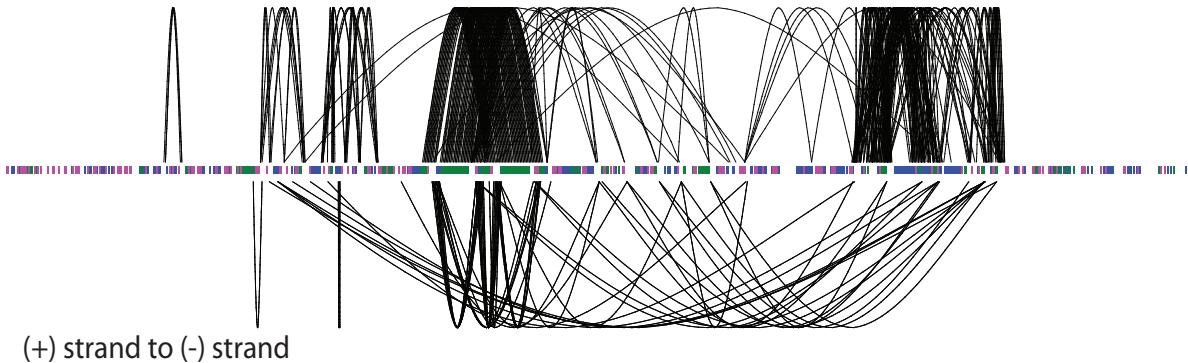


1. Genes



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

(+) strand to (+) strand



(+) strand to (-) strand

3. HapMap SNPs (release 27, merged Phasell+PhaselII genotypes)

SNPs CHB
SNPs CEU
SNPs JPT
SNPs YRI



4. Structural Variations from Database of Genomic Variants

- deletions
- insertions
- both deletions and insertions

