CD4 T cells from patients with 13 different inflammatory diseases and healthy controls demonstrated that these profiles could discriminate between healthy and diseased states and among diseases Furthermore a network based approach identified drugs targeting disease associated changes common to multiple inflammatory diseases Notably one of these drugs bezafibrate successfully dampened inflammation in the AIA mouse model Paper II Building on the insights from Paper I we investigated multicellular network models MNMs with time as an additional dimension Using seasonal allergic rhinitis SAR as a disease model we analyzed time series scRNAseq data to construct MNMs of inflammatory diseases We identified thousands of disease associated expression changes across multiple cell types varying at different disease stages Notably upstream regulators URs of these changes were also stage dependent and multidirectional To prioritize URs for drug discovery we focused on those causing significant expression changes in multiple cell types across all time points This strategy was validated through similar analyses of atopic dermatitis ulcerative colitis and Crohn s disease confirming that ranked URs aligned with the efficacy of existing drugs targeting the URs in the respective diseases Furthermore experimental validation included targeting the top ranked regulatory gene in SAR which was more effective than previously discovered IL4 inhibition Paper III While Paper I established the use of transcriptomic data for therapeutic predictions it focused on overlapping disease related changes across multiple inflammatory diseases and considered transcriptomic changes in only one cell type Paper II indicated a potential benefit in UR prioritization in numerous cell types However it yielded heterogeneous results and was limited by the fact that few drugs directly target URs Neither of these approaches was feasible for individualized drug predictions Drawing on previous insights by us and others we next aimed to develop digital disease models for individual patients termed digital twins with the capability for drug

efficacy screening We proposed scDrugPrio a strategy utilizing single cell scRNA sequencing based multicellular disease models incorporating key biological and pharmacological properties such as varying gene expression levels varying gene interactions within and between cell types and drug effect scDrugPrio was constructed based on a mouse model of arthritis and validated by improved precision recall for known drugs and in vitro studies of predicted drugs that were FDA approved for other diseases and had not yet been tried in rheumatoid arthritis or mouse arthritis For validation scDrugPrio was applied to human multiple sclerosis as well as Crohn s disease data that included tissue samples from healthy and sick tissue of all patients scDrugPrio was able to identify relevant treatments for individual patients and could distinguish anti TNF responders from non responders Conclusion This thesis demonstrates a framework for constructing digital disease models for personalized therapeutic predictions that might hold potential for better clinical treatment decisions By leveraging advanced genome wide analyses and network based approaches we may enhance the precision and efficacy of treatments for immune mediated inflammatory diseases bringing personalized medicine closer to clinical reality *Supplement to the Official Journal of the European Communities* ,1995-10-17 **Complex, Intelligent, and Software Intensive Systems** Leonard Barolli,Olivier Terzo,2017-07-04 This book gathers the proceedings of the 11th International Conference on Complex Intelligent and Software Intensive Systems CISIS 2017 held on June 28 June 30 2017 in Torino Italy Software Intensive Systems are characterized by their intensive interaction with other systems sensors actuators devices and users Further they are now being used in more and more domains e g the automotive sector telecommunication systems embedded systems in general industrial automation systems and business applications Moreover the outcome of web services delivers a new platform for enabling software intensive systems Complex Systems research is focused on the understanding of a system as a whole rather than its components Complex Systems are very much shaped by the changing environments in which they operate and by their multiple internal and external interactions They evolve and adapt through internal and external dynamic interactions The development of Intelligent Systems and agents which invariably involves the use of ontologies and their logical foundations offers a fruitful impulse for both Software Intensive Systems and Complex Systems Recent research in the fields of intelligent systems robotics neuroscience artificial intelligence and cognitive sciences is essential to the future development of and innovations in software intensive and complex systems The aim of the volume Complex Intelligent and Software Intensive Systems is to provide a platform of scientific interaction between the three interwoven and challenging areas of research and development of future Information and Communications Technology ICT enabled applications Software Intensive Systems Complex systems and Intelligent Systems *Systems Engineering of Software-Enabled Systems* Richard E. Fairley,2019-06-17 A comprehensive review of the life cycle processes methods and techniques used to develop and modify software enabled systems Systems Engineering of Software Enabled Systems offers an authoritative review of the most current methods and techniques that can improve the links between systems engineering and software engineering The

