Список литературы

- [Ahmed et al., 2013] Ahmed, S.; Zhang, M.; and Peng, L. (2013). "Feature Selection and Classification of High Dimensional Mass Spectrometry Data: A Genetic Programming Approach". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 43–54, Vienna, Austria. Springer Verlag.
- [Castaldi et al., 2013] Castaldi, D.; Maccagnola, D.; Mari, D.; and Archetti, F. (2013). "Mining for Variability in the Coagulation Pathway: A Systems Biology Approach". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 151–162, Vienna, Austria. Springer Verlag.
- [Darabos et al., 2013] Darabos, C.; Desai, K.; Cowper-Sallari, R.; Giacobini, M.; Graham, B. E.; Lupien, M.; and Moore, J. H. (2013). "Inferring Human Phenotype Networks from Genome-Wide Genetic Associations". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 23–34, Vienna, Austria. Springer Verlag.
- [Fisher et al., 2013] Fisher, J. M.; Andrews, P.; Kiralis, J.; Sinnott-Armstrong, N. A.; and Moore, J. H. (2013). "Alternative Cell-based Metrics Improve the Detection of Multifactor Dimensionality Reduction". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 198–209, Vienna, Austria. Springer Verlag.
- [Gaudesi et al., 2013] Gaudesi, M.; Marion, A.; Musner, T.; Squillero, G.; and Tonda, A. (2013). "An Evolutionary Approach to Wetlands Design". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 175–185, Vienna, Austria. Springer Verlag.
- [Gonzalez-Alvarez and Vega-Rodriguez, 2013] Gonzalez-Alvarez, D. L. and Vega-Rodriguez, M. A. (2013). "Hybrid Multiobjective Artificial Bee Colony with Differential Evolution Applied to Motif Finding". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 67–78, Vienna, Austria. Springer Verlag.
- [Granizo-Mackenzie and Moore, 2013] Granizo-Mackenzie, D. and Moore, J. H. (2013). "Multiple Threshold Spatially Uniform ReliefF for the Genetic Analysis of Complex Human Diseases". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 1–10, Vienna, Austria. Springer Verlag.
- [Manning and Walsh, 2013] Manning, T. and Walsh, P. (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., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 163–174, Vienna, Austria. Springer Verlag.
- [Orsenigo and Vercellis, 2013] Orsenigo, C. and Vercellis, C. (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., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 91–102, Vienna, Austria. Springer Verlag.
- [Pan et al., 2013] Pan, Q.; Hu, T.; Malley, J. D.; Andrew, A. S.; Karagas, M. R.; and Moore, J. H. (2013). "Supervising Random Forest Using Attribute Interaction Networks". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 103–114, Vienna, Austria. Springer Verlag.

- [Rosenthal et al., 2013] Rosenthal, S.; El-Sourani, N.; and Borschbach, M. (2013). "Impact of Different Recombination Methods in a Mutation-Specific MOEA for a Biochemical Application". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 186–197, Vienna, Austria. Springer Verlag.
- [Salama and Freitas, 2013] Salama, K. and Freitas, A. (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., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 79–90, Vienna, Austria. Springer Verlag.
- [Santander-Jimenez and Vega-Rodriguez, 2013] Santander-Jimenez, S. and Vega-Rodriguez, M. A. (2013). "A Multiobjective Proposal Based on the Firefly Algorithm for Inferring Phylogenies". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 139–150, Vienna, Austria. Springer Verlag.
- [Sharma and Gedeon, 2013] Sharma, N. and Gedeon, T. (2013). "Hybrid Genetic Algorithms for Stress Recognition in Reading". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBio 2013, volume 7833 of LNCS, pages 115–126, Vienna, Austria. Springer Verlag.
- [Sivley et al., 2013] Sivley, R. M.; Fish, A. E.; and Bush, W. S. (2013). "Knowledge-constrained K-medoids Clustering of Regulatory Rare Alleles for Burden Tests". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 35–42, Vienna, Austria. Springer Verlag.
- [Sulovari et al., 2013] Sulovari, A.; Kiralis, J.; and Moore, J. H. (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., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 127–138, Vienna, Austria. Springer Verlag.
- [Tan et al., 2013] Tan, J.; Grant, G.; Whitfield, M.; and Greene, C. (2013). "Time-point Specific Weighting Improves Coexpression Networks from Time-course Experiments". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 11–22, Vienna, Austria. Springer Verlag.
- [Vanneschi et al., 2013] Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors (2013). 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, Vienna, Austria. Springer Verlag.
- [Whigham et al., 2013] Whigham, P. A.; Dick, G.; Wright, A.; and Spencer, H. G. (2013). "Structured populations and the maintenance of sex". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 55–66, Vienna, Austria. Springer Verlag.
- [Zagorski, 2013] Zagorski, M. (2013). "Emergence of motifs in model gene regulatory networks". In Vanneschi, L.; Bush, W. S.; and Giacobini, M., editors, 11th European Conference on Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics, EvoBIO 2013, volume 7833 of LNCS, pages 210–213, Vienna, Austria. Springer Verlag.