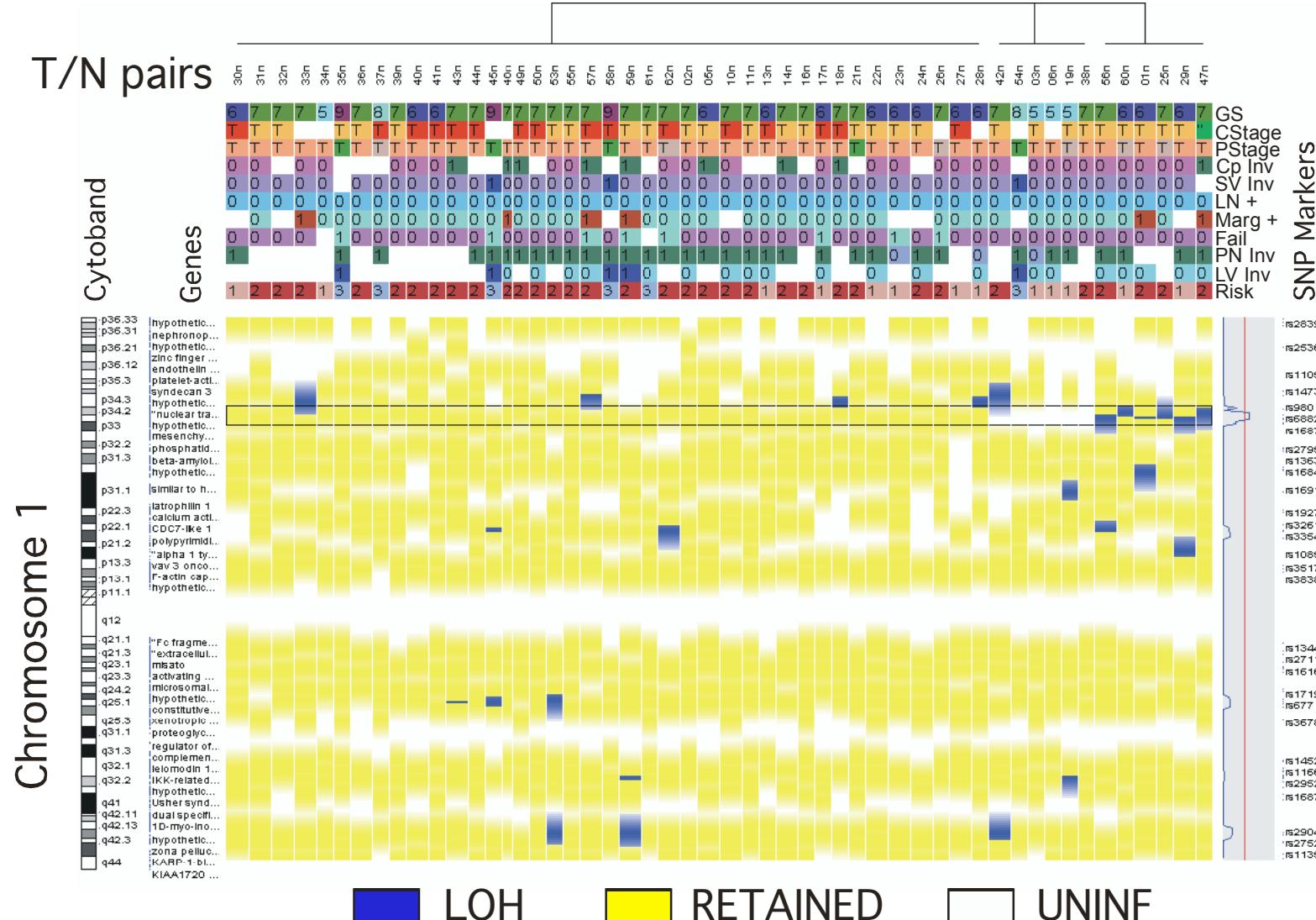
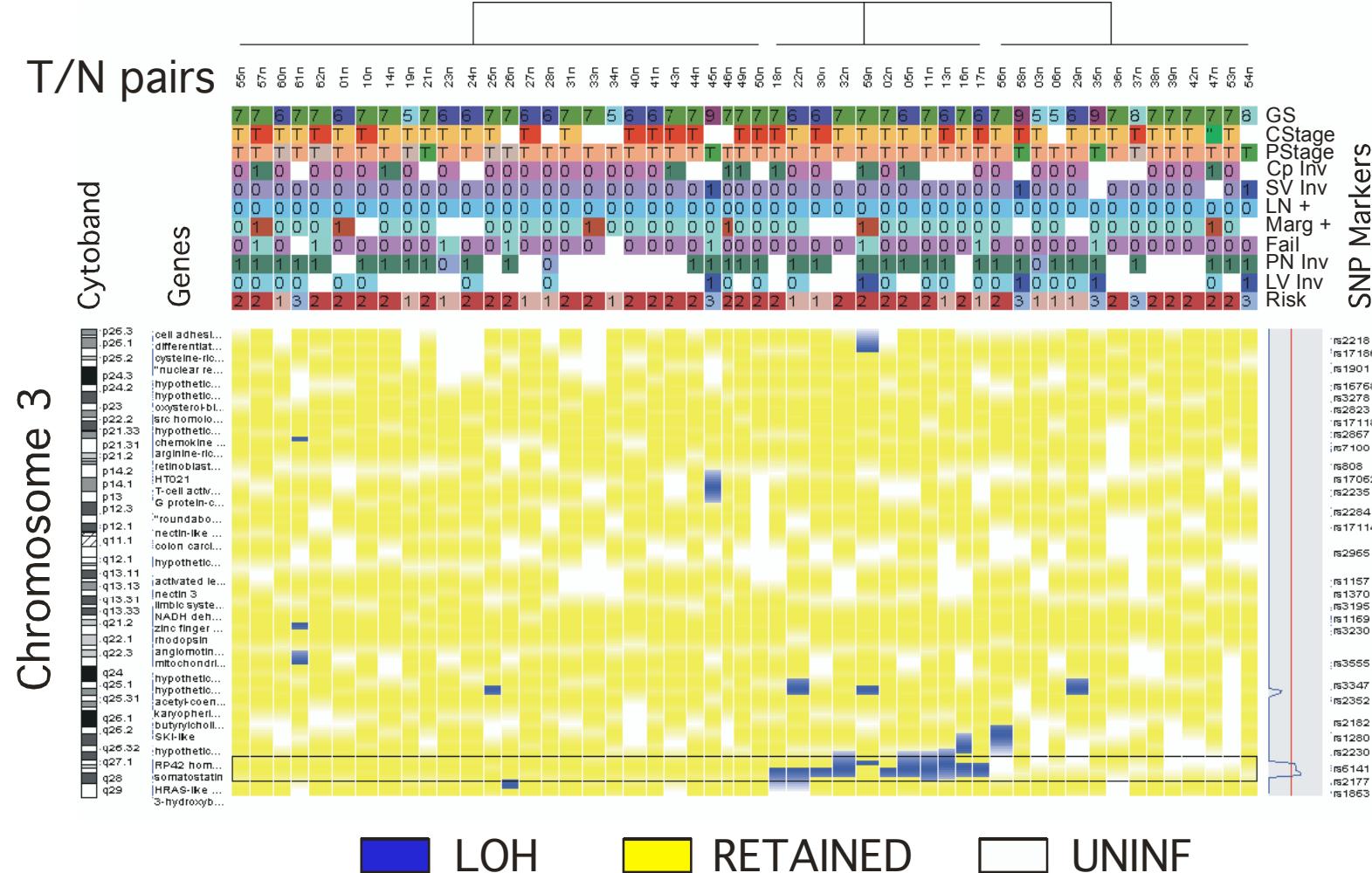


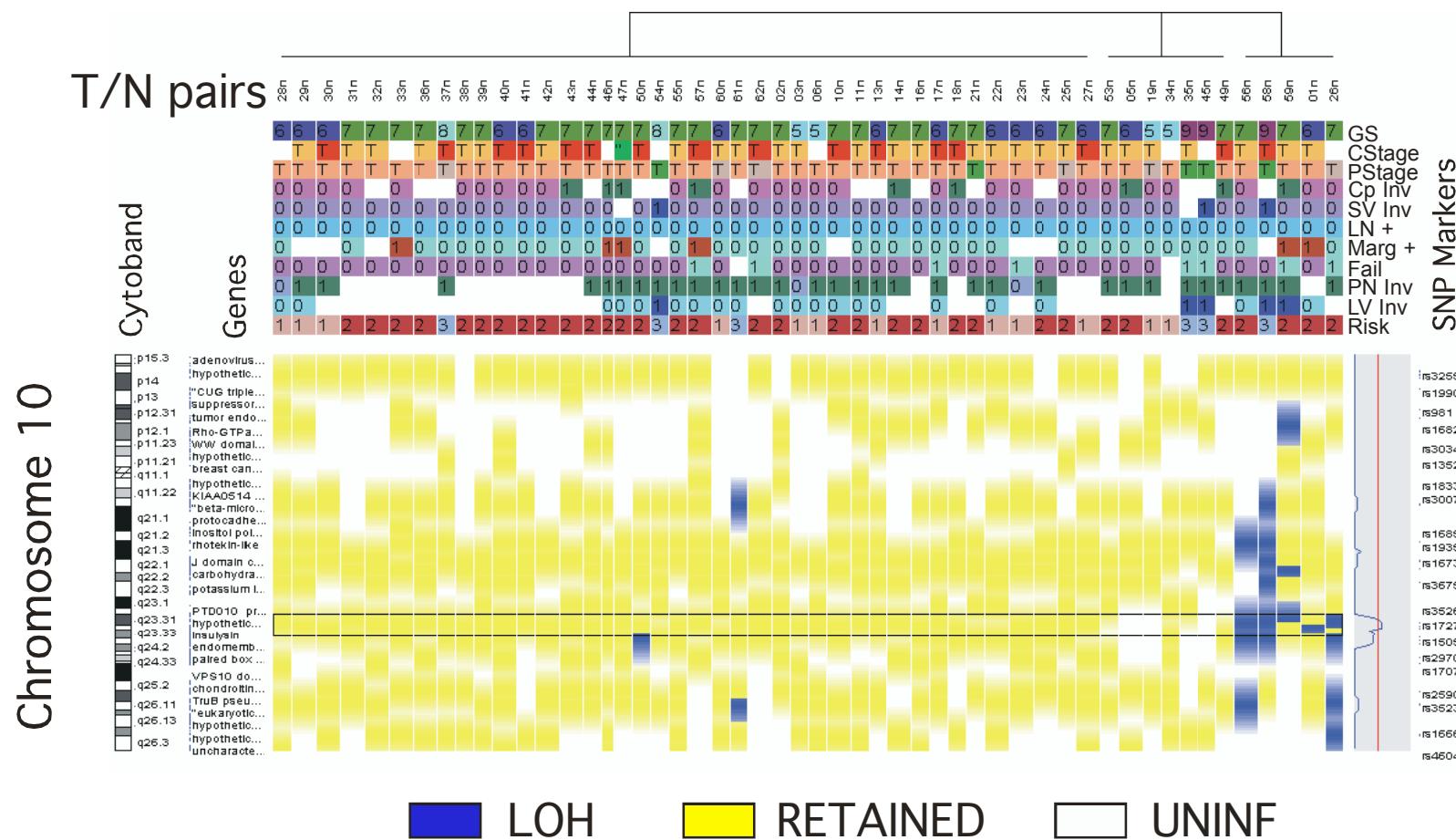
Supplemental Figure 1A



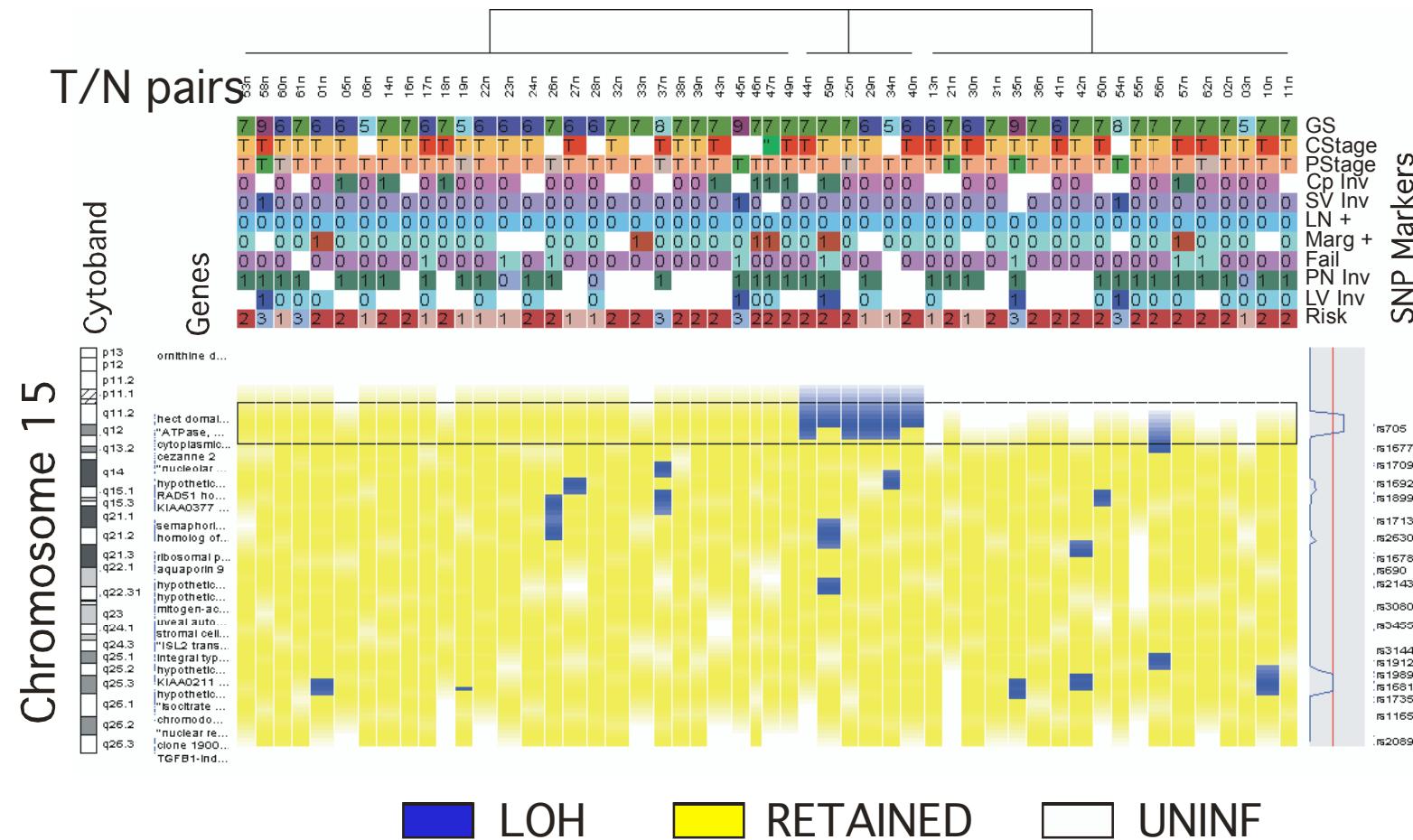
