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| Altri autori (Persone) | VinaTomás |
| Disciplina | 570.285 |
| Soggetti | Biomathematics Bioinformatics Algorithms Computer networks Computer science - Mathematics Application software Mathematical and Computational Biology Computational and Systems Biology Design and Analysis of Algorithms Computer Communication Networks Mathematics of Computing Computer and Information Systems Applications |
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| Nota di contenuto | Classifying the Post-Duplication Fate of Paralogous Genes -- Inferring clusters of orthologous and paralogous transcripts -- On the class of double distance problems -- The Floor is Lava - Halving Genomes with Viaducts, Piers and Pontoon -- Two strikes against the phage recombination problem -- Physical mapping of two nested fixed inversions in the X chromosome of the malaria mosquito Anopheles messeae -- Gene order phylogeny via ancestral genome reconstruction under Dollo -- Prior Density Learning in Variational Bayesian Phylogenetic Parameters Inference -- The Asymmetric Cluster Affinity Cost -- The k-RF Measures for Labeled Trees -- Bounding the Number |

of Reticulations in a Tree-Child Network That Displays a Set of Trees --
Finding agreement cherry-reduced subnetworks in level-1 networks --
CONSULT-II: Taxonomic identification using Locality Sensitive Hashing
-- MAGE: Strain Level Profiling of Metagenome Samples -- MoTERNN:
Classifying the Mode of Cancer Evolution Using Recursive Neural
Networks.

Sommario/riassunto

This book constitutes the refereed proceedings of the 20th Annual RECOMB Satellite Workshop on Comparative Genomics, RECOMB-CG 2023 which took place in Istanbul, Turkey, in April 2023. The 15 full papers included in this book were carefully reviewed and selected from 25 submissions. The papers present cutting edge research in comparative genomics, with an emphasis on computational approaches and novel experimental results. Chapters "Inferring Clusters of Orthologous and Paralogous Transcripts" and "Gene Order Phylogeny via Ancestral Genome Reconstruction under Dollo" are published Open Access under Creative Commons Attribution license (CC BY 4.0). .
