

1. Record Nr.	UNINA9910441060203321
Autore	Pinto Guerra, Edgardo
Titolo	Risanamento di murature umide e degradate : sintomi e cause, rimedi, soluzioni progettuali : per risultati duraturi / Edgardo Pinto Guerra ; [presentazione di Giovanni Carbonara]
Pubbl/distr/stampa	[Palermo] : D. Flaccovio, 2020
ISBN	978-88-579-1057-4
Edizione	[5. ed]
Descrizione fisica	254 p. : ill. ; 24 cm
Locazione	FINBC
Collocazione	13 A 66 11
Lingua di pubblicazione	Italiano
Formato	Materiale a stampa
Livello bibliografico	Monografia

2. Record Nr.	UNISA996466257903316
Titolo	Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics [[electronic resource]] : 5th European Conference, EvoBIO 2007, Valencia, Spain, April 11-13, 2007, Proceedings // edited by Elena Marchiori, Jason H. Moore, Jagath C. Rajapakse
Pubbl/distr/stampa	Berlin, Heidelberg : , : Springer Berlin Heidelberg : , : Imprint : Springer, , 2007
ISBN	3-540-71783-8
Edizione	[1st ed. 2007.]
Descrizione fisica	1 online resource (XI, 302 p.)
Collana	Theoretical Computer Science and General Issues, , 2512-2029 ; ; 4447
Disciplina	572.80285/6
Soggetti	Artificial intelligence Computer programming Computer science Algorithms Bioinformatics Pattern recognition systems Artificial Intelligence Programming Techniques Theory of Computation Computational and Systems Biology Automated Pattern Recognition
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Bibliographic Level Mode of Issuance: Monograph
Nota di bibliografia	Includes bibliographical references and index.
Nota di contenuto	Identifying Regulatory Sites Using Neighborhood Species -- Genetic Programming and Other Machine Learning Approaches to Predict Median Oral Lethal Dose (LD50) and Plasma Protein Binding Levels (% PPB) of Drugs -- Hypothesis Testing with Classifier Systems for Rule-Based Risk Prediction -- Robust Peak Detection and Alignment of nanoLC-FT Mass Spectrometry Data -- One-Versus-One and One-Versus-All Multiclass SVM-RFE for Gene Selection in Cancer Classification -- Understanding Signal Sequences with Machine Learning -- Targeting Differentially Co-regulated Genes by Multiobjective and Multimodal Optimization -- Modeling Genetic

Networks: Comparison of Static and Dynamic Models -- A Genetic Embedded Approach for Gene Selection and Classification of Microarray Data -- Modeling the Shoot Apical Meristem in *A. thaliana*: Parameter Estimation for Spatial Pattern Formation -- Evolutionary Search for Improved Path Diagrams -- Simplifying Amino Acid Alphabets Using a Genetic Algorithm and Sequence Alignment -- Towards Evolutionary Network Reconstruction Tools for Systems Biology -- A Gaussian Evolutionary Method for Predicting Protein-Protein Interaction Sites -- Bio-mimetic Evolutionary Reverse Engineering of Genetic Regulatory Networks -- Tuning ReliefF for Genome-Wide Genetic Analysis -- Dinucleotide Step Parameterization of Pre-miRNAs Using Multi-objective Evolutionary Algorithms -- Amino Acid Features for Prediction of Protein-Protein Interface Residues with Support Vector Machines -- Predicting HIV Protease-Cleavable Peptides by Discrete Support Vector Machines -- Inverse Protein Folding on 2D Off-Lattice Model: Initial Results and Perspectives -- Virtual Error: A New Measure for Evolutionary Bioclustering -- Characterising DNA/RNA Signals with Crisp Hypermotifs: A Case Study on Core Promoters -- Evaluating Evolutionary Algorithms and Differential Evolution for the Online Optimization of Fermentation Processes -- The Role of a Priori Information in the Minimization of Contact Potentials by Means of Estimation of Distribution Algorithms -- Classification of Cell Fates with Support Vector Machine Learning -- Reconstructing Linear Gene Regulatory Networks -- Individual-Based Modeling of Bacterial Foraging with Quorum Sensing in a Time-Varying Environment -- Substitution Matrix Optimisation for Peptide Classification.

3. Record Nr.	UNINA9910813980503321
Autore	Attebery Rachel <1990->
Titolo	The Bluebeam guidebook : game-changing tips and stories for architects, engineers, and contractors // Rachel Attebery, Jason Hascall
Pubbl/distr/stampa	Hoboken, New Jersey : , : Wiley, , 2018 ©2018
ISBN	1-119-39395-7 1-119-39396-5 1-119-39397-3
Descrizione fisica	1 online resource (319 pages) : illustrations
Classificazione	TEC005000
Disciplina	720.285/53
Soggetti	Building information modeling - Computer programs PDF (Computer file format)
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Includes index.
Sommario/riassunto	"Expert tips for the last piece in the paperless puzzle The Bluebeam Guidebook offers comprehensive coverage of the industry's leading PDF tool to help AEC professionals adopt a more efficient digital workflow. With desktop, mobile, and server-based products, Bluebeam makes collaboration and document coordination seamless, and provides a perfect complement to BIM software. This book shows you how to push the boundaries and discover the software's true capabilities. Written expressly for working AEC professionals, this book offers tips, tricks, and ideas that cater to industry-specific needs. Expert instruction and step-by-step guidance helps you get started quickly, and case studies feature users from firms such as Kiewit, Populus, Sundt Construction, and more to show you how Bluebeam is quickly becoming a critical component of design and construction. Master the industry's leading PDF software and alternative to Adobe Acrobat Create, edit, and markup documents in a way that suits the architecture and engineering workflow Learn how major AEC firms have transitioned seamlessly to digital workflows Integrate Bluebeam into estimating, quality control, field applications,

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"This book offers comprehensive coverage of the industry's leading PDF tool to help AEC professionals adopt a more efficient digital workflow. With desktop, mobile, and server-based products, Bluebeam makes collaboration and document coordination seamless, and provides a perfect complement to BIM software. This book shows you how to push the boundaries and discover the software's true capabilities"--
