Scale
chr28:
100 kb
canFam3
10,250,000
10,300,000
10,350,000
10,400,000
10,450,000
10,500,000
10,550,000
10,600,000
A
11variants_LR
BedGraph format
B
25variants_LR
BedGraph format
C
11variants_GSD
BedGraph format
D
25variants_GSD
BedGraph format
E
top SP SNPs
User Supplied Track
RefSeq Genes
Non-Dog RefSeq Genes
Arabido TMN1
Homo LCOR
Homo LCOR
Homo LCOR
Pongo C10H10orf12
Rattus Eif4e
Oryctol EIF4E
Taeniop LOC100189975
Pan EIF4E
Danio eif4e1c
Ictalur if4e
Danio eif4eb
Danio eif4e1b
Salmo if4e
Danio eif4ea
Strongy eif4e1
Aplysia LOC100533326
Acyrtho LOC100160676
Acyrtho LOC100169333
Drosoph eIF4E-6
Drosoph eIF4E-6
Drosoph eIF4E-6
Bombyx LOC732913
Caenorh ife-3
Caenorh ife-3
Acyrtho LOC100144893
Glycine LOC100499680
Ashbya AGOS_AFR478W
Xenopus lcor
Danio lcor
Mus Lcor
Homo C10orf12
Gallus SLIT1
Homo SLIT1
Danio slit1b
Xenopus slit1
Rattus Slit1
Xenopus slit1
Mus Slit1
Danio slit2
Danio slit1a
Danio slit3
Homo ARHGAP19-SLIT1
Homo ARHGAP19
Homo ARHGAP19
Homo ARHGAP19
Mus Arhgap19
Mus Arhgap19
Macaca ARHGAP19
Gallus ARHGAP19
Xenopus arhgap19