Список литературы

- [1] 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, Vienna, Austria, 2013, Springer Verlag.
- [2] GRANIZO-MACKENZIE, D. and MOORE, J. H., 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, pp. 1–10, Vienna, Austria, 2013, Springer Verlag.
- [3] TAN, J., GRANT, G., WHITFIELD, M., and GREENE, C., 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, pp. 11–22, Vienna, Austria, 2013, Springer Verlag.
- [4] DARABOS, C., DESAI, K., COWPER-SALLARI, R., et al., 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, pp. 23–34, Vienna, Austria, 2013, Springer Verlag.
- [5] SIVLEY, R. M., FISH, A. E., and BUSH, W. S., 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, pp. 35–42, Vienna, Austria, 2013, Springer Verlag.
- [6] AHMED, S., ZHANG, M., and PENG, L., 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, pp. 43–54, Vienna, Austria, 2013, Springer Verlag.
- [7] WHIGHAM, P. A., DICK, G., WRIGHT, A., and SPENCER, H. G., 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, pp. 55–66, Vienna, Austria, 2013, Springer Verlag.
- [8] GONZALEZ-ALVAREZ, D. L. and VEGA-RODRIGUEZ, M. A., 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, pp. 67–78, Vienna, Austria, 2013, Springer Verlag.
- [9] SALAMA, K. and FREITAS, A., 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, pp. 79–90, Vienna, Austria, 2013, Springer Verlag.
- [10] ORSENIGO, C. and VERCELLIS, C., 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, pp. 91–102, Vienna, Austria, 2013, Springer Verlag.

- [11] PAN, Q., HU, T., MALLEY, J. D., et al., 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, pp. 103–114, Vienna, Austria, 2013, Springer Verlag.
- [12] SHARMA, N. and GEDEON, T., 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, pp. 115–126, Vienna, Austria, 2013, Springer Verlag.
- [13] SULOVARI, A., KIRALIS, J., and MOORE, J. H., 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, pp. 127–138, Vienna, Austria, 2013, Springer Verlag.
- [14] SANTANDER-JIMENEZ, S. and VEGA-RODRIGUEZ, M. A., 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, pp. 139– 150, Vienna, Austria, 2013, Springer Verlag.
- [15] CASTALDI, D., MACCAGNOLA, D., MARI, D., and ARCHETTI, F., 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, pp. 151–162, Vienna, Austria, 2013, Springer Verlag.
- [16] MANNING, T. and WALSH, P., 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, pp. 163–174, Vienna, Austria, 2013, Springer Verlag.
- [17] GAUDESI, M., MARION, A., MUSNER, T., SQUILLERO, G., and TONDA, A., 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, pp. 175–185, Vienna, Austria, 2013, Springer Verlag.
- [18] ROSENTHAL, S., EL-SOURANI, N., and BORSCHBACH, M., 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, pp. 186–197, Vienna, Austria, 2013, Springer Verlag.
- [19] FISHER, J. M., ANDREWS, P., KIRALIS, J., SINNOTT-ARMSTRONG, N. A., and MOORE, J. H., 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, pp. 198–209, Vienna, Austria, 2013, Springer Verlag.
- [20] ZAGORSKI, M., 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, pp. 210–213, Vienna, Austria, 2013, Springer Verlag.