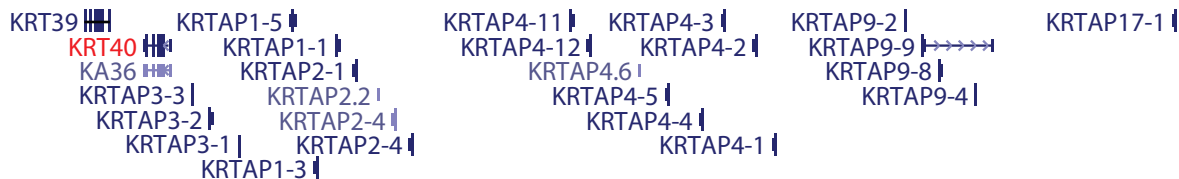
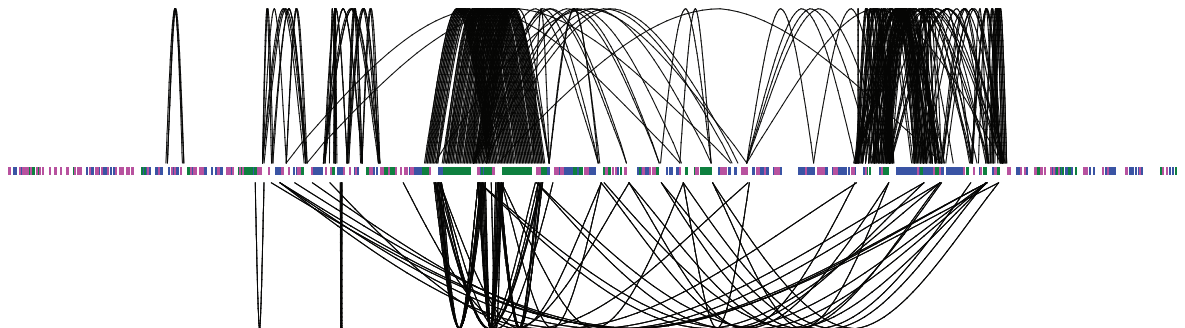


1. Genes

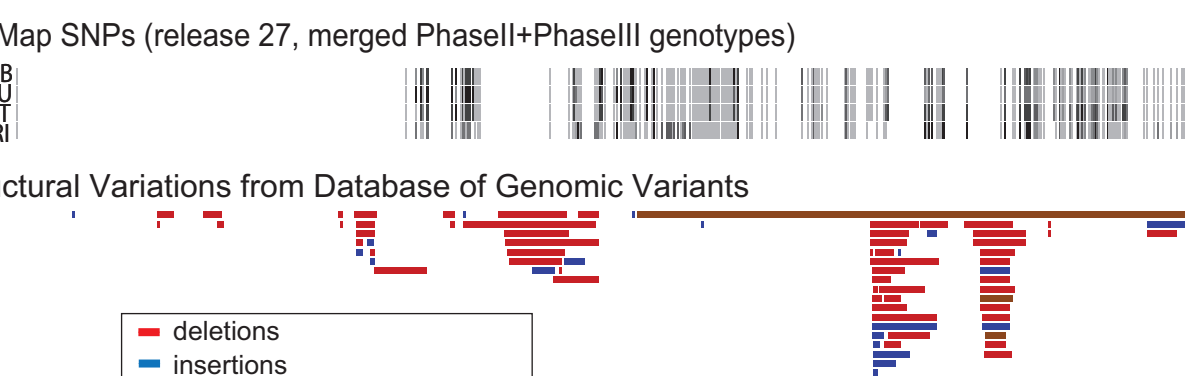


2. Duplications (Sequence homology > 90 %, size > 100bp)

(+) strand to (+) strand



(+) strand to (-) strand



3. HapMap SNPs (release 27, merged PhaseII+PhaseIII genotypes)



4. Structural Variations from Database of Genomic Variants

