References

[1] Sèverine Bérard, Anne Bergeron, and Cedric Chauve. Conservation of combinatorial structures in evolution scenarios. In Jens Lagergren, editor, *Proceedings of the RECOMB 2004 International Workshop on Comparative Genomics, RCG'2004 (Bertinoro, Italy, October 16-19, 2004)*, volume 3388 of *LNCS/LNBI*, pages 1–14. Springer-Verlag, Berlin-Heidelberg, 2005.

[2] Jeremy Buhler and Rachel Nordgren. Toward a phylogenetically aware algorithm for fast dna similarity search. In Jens Lagergren, editor, *Proceedings of the RECOMB 2004 International Workshop on Comparative Genomics, RCG'2004 (Bertinoro, Italy, October 16-19, 2004)*, volume 3388 of *LNCS/LNBI*, pages 15–29. Springer-Verlag, Berlin-Heidelberg, 2005.

[3] Jeong-Hyeon Choi, Kwangmin Choi, Hwan-Gue Cho, and Sun Kim. Multiple genome alignment by clustering pairwise matches. In Jens Lagergren, editor, *Proceedings of the RECOMB 2004 International Workshop on Comparative Genomics, RCG'2004 (Bertinoro, Italy, October 16-19, 2004)*, volume 3388 of *LNCS/LNBI*, pages 30–41. Springer-Verlag, Berlin-Heidelberg, 2005.

[4] Paweł Górecki and Jerzy Tiuryn. On the structure of reconciliations. In Jens Lagergren, editor, *Proceedings of the RECOMB 2004 International Workshop on Comparative Genomics, RCG'2004 (Bertinoro, Italy, October 16-19, 2004)*, volume 3388 of *LNCS/LNBI*, pages 42–54. Springer-Verlag, Berlin-Heidelberg, 2005.

[5] Rose Hoberman, David Sankoff, and Dannie Durand. The statistical significance of max-gap clusters. In Jens Lagergren, editor, *Proceedings of the RECOMB 2004 International Workshop on Comparative Genomics, RCG'2004 (Bertinoro, Italy, October 16-19, 2004)*, volume 3388 of *LNCS/LNBI*, pages 55–71. Springer-Verlag, Berlin-Heidelberg, 2005.

[6] Bob Mau, Aaron E. Darling, and Nicole T. Perna. Identifying evolutionarily conserved segments among multiple divergent and rearranged genomes. In Jens Lagergren, editor, *Proceedings of the RECOMB 2004*

International Workshop on Comparative Genomics, RCG'2004 (Bertinoro, Italy, October 16-19, 2004), volume 3388 of *LNCS/LNBI*, pages 72–84. Springer-Verlag, Berlin-Heidelberg, 2005.

[7] István Miklós and Jotun Hein. Genome rearrangement in mitochondria and its computational biology. In Jens Lagergren, editor, *Proceedings of the RECOMB 2004 International Workshop on Comparative Genomics, RCG'2004 (Bertinoro, Italy, October 16-19, 2004)*, volume 3388 of *LNCS/LNBI*, pages 85–96. Springer-Verlag, Berlin-Heidelberg, 2005.

[8] David Sankoff, Jean-François Lefebvre, Elisabeth Tillier, Adrian Maler, and Nadia El-Mabrouk. The distribution of inversion lengths in bacteria. In Jens Lagergren, editor, *Proceedings of the RECOMB 2004 International Workshop on Comparative Genomics, RCG'2004 (Bertinoro, Italy, October 16-19, 2004)*, volume 3388 of *LNCS/LNBI*, pages 97–108. Springer-Verlag, Berlin-Heidelberg, 2005.

[9] David Sankoff and Matthew Mazowita. Estimators of translocations and inversions in comparative maps. In Jens Lagergren, editor, *Proceedings of the RECOMB 2004 International Workshop on Comparative Genomics, RCG'2004 (Bertinoro, Italy, October 16-19, 2004)*, volume 3388 of *LNCS/LNBI*, pages 109–122. Springer-Verlag, Berlin-Heidelberg, 2005.

[10] Bahar Taneri, Alexey Novoradovsky, Ben Snyder, and Terry Gaasterland. Databases for comparative analysis of human-mouse orthologous alternative splicing. In Jens Lagergren, editor, *Proceedings of the RECOMB 2004 International Workshop on Comparative Genomics, RCG'2004 (Bertinoro, Italy, October 16-19, 2004)*, volume 3388 of *LNCS/LNBI*, pages 123–131. Springer-Verlag, Berlin-Heidelberg, 2005.