author a noted expert on the topic offers an introduction to systems engineering and software engineering and presents the issues caused by the differences between the two during development process The book reviews the traditional approaches used by systems engineers and software engineers and explores how they differ The book presents an approach to developing software enabled systems that integrates the incremental approach used by systems engineers and the iterative approach used by software engineers This unique approach is based on developing system capabilities that will provide the features behaviors and quality attributes needed by stakeholders based on model based system architecture In addition the author covers the management activities that a systems engineer or software engineer must engage in to manage and lead the technical work to be done This important book Offers an approach to improving the process of working with systems engineers and software engineers Contains information on the planning and estimating measuring and controlling managing risk and organizing and leading systems engineering teams Includes a discussion of the key points of each chapter and exercises for review Suggests numerous references that provide additional readings for development of software enabled physical systems Provides two case studies as running examples throughout the text Written for advanced undergraduates graduate students and practitioners Systems Engineering of Software Enabled Systems offers a comprehensive resource to the traditional and current techniques that can improve the links between systems engineering and software engineering

Cumulated Index Medicus ,1979   Commerce Business Daily ,1999-10   *Cancer and inflammatory diseases research: from the basics to the precision medicine* Mariana Quezado,Whocely Victor de Castro,Masamitsu Kanada,Adriano Sabino,Elaine Leite,2024-10-07 Precision medicine or personalized medicine is related to treatment based on the patient s individual characteristics To promote this personalized approach diseases must accurately be diagnosed specific biomarkers can predict disease evolution and optimize therapy based on each patient s characteristics such as their genetic background lifestyle and environmental risk factors Over the years we have seen a revolution in pharmacotherapy and management of patients receiving personalized medicine especially when we consider immunotherapy and specific tyrosine kinases inhibitors Patients with inflammatory diseases such as monogenetic diabetes and chronic kidney disease have benefited from the progress made within the last decade related to our understanding of the genetic basis of diabetes Similarly patients in precision oncology with breast prostate and blood cancer have experienced better overall survival rates with targeted therapy such as trastuzumab breast cancer PARPs inhibitors prostate cancer imatinib ibrutinib and venetoclax leukemia

Molecular Microbiology David H. Persing,Fred C. Tenover,Randall T. Hayden,Margareta Ieven,Melissa B. Miller,Frederick S. Nolte,Yi-Wei Tang,Alex van Belkum,2020-07-24 Presenting the latest molecular diagnostic techniques in one comprehensive volume The molecular diagnostics landscape has changed dramatically since the last edition of Molecular Microbiology Diagnostic Principles and Practice in 2011 With the spread of molecular testing and the development of new technologies and their opportunities laboratory professionals and physicians more than ever need a resource to help them

navigate this rapidly evolving field Editors David Persing and Fred Tenover have brought together a team of experienced researchers and diagnosticians to update this third edition comprehensively to present the latest developments in molecular diagnostics in the support of clinical care and of basic and clinical research including next generation sequencing and whole genome analysis These updates are provided in an easy to read format and supported by a broad range of practical advice such as determining the appropriate type and quantity of a specimen releasing and concentrating the targets and eliminating inhibitors Molecular Microbiology Diagnostic Principles and Practice Presents the latest basic scientific theory underlying molecular diagnostics Offers tested and proven applications of molecular diagnostics for the diagnosis of infectious diseases including point of care testing Illustrates and summarizes key concepts and techniques with detailed figures and tables Discusses emerging technologies including the use of molecular typing methods for real time tracking of infectious outbreaks and antibiotic resistance Advises on the latest quality control and quality assurance measures Explores the increasing opportunities and capabilities of information technology Molecular Microbiology Diagnostic Principles and Practice is a textbook for molecular diagnostics courses that can also be used by anyone involved with diagnostic test selection and interpretation It is also a useful reference for laboratories and as a continuing education resource for physicians If you are looking for online access to the latest clinical microbiology content please visit www wiley com learn clinmicronow

Virtual Colonoscopy Judy Yee,2008 This text is a complete and practical guide to performing virtual colonoscopy examinations and interpreting the results The book provides precise instructions on how to prepare the patient and specific protocols for the various CT scanners currently in use Detailed chapters on interpretation demonstrate normal anatomy anatomic variants CT appearances of polyps and cancer and pitfalls of interpretation on two dimensional and three dimensional views Additional chapters cover extracolonic findings reporting and fecal and fluid tagging More than 700 illustrations show examination techniques and normal and pathologic findings A companion Website will offer teaching cases to further refine the reader s interpretive skills

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