

SK3R11_Tdil22_10

%features

60
40
20
0

■ % in genome
▨ detected
■ % in sample

2.1
1.4
0.7
0

protein_coding
processed_pseudogene
lincRNA
TEC
antisense
miRNA
snoRNA
processed_transcript
snRNA
unprocessed_pseudogene
processed_pseudogene
sense_intronic
misc_RNA
unprocessed_pseudogene
pseudogene
functional_promoter_lincRNA
scaRNA
rRNA
polymorphic_pseudogene
IG_V_gene
TR_J_gene
sense_overlapping
sense_unitary_pseudogene
unitary_pseudogene
IG_V_pseudogene
TR_C_gene
ribozyme
TR_V_gene
prime_overlapping_ncRNA
IG_D_gene
TR_J_pseudogene

