



Integrating Omics Data

George Tseng, Debashis Ghosh, Xianghong Jasmine Zhou

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In most modern biomedical research projects, application of high-throughput genomic, proteomic and transcriptomic experiments has gradually become an inevitable component. Popular technologies include microarray and next-generation sequencing such as CHiP and RNA-Seq. As the technologies have become mature and the price affordable, omics data are rapidly generated and the problem of information integration and modeling of multi-lab and/or multi-omics data is becoming a growing one in the bioinformatics field. This book provides comprehensive coverage of these topics, and will have a long-lasting impact on this evolving subject. Each chapter, written by a leader in the field, introduces state-of-the-art methods to handle information integration, experimental data, and database problems of omics data.

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