

WABI Accepted Posters

1. *Panteleimon Zotos, Ioannis Xenarios, and Li Long.* **Applying Spectrum Graph Approach in MS/MS Peptide Libraries**
2. *Md. Shafiul Alam and Asish Mukhopadhyay.* **Restriction Site Mapping and The Point Placement Problem: Improved Upper and Lower Bounds.**
3. *Mark X. Caddick, Chris Sibthorp, Huihai Wu, Prudence W.H. Wong, and Neil Hall.* **High Throughput Sequencing and Motif Analysis of the *A. nidulans* Transcriptome**
4. *Georg Sauthoff, Stefan Janssen, Robert Giegerich.* **No Subscripts, No Errors -- Enjoy Dynamic Programming in Bellman's GAP**
5. *Yoo-Ah Kim, Stefan Wuchty, Teresa Przytycka.* **Simultaneous Identification of Causal Genes and Dys-regulated Pathways in Complex Diseases**
6. *Andrew Bromage, Thomas Conway, Bryan Beresford-Smith, Justin Zobel.* **Space-Efficient Fragment Assembly using Succinct Data Structures**
7. *Ugochukwa C. Anokwuru, Stanley Dappa, Sikhongi S. Phungwayo, Stefan Spears, Rahul Suri, M. Shel Swenson, C. Randal Linder, and Tandy Warnow.* **Analyzing a 50,174-taxon Dataset using Maximum Likelihood**
8. *Ghada Badr, Krister M. Swenson, and David Sankoff.* **Listing All Sorting Sequences by Reversals Problem Revisited with a New Framework**
9. *Mona Singh.* **Predicting and Analyzing Protein Interactions**
10. *Bonnie Kirkpatrick, Hilary Finucane, Yakir Reshef, and Richard Karp.* **Algorithms for Comparing Pedigree Graphs**
11. *Radoslaw Suchacki, Pierre Chardaire, Katharina T. Huber.* **A GRASP-based Approach for the Pure Parsimony Haplotype Inference Problem**
12. *F. Schwach, S. Moxon, T. Dalmay, V. Moulton.* **Computational Tools for the Analysis of Large-scale Plant Small RNA Data Sets**

Presenting authors