Supplemental Figure 1B



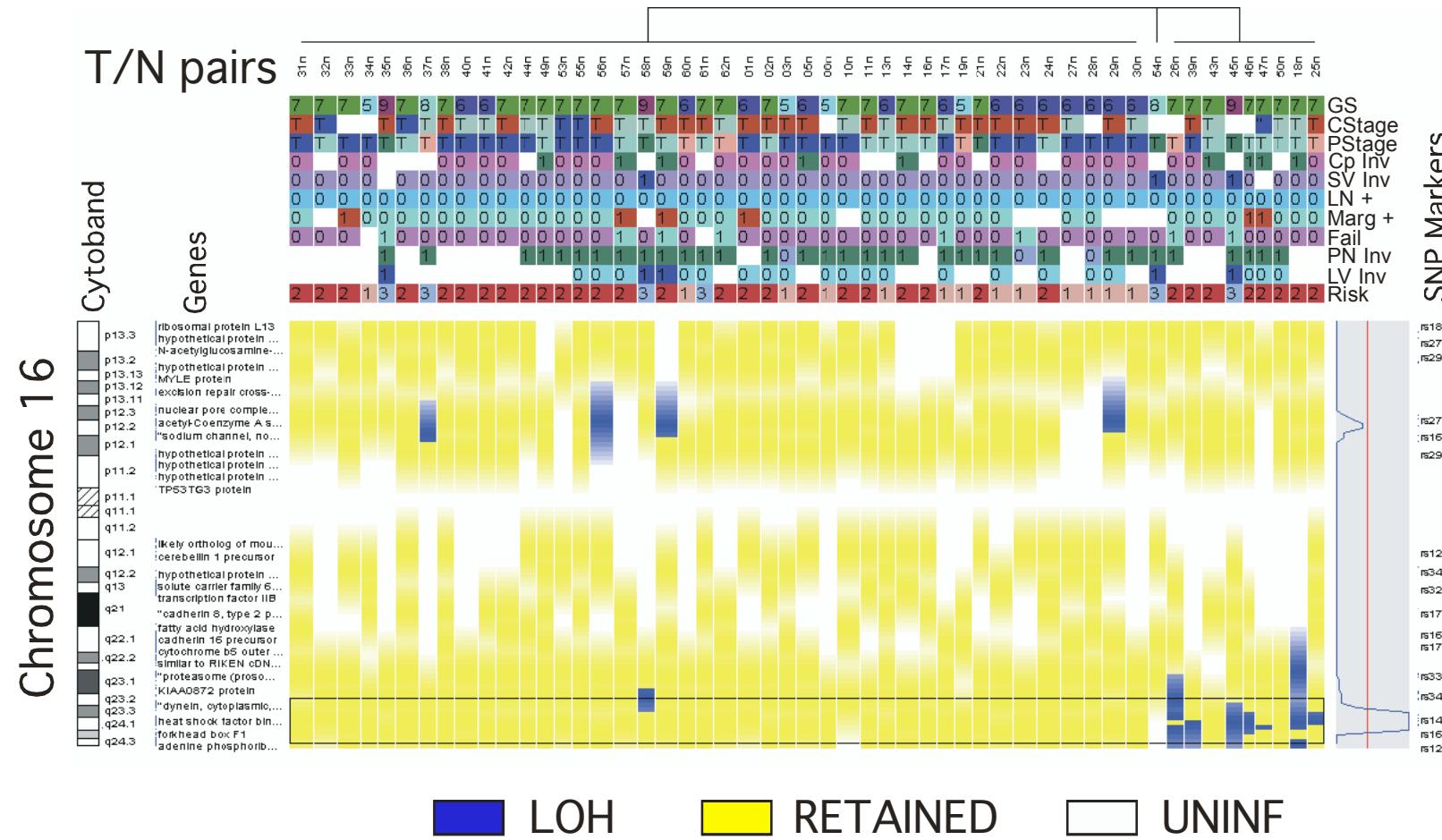
Supplemental Figure 1C



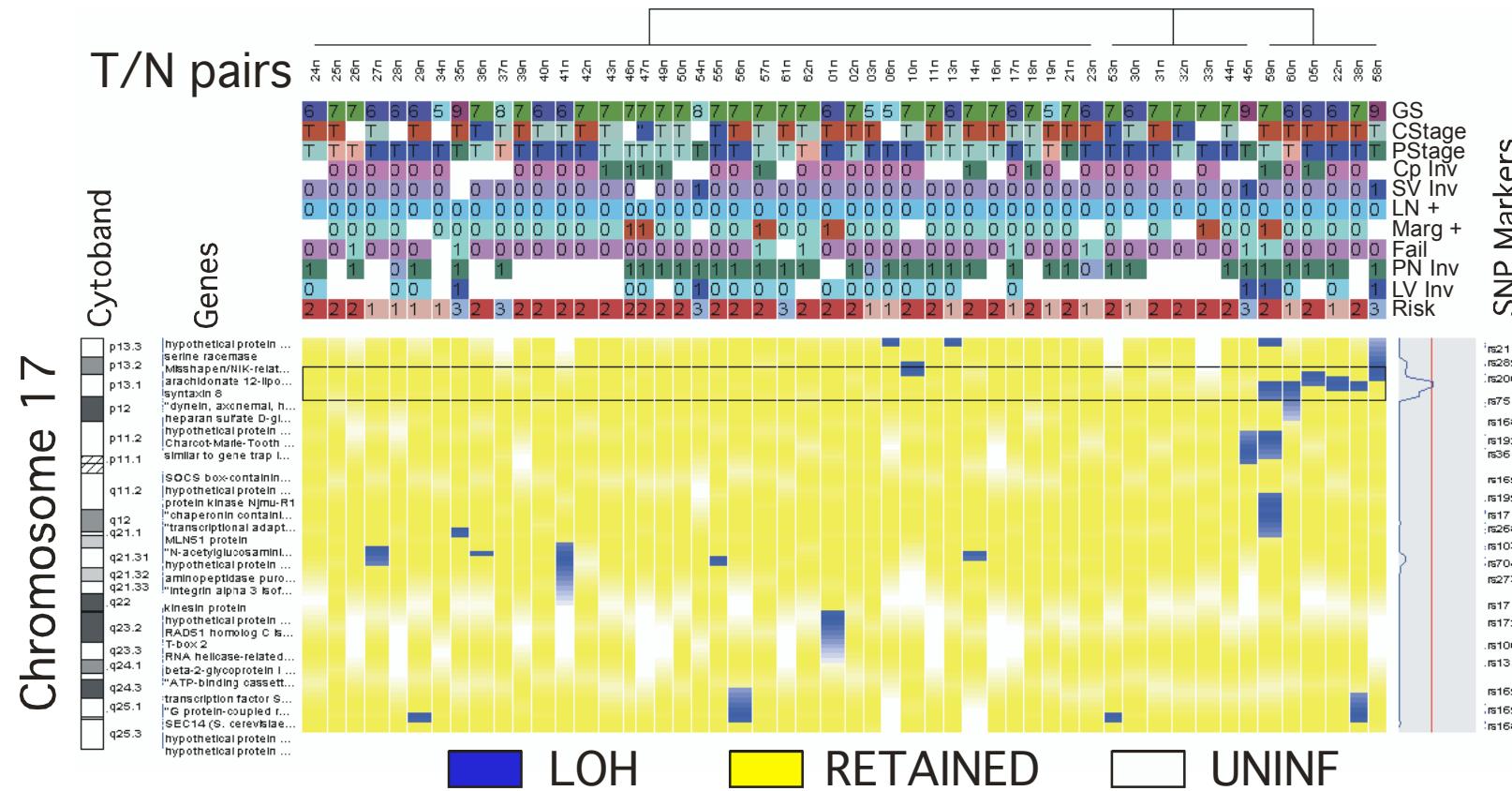
Supplemental Figure 1D



Supplemental Figure 1E



Supplemental Figure 1F



Supplemental Figure Legends

Supplemental Figure 1: Chromosome maps for the regions of LOH passing permutation.

For each region of loss, the chromosome view in dChipSNP is shown. Cancer-normal pairs are shown across the top along with associated clinical data while chromosomal regions and the ideogram map are shown on the left. On the right of each map is the plot of the summary LOH score. Hierarchical clustering based only on the data for each chromosomal region is shown. Blue indicates a region of LOH, yellow indicates Retention and white indicates the absence of informative alleles. A) Chromosome 1, B) Chromosome 3, C) Chromosome 10, D) Chromosome 15, E) Chromosome 16, F) Chromosome 17.