Foreword

Special issue on computational molecular biology

This is the third JCSS special issue devoted to computational molecular biology, an emerging field with many exciting developments. This issue contains a collection of papers covering HMM and gene finding, phylogenetic studies, pooling design, alignment algorithms, optimal spaced seeds, and genome rearrangement algorithms. All papers were refereed according to the standards of JCSS.

We are grateful to the authors and especially to the referees for making this special issue possible.

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