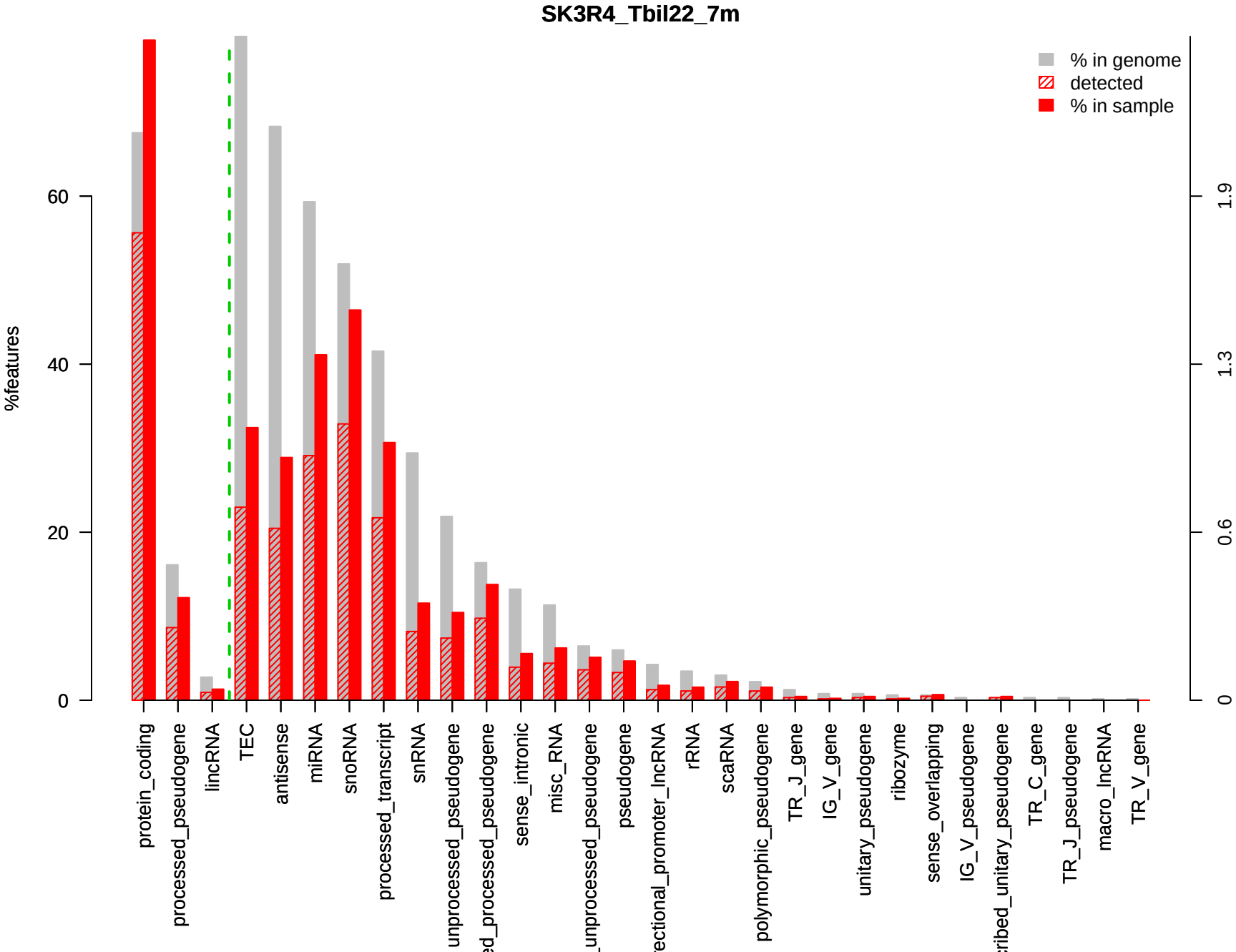
SK3R4_Tbil22_7m
% in genome
detected
% in sample
%features
0
20
40
60
0
0.6
1.3
1.9
protein_coding
processed_pseudogene
lincRNA
TEC
antisense
miRNA
snoRNA
processed_transcript
snRNA
unprocessed_pseudogene
processed_pseudogene
sense_intronic
misc_RNA
unprocessed_pseudogene
pseudogene
ectional_promoter_lncRNA
rRNA
scaRNA
polymorphic_pseudogene
TR_J_gene
IG_V_gene
unitary_pseudogene
ribozyme
sense_overlapping
IG_V_pseudogene
ribed_unitary_pseudogene
TR_C_gene
TR_J_pseudogene
macro_lncRNA
TR_V_gene