

Список литературы

- [Yeu et al.(2011)Yeu, Ahn, Yoon, and Park] Yunku Yeu, Jaegyo Ahn, Youngmi Yoon, and Sanghyun Park. Protein complex discovery from protein interaction network with high false-positive rate. In Clara Pizzuti, Marylyn D. Ritchie, and Mario Giacobini, editors, *9th European Conference Evolutionary Computation on Machine Learning and Data Mining in Bioinformatics: EvoBIO 2011*, volume 6623 of *LNCS*, pages 181–186, Turin, Italy, 27-29 April 2011. Springer Verlag.
- [Toti et al.(2011)Toti, Atzeni, and Polticelli] Daniele Toti, Paolo Atzeni, and Fabio Polticelli. An automatic identification and resolution system for protein-related abbreviations in scientific papers. In Clara Pizzuti, Marylyn D. Ritchie, and Mario Giacobini, editors, *9th European Conference Evolutionary Computation on Machine Learning and Data Mining in Bioinformatics: EvoBIO 2011*, volume 6623 of *LNCS*, pages 175–180, Turin, Italy, 27-29 April 2011. Springer Verlag.
- [Silva et al.(2011)Silva, Anunciao, and Lotz] Sara Silva, Orlando Anunciao, and Marco Lotz. A comparison of machine learning methods for the prediction of breast cancer. In Clara Pizzuti, Marylyn D. Ritchie, and Mario Giacobini, editors, *9th European Conference Evolutionary Computation on Machine Learning and Data Mining in Bioinformatics: EvoBIO 2011*, volume 6623 of *LNCS*, pages 163–174, Turin, Italy, 27-29 April 2011. Springer Verlag.
- [Toca et al.(2011)Toca, Márquez-Chamorro, Asencio-Cortés, and Aguilar-Ruiz] Cosme Ernesto Santiesteban Toca, Alfonso E. Márquez-Chamorro, Gualberto Asencio-Cortés, and Jesús S. Aguilar-Ruiz. A decision tree-based method for protein contact map prediction. In Clara Pizzuti, Marylyn D. Ritchie, and Mario Giacobini, editors, *9th European Conference Evolutionary Computation on Machine Learning and Data Mining in Bioinformatics: EvoBIO 2011*, volume 6623 of *LNCS*, pages 157–162, Turin, Italy, 27-29 April 2011. Springer Verlag.
- [Leonarski et al.(2011)Leonarski, Trovato, Tozzini, and Trylska] Filip Leonarski, Fabio Trovato, Valentina Tozzini, and Joanna Trylska. Genetic algorithm optimization of force field parameters. application to a coarse-grained model of RNA. In Clara Pizzuti, Marylyn D. Ritchie, and Mario Giacobini, editors, *9th European Conference Evolutionary Computation on Machine Learning and Data Mining in Bioinformatics: EvoBIO 2011*, volume 6623 of *LNCS*, pages 151–156, Turin, Italy, 27-29 April 2011. Springer Verlag.