Window Position
Scale
chrX:

Human Mar. 2006 (NCBI36/hg18)
100 kb
hg18
151,150,000 151,200,000 151,250,000 151,300,000 151,350,000

UCSC Genes (RefSeq, GenBank, tRNAs & Comparative Genomics)
GABRA3
CS266382

Human Genome Diversity Project SNP Population Allele Frequencies
rs4828684 rs5925128 rs5970231 rs1602624 rs1492296 rs11798711 rs5970292 rs6627588
rs11094547 rs1907600 rs5970232 rs1492302 rs1565610 rs389292 rs6526107
rs5970220 rs5925140 rs5970242 rs5925155 rs4828589 rs1109840
rs6653441 rs994423 rs2131190 rs5970266 rs5969896
rs6526109
rs6526110
rs7065528

RefSeq genes from NCBI
UCSC RefSeq

Human mRNAs from GenBank
Human mRNAs

Human ESTs That Have Been Spliced
Spliced ESTs

Simple Nucleotide Polymorphisms (dbSNP, build 130)
rs58074393 rs4828689 rs5970228 rs73639075 rs4431735 rs970417 rs3830670 rs4828694 rs5970299 rs5925169
rs61444466 rs7049421 rs1491788 rs2162140 rs6653475 rs7050378 rs584473 rs4828588 rs5970300 rs6627591
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ENCODE Gencode Manual Gene Annotations (level 1+2) (Oct 2009)
AC116666.1
GABRA3
GABRA3
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GABRA3

ENCODE Gencode Automated Gene Annotations (level 3) (Oct 2009)
hsa-mir-105-1
hsa-mir-767
hsa-mir-105-2

Repeating Elements by RepeatMasker
RepeatMasker

ENCODE Digital DNaseI Hypersensitivity Clusters
DNase Clusters

Multiz Alignments of 44 Vertebrates
Rhesus
Mouse
Dog
Elephant
Opossum
Platypus
Chicken
Lizard
X_tropicalis
Stickleback

ln(x+1) 4 _
ENCODE Transcription Levels Assayed by RNA-seq on 6 Cell Lines
Transcription
0