

Translational Bioinformatics

Volume 14

Series editor

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Aims and Scope

The Book Series in Translational Bioinformatics is a powerful and integrative resource for understanding and translating discoveries and advances of genomic, transcriptomic, proteomic and bioinformatic technologies into the study of human diseases. The Series represents leading global opinions on the translation of bioinformatics sciences into both the clinical setting and descriptions to medical informatics. It presents the critical evidence to further understand the molecular mechanisms underlying organ or cell dysfunctions in human diseases, the results of genomic, transcriptomic, proteomic and bioinformatic studies from human tissues dedicated to the discovery and validation of diagnostic and prognostic disease biomarkers, essential information on the identification and validation of novel drug targets and the application of tissue genomics, transcriptomics, proteomics and bioinformatics in drug efficacy and toxicity in clinical research.

The Book Series in Translational Bioinformatics focuses on outstanding articles/chapters presenting significant recent works in genomic, transcriptomic, proteomic and bioinformatic profiles related to human organ or cell dysfunctions and clinical findings. The Series includes bioinformatics-driven molecular and cellular disease mechanisms, the understanding of human diseases and the improvement of patient prognoses. Additionally, it provides practical and useful study insights into and protocols of design and methodology.

Series Description

Translational bioinformatics is defined as the development of storage-related, analytic, and interpretive methods to optimize the transformation of increasingly voluminous biomedical data, and genomic data in particular, into proactive, predictive, preventive, and participatory health. Translational bioinformatics includes research on the development of novel techniques for the integration of biological and clinical data and the evolution of clinical informatics methodology to encompass biological observations. The end product of translational bioinformatics is the newly found knowledge from these integrative efforts that can be disseminated to a variety of stakeholders including biomedical scientists, clinicians, and patients. Issues related to database management, administration, or policy will be coordinated through the clinical research informatics domain. Analytic, storage-related, and interpretive methods should be used to improve predictions, early diagnostics, severity monitoring, therapeutic effects, and the prognosis of human diseases.

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ISSN 2213-2775

Translational Bioinformatics

ISBN 978-981-13-0619-8

<https://doi.org/10.1007/978-981-13-0620-4>

ISSN 2213-2783 (electronic)

ISBN 978-981-13-0620-4 (eBook)

Library of Congress Control Number: 2018948704

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Printed on acid-free paper

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The registered company address is: 152 Beach Road, #21-01/04 Gateway East, Singapore 189721, Singapore

Contents

1	Clinical Lipidomics: A Critical Approach for Disease Diagnosis and Therapy	1
	Xiangdong Wang	
2	The Role of Lipid Metabolism in the Development of Lung Cancer	7
	Lixin Wang, Weiling Huang, and Xiu-Min Li	
3	Bioinformatics of Embryonic Exposures: Lipid Metabolism and Gender as Biomedical Variables	21
	K. K. Linask	
4	An Evaluation of Multivariate Data Analysis Models for Lipidomic Parameters from Patients with Metabolic Syndrome Undergoing Remedial Treatment	39
	D. Farabos, C. Wolf, R. Chapier, A. Lamaziere, and Peter J. Quinn	
5	Lipidomics in Carotid Artery Stenosis: Further Understanding of Pathology and Treatment	55
	Wei Zhang, Xiushi Zhou, Daqiao Guo, Weiguo Fu, and Lixin Wang	
6	Metabolomics of Immunity and Its Clinical Applications	73
	Jing Qiu, Fangming Liu, and Duoqiao Wu	
7	Urinary Lipidomics	97
	Phornpimon Tiphara and Visith Thongboonkerd	
8	Breast Cancer and Lipid Metabolism	113
	Chunfa Huang, Yuntao Li, Yifan Tu, and Carl E. Freter	
9	Association of Circulating Oxidized Lipids with Cardiovascular Outcomes	137
	Irena Levitan, Ibra S. Fancher, and Evgeny Berdyshev	

10 Lipidomics: Mass Spectrometry Based Untargeted Profiling and False Positives.....	155
Xiaohui Liu, Lina Xu, Xueying Wang, and Yupei Jiao	
11 Phospholipid and Phospholipidomics in Health and Diseases.....	177
Tanxi Cai and Fuquan Yang	
Index.....	203