

DK3R3r_C3

%features

60
40
20
0

■ % in genome
▨ detected
■ % in sample

1.7
1.1
0.6
0

protein_coding
processed_pseudogene
lincRNA
antisense
TEC
miRNA
snoRNA
processed_transcript
snRNA
unprocessed_pseudogene
processed_pseudogene
misc_RNA
sense_intronic
unprocessed_pseudogene
rRNA
pseudogene
regional_promoter_lncRNA
polymorphic_pseudogene
scaRNA
IG_V_gene
IG_C_gene
IG_V_pseudogene
sense_overlapping
TR_V_gene
TR_C_gene
unitary_pseudogene
ribozyme
ribosomal_pseudogene
prime_overlapping_ncRNA
sRNA
TR_V_pseudogene