

DK3R8_C2L4

%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

regional_promoter_lncRNA

scaRNA

polymorphic_pseudogene

IG_C_gene

IG_V_gene

IG_V_pseudogene

sense_overlapping

ribozyme

unitary_pseudogene

ribbed_unitary_pseudogene

TR_C_gene

TR_V_gene

TR_V_pseudogene

IG_D_gene

IG_J_gene

macro_lncRNA

sRNA

TR_J_gene