Figure S6: Variation in recombination rates in relation to GC content for cross 1A5x1E4. Recombination rates were estimated in non-overlapping 20 kb sequence segments. The % GC content of each of the 20 kb sequence segments was used to correlate recombination rates and GC content. Panels distinguish correlations for core and accessory chromosomes. The highest recombination rates were observed in high GC-content segments. Spearman’s rank correlation rho = 0.324 and p-value < 2.2e-16.