Список литературы

- 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. — V. 7833 of LNCS, Vienna, Austria, 2013. — April 3-5. Springer Verlag.
- [2] Ahmed, S. Feature Selection and Classification of High Dimensional Mass Spectrometry Data: A Genetic Programming Approach / S. Ahmed, M. Zhang, L. Peng // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 43-54.
- [3] Alternative Cell-based Metrics Improve the Detection of Multifactor Dimensionality Reduction / J. M. Fisher, P. Andrews, J. Kiralis et al. // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 198–209.
- [4] An Evolutionary Approach to Wetlands Design / M. Gaudesi, A. Marion, T. Musner et al. // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. — V. 7833 of LNCS. — Vienna, Austria: Springer Verlag, 2013. — April 3-5. — P. 175–185.
- [5] Gonzalez-Alvarez, D. L. Hybrid Multiobjective Artificial Bee Colony with Differential Evolution Applied to Motif Finding / D. L. Gonzalez-Alvarez, M. A. Vega-Rodriguez // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. — V. 7833 of LNCS. — Vienna, Austria: Springer Verlag, 2013. — April 3-5. — P. 67–78.
- [6] Granizo-Mackenzie, D. Multiple Threshold Spatially Uniform ReliefF for the Genetic Analysis of Complex Human Diseases / D. Granizo-Mackenzie, J. H. Moore // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. — V. 7833 of LNCS. — Vienna, Austria: Springer Verlag, 2013. — April 3-5. — P. 1-10.
- [7] Inferring Human Phenotype Networks from Genome-Wide Genetic Associations / C. Darabos, K. Desai, R. Cowper-Sallari et al. // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 23–34.
- [8] Manning, T. Improving the Performance of CGPANN for Breast Cancer Diagnosis using Crossover and Radial Basis Functions / T. Manning, P. Walsh // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 163–174.
- [9] Mining for Variability in the Coagulation Pathway: A Systems Biology Approach / D. Castaldi, D. Maccagnola, D. Mari, F. Archetti // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 151–162.
- [10] Orsenigo, C. Dimensionality reduction via Isomap with lock-step and elastic measures for time series gene expression classification / C. Orsenigo, C. Vercellis // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 91–102.

- [11] Rosenthal, S. Impact of Different Recombination Methods in a Mutation-Specific MOEA for a Biochemical Application / S. Rosenthal, N. El-Sourani, M. Borschbach // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 186–197.
- [12] Salama, K. Dimensionality reduction via Isomap with lock-step and elastic measures for time series gene expression classification / K. Salama, A. Freitas // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 79–90.
- [13] Santander-Jimenez, S. A Multiobjective Proposal Based on the Firefly Algorithm for Inferring Phylogenies / S. Santander-Jimenez, M. A. Vega-Rodriguez // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. — V. 7833 of LNCS. — Vienna, Austria: Springer Verlag, 2013. — April 3-5. — P. 139–150.
- [14] Sharma, N. Hybrid Genetic Algorithms for Stress Recognition in Reading / N. Sharma, T. Gedeon // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 115–126.
- [15] Sivley, R. M. Knowledge-constrained K-medoids Clustering of Regulatory Rare Alleles for Burden Tests / R. M. Sivley, A. E. Fish, W. S. Bush // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. — V. 7833 of LNCS. — Vienna, Austria: Springer Verlag, 2013. — April 3-5. — P. 35–42.
- [16] Structured populations and the maintenance of sex / P. A. Whigham, G. Dick, A. Wright, H. G. Spencer // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 55–66.
- [17] Sulovari, A. Optimal Use of Biological Expert Knowledge from Literature Mining in Ant Colony Optimization for Analysis of Epistasis in Human Disease / A. Sulovari, J. Kiralis, J. H. Moore // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 127–138.
- [18] Supervising Random Forest Using Attribute Interaction Networks / Q. Pan, T. Hu, J. D. Malley et al. // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 103—114.
- [19] Time-point Specific Weighting Improves Coexpression Networks from Time-course Experiments / J. Tan, G. Grant, M. Whitfield, C. Greene // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 11–22.
- [20] Zagorski, M. Emergence of motifs in model gene regulatory networks / M. Zagorski // 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 / ed. by L. Vanneschi, W. S. Bush, M. Giacobini. V. 7833 of LNCS. Vienna, Austria: Springer Verlag, 2013. April 3-5. P. 210–213.