

DK3R1r_C1

%features

60
40
20
0

■ % in genome
▨ detected
■ % in sample

1.7
1.2
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
pseudogene
rRNA
functional_promoter_lincRNA
scaRNA
polymorphic_pseudogene
IG_C_gene
IG_V_gene
sense_overlapping
TR_V_gene
unitary_pseudogene
IG_V_pseudogene
ribozyme
ribbed_unitary_pseudogene
TR_C_gene
prime_overlapping_ncRNA
IG_D_gene
macro_lincRNA
sRNA
TR_J_gene

