

Список литературы

- [1] L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, editors, *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, volume 7833 of *LNCS*, Vienna, Austria, 2013, Springer Verlag.
- [2] D. GRANIZO-MACKENZIE and J. H. MOORE, Multiple Threshold Spatially Uniform ReliefF for the Genetic Analysis of Complex Human Diseases, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 1–10, Vienna, Austria, 2013, Springer Verlag.
- [3] J. TAN, G. GRANT, M. WHITFIELD, and C. GREENE, Time-point Specific Weighting Improves Coexpression Networks from Time-course Experiments, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 11–22, Vienna, Austria, 2013, Springer Verlag.
- [4] C. DARABOS, K. DESAI, R. COWPER-SALLARI, M. GIACOBINI, B. E. GRAHAM, M. LUPIEN, and J. H. MOORE, Inferring Human Phenotype Networks from Genome-Wide Genetic Associations, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 23–34, Vienna, Austria, 2013, Springer Verlag.
- [5] R. M. SIVLEY, A. E. FISH, and W. S. BUSH, Knowledge-constrained K-medoids Clustering of Regulatory Rare Alleles for Burden Tests, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 35–42, Vienna, Austria, 2013, Springer Verlag.
- [6] S. AHMED, M. ZHANG, and L. PENG, Feature Selection and Classification of High Dimensional Mass Spectrometry Data: A Genetic Programming Approach, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 43–54, Vienna, Austria, 2013, Springer Verlag.
- [7] P. A. WHIGHAM, G. DICK, A. WRIGHT, and H. G. SPENCER, Structured populations and the maintenance of sex, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 55–66, Vienna, Austria, 2013, Springer Verlag.
- [8] D. L. GONZALEZ-ALVAREZ and M. A. VEGA-RODRIGUEZ, Hybrid Multiobjective Artificial Bee Colony with Differential Evolution Applied to Motif Finding, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 67–78, Vienna, Austria, 2013, Springer Verlag.
- [9] K. SALAMA and A. FREITAS, Dimensionality reduction via Isomap with lock-step and elastic measures for time series gene expression classification, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 79–90, Vienna, Austria, 2013, Springer Verlag.
- [10] C. ORSENIGO and C. VERCELLIS, Dimensionality reduction via Isomap with lock-step and elastic measures for time series gene expression classification, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 91–102, Vienna, Austria, 2013, Springer Verlag.

- [11] Q. PAN, T. HU, J. D. MALLEY, A. S. ANDREW, M. R. KARAGAS, and J. H. MOORE, Supervising Random Forest Using Attribute Interaction Networks, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 103–114, Vienna, Austria, 2013, Springer Verlag.
- [12] N. SHARMA and T. GEDEON, Hybrid Genetic Algorithms for Stress Recognition in Reading, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 115–126, Vienna, Austria, 2013, Springer Verlag.
- [13] A. SULOVARI, J. KIRALIS, and J. H. MOORE, Optimal Use of Biological Expert Knowledge from Literature Mining in Ant Colony Optimization for Analysis of Epistasis in Human Disease, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 127–138, Vienna, Austria, 2013, Springer Verlag.
- [14] S. SANTANDER-JIMENEZ and M. A. VEGA-RODRIGUEZ, A Multiobjective Proposal Based on the Firefly Algorithm for Inferring Phylogenies, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 139–150, Vienna, Austria, 2013, Springer Verlag.
- [15] D. CASTALDI, D. MACCAGNOLA, D. MARI, and F. ARCHETTI, Mining for Variability in the Coagulation Pathway: A Systems Biology Approach, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 151–162, Vienna, Austria, 2013, Springer Verlag.
- [16] T. MANNING and P. WALSH, Improving the Performance of CGPANN for Breast Cancer Diagnosis using Crossover and Radial Basis Functions, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 163–174, Vienna, Austria, 2013, Springer Verlag.
- [17] M. GAUDES, A. MARION, T. MUSNER, G. SQUILLERO, and A. TONDA, An Evolutionary Approach to Wetlands Design, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 175–185, Vienna, Austria, 2013, Springer Verlag.
- [18] S. ROSENTHAL, N. EL-SOURANI, and M. BORSCHBACH, Impact of Different Recombination Methods in a Mutation-Specific MOEA for a Biochemical Application, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 186–197, Vienna, Austria, 2013, Springer Verlag.
- [19] J. M. FISHER, P. ANDREWS, J. KIRALIS, N. A. SINNOTT-ARMSTRONG, and J. H. MOORE, Alternative Cell-based Metrics Improve the Detection of Multifactor Dimensionality Reduction, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 198–209, Vienna, Austria, 2013, Springer Verlag.
- [20] M. ZAGORSKI, Emergence of motifs in model gene regulatory networks, in *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, edited by L. VANNESCHI, W. S. BUSH, and M. GIACOBINI, volume 7833 of *LNCS*, pp. 210–213, Vienna, Austria, 2013, Springer Verlag.