Список литературы

[1] Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.) 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013 Vol. 7833 of LNCS, Vienna, Austria (April 3-5, 2013) Springer Verlag.

[2] Granizo-Mackenzie, D. and Moore, J. H. (April 3-5, 2013) Multiple Threshold Spatially Uniform ReliefF for the Genetic Analysis of Complex Human Diseases. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 1–10.

[3] Tan, J., Grant, G., Whitfield, M., and Greene, C. (April 3-5, 2013) Time-point Specific Weighting Improves Coexpression Networks from Time-course Experiments. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 11–22.

[4] Darabos, C., Desai, K., Cowper-Sallari, R., Giacobini, M., Graham, B. E., Lupien, M., and Moore, J. H. (April 3-5, 2013) Inferring Human Phenotype Networks from Genome-Wide Genetic Associations. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 23–34.

[5] Sivley, R. M., Fish, A. E., and Bush, W. S. (April 3-5, 2013) Knowledge-constrained K-medoids Clustering of Regulatory Rare Alleles for Burden Tests. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 35–42.

[6] Ahmed, S., Zhang, M., and Peng, L. (April 3-5, 2013) Feature Selection and Classification of High Dimensional Mass Spectrometry Data: A Genetic Programming Approach. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 43–54.

[7] Whigham, P. A., Dick, G., Wright, A., and Spencer, H. G. (April 3-5, 2013) Structured populations and the maintenance of sex. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 55–66.

[8] Gonzalez-Alvarez, D. L. and Vega-Rodriguez, M. A. (April 3-5, 2013) Hybrid Multiobjective Artificial Bee Colony with Differential Evolution Applied to Motif Finding. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 67–78.

[9] Salama, K. and Freitas, A. (April 3-5, 2013) Dimensionality reduction via Isomap with lock-step and elastic measures for time series gene expression classification. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 79–90.

[10] Orsenigo, C. and Vercellis, C. (April 3-5, 2013) Dimensionality reduction via Isomap with lock-step and elastic measures for time series gene expression classification. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 91–102.

[11] Pan, Q., Hu, T., Malley, J. D., Andrew, A. S., Karagas, M. R., and Moore, J. H. (April 3-5, 2013) Supervising Random Forest Using Attribute Interaction Networks. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 103–114.

[12] Sharma, N. and Gedeon, T. (April 3-5, 2013) Hybrid Genetic Algorithms for Stress Recognition in Reading. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 115–126.

[13] Sulovari, A., Kiralis, J., and Moore, J. H. (April 3-5, 2013) Optimal Use of Biological Expert Knowledge from Literature Mining in Ant Colony Optimization for Analysis of Epistasis in Human Disease. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 127–138.

[14] Santander-Jimenez, S. and Vega-Rodriguez, M. A. (April 3-5, 2013) A Multiobjective Proposal Based on the Firefly Algorithm for Inferring Phylogenies. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 139–150.

[15] Castaldi, D., Maccagnola, D., Mari, D., and Archetti, F. (April 3-5, 2013) Mining for Variability in the Coagulation Pathway: A Systems Biology Approach. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 151–162.

[16] Manning, T. and Walsh, P. (April 3-5, 2013) Improving the Performance of CGPANN for Breast Cancer Diagnosis using Crossover and Radial Basis Functions. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 163–174.

[17] Gaudesi, M., Marion, A., Musner, T., Squillero, G., and Tonda, A. (April 3-5, 2013) An Evolutionary Approach to Wetlands Design. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 175–185.

[18] Rosenthal, S., El-Sourani, N., and Borschbach, M. (April 3-5, 2013) Impact of Different Recombination Methods in a Mutation-Specific MOEA for a Biochemical Application. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 186–197.

[19] Fisher, J. M., Andrews, P., Kiralis, J., Sinnott-Armstrong, N. A., and Moore, J. H. (April 3-5, 2013) Alternative Cell-based Metrics Improve the Detection of Multifactor Dimensionality Reduction. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 198–209.

[20] Zagorski, M. (April 3-5, 2013) Emergence of motifs in model gene regulatory networks. In Vanneschi, L., Bush, W. S., and Giacobini, M., (eds.), *11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013*, Vienna, Austria: Springer Verlag Vol. 7833 of LNCS, pp. 210–213.