

References

- [1] Daniel Fasulo, Anne-Katrin Emde, Lu-Yong Wang, Karin Noy, and Nathan Edwards. Alignment of mass spectrometry data by clique finding and optimization. In Trey Ideker and Vineet Bafna, editors, *Proceedings of the Joint RECOMB 2006 Satellite Workshops on Systems Biology and on Computational Proteomics (San Diego, CA, USA, December 1-3, 2006)*, volume 4532 of *LNCS/LNBI*, pages 119–129. Springer-Verlag, Berlin-Heidelberg, 2007.
- [2] Fereydoun Hormozdiari, Petra Berenbrink, Nataša Pržulj, and Cenk Sahinalp. Not all scale free networks are born equal: The role of the seed graph in ppi network emulation. In Trey Ideker and Vineet Bafna, editors, *Proceedings of the Joint RECOMB 2006 Satellite Workshops on Systems Biology and on Computational Proteomics (San Diego, CA, USA, December 1-3, 2006)*, volume 4532 of *LNCS/LNBI*, pages 1–13. Springer-Verlag, Berlin-Heidelberg, 2007.
- [3] Hailiang Huang, Lan V. Zhang, Frederick P. Roth, and Joel S. Bader. Probabilistic paths for protein complex inference. In Trey Ideker and Vineet Bafna, editors, *Proceedings of the Joint RECOMB 2006 Satellite Workshops on Systems Biology and on Computational Proteomics (San Diego, CA, USA, December 1-3, 2006)*, volume 4532 of *LNCS/LNBI*, pages 14–28. Springer-Verlag, Berlin-Heidelberg, 2007.
- [4] Hans-Michael Kaltenbach, Sebastian Böcker, and Sven Rahmann. Markov additive chains and applications to fragment statistics for peptide mass fingerprinting. In Trey Ideker and Vineet Bafna, editors, *Proceedings of the Joint RECOMB 2006 Satellite Workshops on Systems Biology and on Computational Proteomics (San Diego, CA, USA, December 1-3, 2006)*, volume 4532 of *LNCS/LNBI*, pages 29–41. Springer-Verlag, Berlin-Heidelberg, 2007.
- [5] Celine Lefebvre, Wei Keat Lim, Katia Basso, Riccardo Dalla Favera, and Andrea Califano. A context-specific network of protein-dna and protein-protein interactions reveals new regulatory motifs in human b cells. In Trey Ideker and Vineet Bafna, editors, *Proceedings of the Joint RECOMB 2006 Satellite Workshops on Systems Biology and on Computational Pro-*

teomics (San Diego, CA, USA, December 1-3, 2006), volume 4532 of *LNCS/LNBI*, pages 42–56. Springer-Verlag, Berlin-Heidelberg, 2007.

- [6] Jianhua Ruan and Weixiong Zhang. Identification and evaluation of functional modules in gene co-expression networks. In Trey Ideker and Vineet Bafna, editors, *Proceedings of the Joint RECOMB 2006 Satellite Workshops on Systems Biology and on Computational Proteomics (San Diego, CA, USA, December 1-3, 2006)*, volume 4532 of *LNCS/LNBI*, pages 57–76. Springer-Verlag, Berlin-Heidelberg, 2007.
- [7] Derek Ruths, Jen-Te Tseng, Luay Nakhleh, and Prahlad T. Ram. De novo signaling pathway predictions based on protein-protein interaction, targeted therapy and protein microarray analysis. In Trey Ideker and Vineet Bafna, editors, *Proceedings of the Joint RECOMB 2006 Satellite Workshops on Systems Biology and on Computational Proteomics (San Diego, CA, USA, December 1-3, 2006)*, volume 4532 of *LNCS/LNBI*, pages 108–118. Springer-Verlag, Berlin-Heidelberg, 2007.
- [8] Mingzhou (Joe) Song and Z. Lewis Liu. A linear discrete dynamic system model for temporal gene interaction and regulatory network influence in response to bioethanol conversion inhibitor hmf for ethanologenic yeast. In Trey Ideker and Vineet Bafna, editors, *Proceedings of the Joint RECOMB 2006 Satellite Workshops on Systems Biology and on Computational Proteomics (San Diego, CA, USA, December 1-3, 2006)*, volume 4532 of *LNCS/LNBI*, pages 77–95. Springer-Verlag, Berlin-Heidelberg, 2007.
- [9] Yin Wu, Yehia Mechref, Iveta Klouckova, Milos V. Novotny, and Haixu Tang. A computational approach for the identification of site-specific protein glycosylations through ion-trap mass spectrometry. In Trey Ideker and Vineet Bafna, editors, *Proceedings of the Joint RECOMB 2006 Satellite Workshops on Systems Biology and on Computational Proteomics (San Diego, CA, USA, December 1-3, 2006)*, volume 4532 of *LNCS/LNBI*, pages 96–107. Springer-Verlag, Berlin-Heidelberg, 2007.