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3. Genome-Scale Algorithm Design (Biological
sequence analysis in the era of high-
throughput sequencing) by Veli Mäkinen,
Djamal Belazzougui, Fabio Cunial and
Alexandru I. Tomescu. A graduate-level text
about state-of-the-art data structures and
algorithms for modern sequence analysis,

Genome-Scale Algorithm Design - Veli Mäkinen,
Djamal ...

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*Using Genome-Scale Models to Predict
Biological Capabilities*

*Genome-Scale Algorithm Design: Biological
Sequence Analysis in the Era of High-
Throughput Sequencing. The chapters feature
numerous examples, algorithm visualisations,
exercises and problems, each chosen to
reflect the steps of large-scale sequencing
projects, including read alignment, variant
calling, haplotyping, fragment assembly,...*

*Genome-Scale Algorithm Design: Biological
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*Genome scale metabolic models have emerged as
a valuable tool for illustrating whole cell
function, based on a complete set of
reactions of biochemical networks. These
models are used for the prediction of
organism's behavior. All information we need
in this modeling is a list of biochemical
reactions and their stoichiometry .*

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Textbooks Required •Bioinformatics

Algorithms: An Active Learning Approach

Volume I (Compeau and Pevzner 2015)

•Bioinformatics Algorithms: An Active Learning Approach Volume II (Compeau and Pevzner 2015) Other great resources

•Biological Sequence Analysis (Durbin, Eddy, Krogh, Mitchinson 1998) •Genome Scale

Algorithm Design (Mäkinen, Belazzougui, Cunial,

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High-throughput sequencing has revolutionized the ?eld of biological sequence anal- ysis.

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The Molecular Biology and Evolution toolbox. Click here for a paper that describes the

tool. The Population Genetics and Evolution toolbox. The ms tool for generating samples

under neutral models. INTENDED AUDIENCE: Anyone interested in learning about

algorithms and their use in biological sequence analysis.

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Bottom-up approaches to systems biology rely on constructing a mechanistic basis for the biochemical and genetic processes that underlie cellular functions. Genome-scale network reconstructions of metabolism are built from all known metabolic reactions and metabolic genes in a target organism.

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M{"a}kinen and Djamal Belazzougui and Fabio
Cunial and Alexandru I. Tomescu}, year={2015
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High-throughput sequencing has revolutionised the field of biological sequence analysis. Its application has enabled researchers to address important biological questions, often for the first time. This book provides an integrated presentation of the fundamental algorithms and data structures that power modern sequence analysis workflows.

Genome-Scale Algorithm Design: Biological Sequence ...

Genome-Scale Algorithm Design provides an integrated presentation of the fundamental algorithms and data structures that power modern sequence analysis workflows. The topics covered range from the foundations of biological sequence analysis (alignments and hidden Markov models), to classical index structures (k -mer indexes, suffix arrays and suffix trees), Burrows-Wheeler indexes, graph algorithms and a number of advanced omics applications.

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