

DK3R11_K2L4

%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
unprocessed_pseudogene
misc_RNA
sense_intronic
unprocessed_pseudogene
pseudogene
rRNA
structural_promoter_lncRNA
scaRNA
polymorphic_pseudogene
IG_C_gene
IG_V_gene
IG_V_pseudogene
sense_overlapping
ribozyme
unitary_pseudogene
tribed_unitary_pseudogene
TR_C_gene
TR_V_gene
TR_V_pseudogene
IG_D_gene
IG_J_gene
macro_lncRNA
sRNA
TR_J_gene