

MF3R2_Prox1-2

%features

60
50
40
30
20
10
0

■ % in genome
▨ detected
■ % in sample

3.2
2.7
2.2
1.6
1.1
0.5
0

protein_coding
processed_pseudogene
lincRNA
TEC
antisense
miRNA
snoRNA
processed_transcript
snRNA
unprocessed_pseudogene
sense_intronic
misc_RNA
processed_pseudogene
rRNA
unprocessed_pseudogene
pseudogene
functional_promoter_lincRNA
scaRNA
polymorphic_pseudogene
IG_V_gene
sense_overlapping
TR_J_gene
unitary_pseudogene
ribozyme
IG_V_pseudogene
tribed_unitary_pseudogene
IG_C_gene
sRNA
macro_lincRNA
scRNA
TR_C_gene
TR_D_gene
TR_J_pseudogene