UCSC Genes Based on RefSeq, UniProt, GenBank, CCDS and Comparative Genomics

Duplications of >1000 bases of non-RepeatMasked sequence

Human chained self alignments

Mapability - CRG GEM Alignability of 75mers with no more than 2 mismatches

19 individually genotyped SNPS

Common SNPs(132)

Simple Nucleotide Polymorphisms (dbSNP 132) Found in >= 1% of samples