Figure S3: Accessory chromosomes showed large non-recombining segments (in red) in both cross 3D1x3D7 and 1A5x1E4. Recombination fractions were calculated among all marker pairs per chromosome. Interpolation was used to report recombination fractions in a regular grid spanning the chromosome. Large red blocks indicate regions of low recombination rates. Green or blue blocks near the diagonal indicate the presence of a recombination hotspot (see Figure 1). White areas in the upper triangles show regions of the chromosomes that were not sufficiently covered by SNP markers to interpolate recombination fractions.