

DK3R9r-C3L4

%features

60
40
20
0

■ % in genome
▨ detected
■ % in sample

2.4
1.6
0.8
0

protein_coding
processed_pseudogene
lincRNA
antisense
TEC
miRNA
snoRNA
processed_transcript
snRNA
unprocessed_pseudogene
misc_RNA
processed_pseudogene
sense_intronic
unprocessed_pseudogene
rRNA
transcriptional_promoter_lincRNA
pseudogene
polymorphic_pseudogene
scaRNA
IG_V_gene
IG_C_gene
IG_V_pseudogene
ribozyme
unitary_pseudogene
sense_overlapping
TR_C_gene
TR_V_gene
TR_V_pseudogene
ribosomal_pseudogene
IG_D_gene
IG_J_gene
macro_lincRNA
sRNA
TR_J_gene