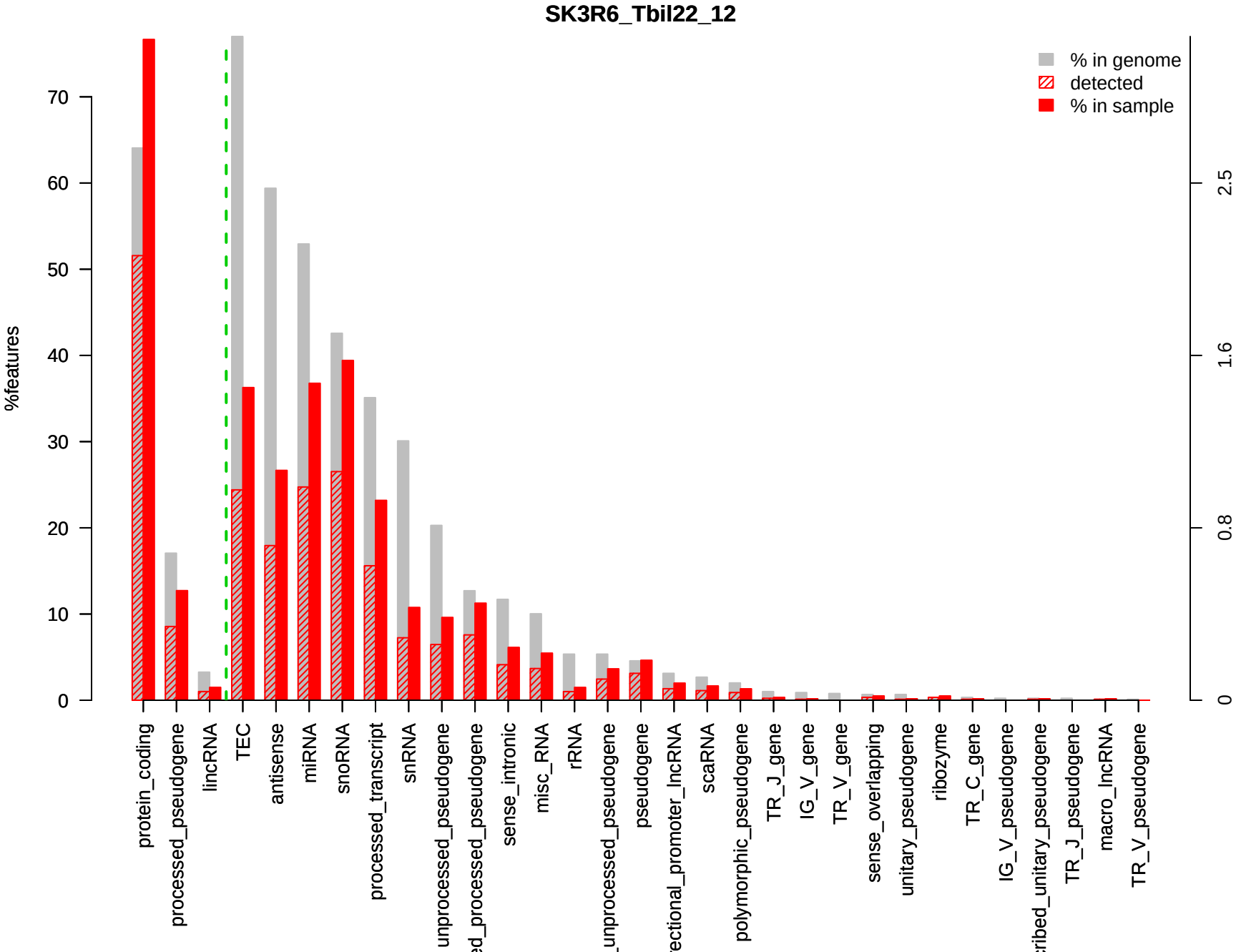
SK3R6_Tbil22_12
% in genome
detected
% in sample
%features
0
10
20
30
40
50
60
70
0
0.8
1.6
2.5
protein_coding
processed_pseudogene
lincRNA
TEC
antisense
miRNA
snoRNA
processed_transcript
snRNA
unprocessed_pseudogene
processed_pseudogene
sense_intronic
misc_RNA
rRNA
unprocessed_pseudogene
pseudogene
ectional_promoter_lncRNA
scaRNA
polymorphic_pseudogene
TR_J_gene
IG_V_gene
TR_V_gene
sense_overlapping
unitary_pseudogene
ribozyme
TR_C_gene
IG_V_pseudogene
ribed_unitary_pseudogene
TR_J_pseudogene
macro_lncRNA
TR_V_pseudogene