## Список литературы

- [1] L. Vanneschi, W. S. Bush, and M. Giacobini, eds., 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, vol. 7833 of LNCS. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 1–10. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 11–22. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 23–34. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 35–42. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 43–54. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 55–66. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 67–78. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 79–90. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 91–102. Springer Verlag, Vienna, Austria, April 3-5, 2013.

- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 103–114. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 115–126. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 127–138. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 139–150. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 151–162. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 163–174. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [17] M. Gaudesi, 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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 175–185. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 186–197. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 198–209. Springer Verlag, Vienna, Austria, April 3-5, 2013.
- [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, L. Vanneschi, W. S. Bush, and M. Giacobini, eds., vol. 7833 of LNCS, pp. 210–213. Springer Verlag, Vienna, Austria, April 3-5, 2013.