BIOKDD02: Workshop on Data Mining in Bioinformatics July 23rd, 2002 Edmonton, Alberta, Canada

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Foreword

Bioinformatics provides opportunities for developing novel data mining methods. Some of the grand challenges in bioinformatics include protein structure prediction, homology search, multiple alignment and phylogeny construction, genomic sequence analysis, gene finding and gene mapping, as well as applications in gene expression data analysis, drug discovery in pharmaceutical industry, etc. This workshop aims to present latest results in this important area at the intersection of biology and KDD.

The proceedings contain 10 papers (out of 16 submissions) that were accepted for presentation at the workshop. Each paper was reviewed by two or three members of the program committee. Along with one invited speaker, we were able to assemble a very exciting program.

We would like to thank all the authors, invited speaker, and attendees for contributing to the success of the workshop. Special thanks are due to the program committee and external referees for help in reviewing the submissions.

Sincerely,

Mohammed Zaki, Jason Wang, Hannu Toivonen (editors)

Workshop Co-Chairs

Mohammed J. Zaki, Rensselaer Polytechnic Institute, USA Jason T. L. Wang, New Jersey Institute of Technology, USA Hannu T.T. Toivonen, University of Helsinki, Finland

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