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Machine learning and data mining for yeast functional genomics

Machine learning and data mining for yeast functional genomics Amanda Clare Department of Computer Science University of Wales Aberystwyth February 2003 This thesis is submitted in partial fulfilment of the requirements for the degree of Doctor of Philosophy of The University of Wales

The Yeast Deletion Collection: A Decade of Functional Genomics

The Yeast Deletion Collection: A Decade of Functional Genomics Guri Giaever and Corey Nislow University of British Columbia, Vancouver, British Columbia, Canada V6T1Z3 ABSTRACT The yeast deletion collections comprise 21,000 mutant strains that carry precise start-to-stop deletions of 6000 open reading frames

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functional genomics and proteomics in yeast have appeared, but no comprehensive yeast-based functional genomics and proteomics textbook has yet been written and published This book aims to be the standard textbook in the field of yeast-based functional genomics and proteomics and should serve as a stand-alone protocols handbook suitable for daily

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TAR Cloning: Perspectives for Functional Genomics ...

of a circular molecule in a YAC (yeast artificial chromosome) form containing the desired genomic fragment carrying a gene or locus When TAR cloning is applied to complex genomes, such as the human genome, the technique produces YAC clones containing the desired genomic fragment at a frequency of 0.5%–2% of all yeast colonies screened

High-dimensional and large-scale phenotyping of yeast mutants

enabling a functional assignment of a locus to a specific cellular pathway The high-dimensional phenotypic analysis of defined yeast mutant strains provides another step toward attributing gene function to all of the genes in the yeast genome cell morphology functional genomics high-dimensional phenotyping phenome O

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Integrating large-scale functional genomic data to dissect ...

Integrating large-scale functional genomic data to dissect the complexity of yeast regulatory networks Jun Zhu¹, Bin Zhang¹, Erin N Smith^{2,3}, Becky Drees⁴, Rachel B Brem⁵, Leonid Kruglyak², Roger E Bumgarner⁴ & Eric E Schadt¹ A key goal of biology is to construct networks that ...

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Chapter 7

The development of this pooled fitness assay has greatly facilitated the functional annotation of the yeast genome by making genome-wide gene-deletion studies faster and easier, and has led to the development of high throughput methods for studying drug action in yeast

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Fungal Genomics, Computational Biology and Systems Biology Crop Genetics and Genomics The goal of this nine-week US Dept of Agriculture REEU (the extra 'E' is for extension) is to develop plant scientists that can address the challenges facing US agricultural competitiveness and food production