



Scale
chr5:

117,355,000

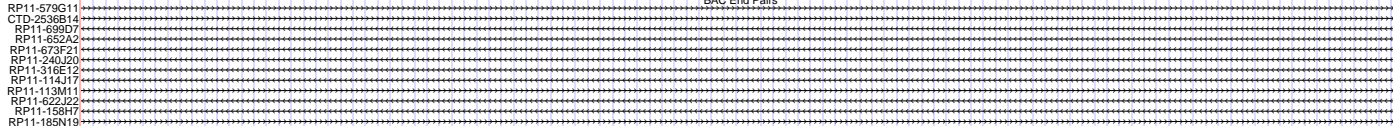
5 kb

117,360,000

hg19

117,365,000

BAC End Pairs



UCSC Genes (RefSeq, GenBank, CCDS, Rfam, tRNAs & Comparative Genomics)
RefSeq Genes



Basic Gene Annotation Set from GENCODE Version 24lift37 (Ensembl 83)

Pseudogene Annotation Set from GENCODE Version 24lift37 (Ensembl 83)

Pseudogene Annotation Set from GENCODE Version 19

AceView Gene Models With Alt-Splicing

Ensembl Gene Predictions - archive 75 - feb2014



Database of Genomic Variants: Structural Variation (CNV, Inversion, In/del)



