

CON_314_BR1

%features

50
40
30
20
10
0

■ % in genome
▨ detected
■ % in sample

8.3
6.6
5
3.3
1.7
0

protein_coding
pseudogene
lincRNA
antisense
miRNA
misc_RNA
sense_intronic
snoRNA
processed_transcript
snRNA
sense_overlapping
rRNA
IG_V_gene
IG_V_pseudogene
TR_V_gene
polymorphic_pseudogene
3prime_overlapping_ncrna
IG_C_gene
TR_V_pseudogene
TR_C_gene
IG_C_pseudogene
IG_J_gene
TR_J_gene
IG_J_pseudogene
IG_D_gene

