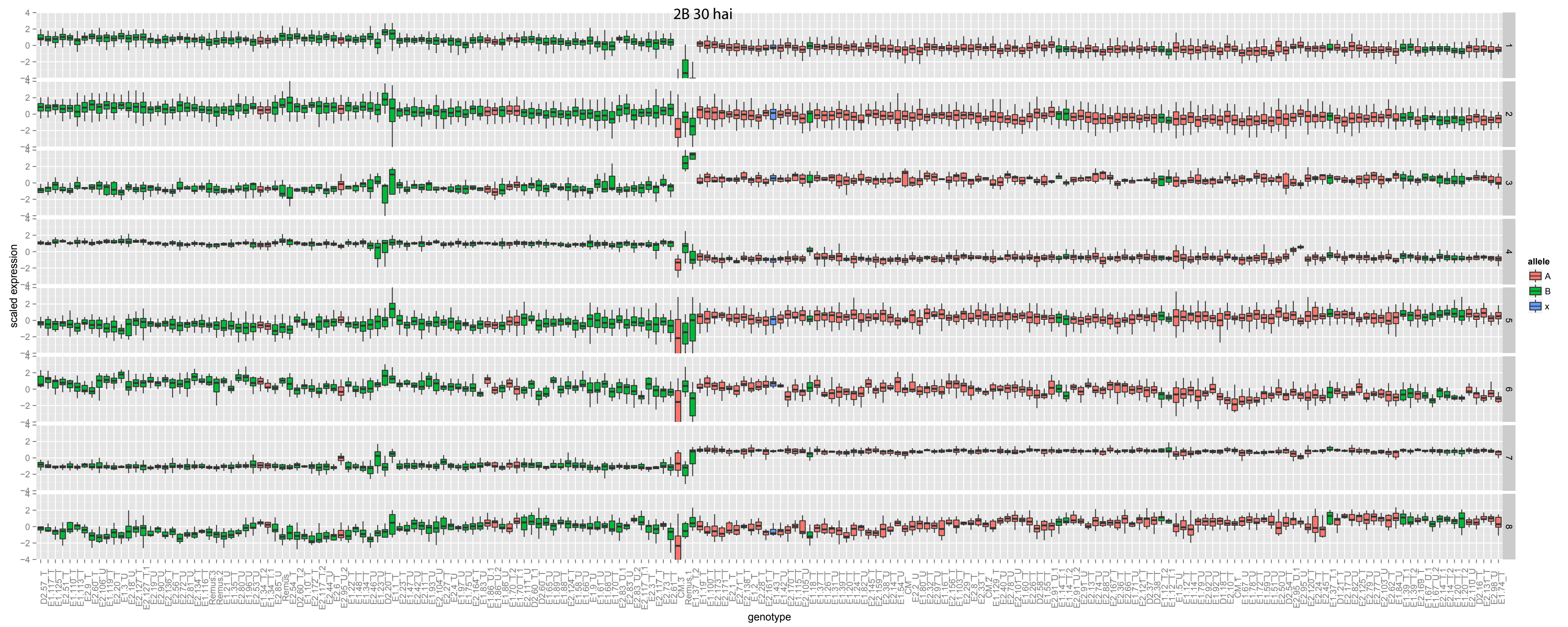


Figure S4 Hierarchical clustering of eQTL expression profiles for hotspots 2B at 30 hai (page 2), 2B at 50 hai (page 3), 4A at 30 hai and 50 hai (pages 4 and 5) and 5A at 30 hai and 50 hai (pages 6 and 7). Parental genotypes (A = CM-82036, B = Remus) at the locus are represented in color for each line. The plotted signal intensities have been normalized and transformed prior clustering.

2B 30 hai



hotspot 2B 50 hai

