global
sense_intronic
lincRNA
processed_pseudogene
protein_coding
TEC
antisense
miRNA
processed_transcript
snoRNA
misc_RNA
snRNA
unprocessed_pseudogene
transcribed_unprocessed_pseudogene
sense_overlapping
transcribed_processed_pseudogene
scaRNA
rRNA
pseudogene
unitary_pseudogene
bidirectional_promoter_lncRNA
ribozyme
polymorphic_pseudogene
IG_C_gene
IG_D_pseudogene
IG_V_pseudogene
IG_V_gene
TR_C_gene
TR_V_gene
TR_V_pseudogene
TR_J_gene
sRNA
macro_lncRNA
transcribed_unitary_pseudogene
Detected features
Depth in millions of reads
YK3R1−5291−4aplus
YK3R2−5291−4aplus
YK3R3−5291−4aplus
YK3R4−5291−4aminus
YK3R5−5291−4aminus
YK3R6−5269−4fplus
YK3R7−5269−4fplus
YK3R8−5269−4fplus
YK3R9−5269−4fminus
YK3R10−5269−4fminus
YK3R11−Cplus−1plus