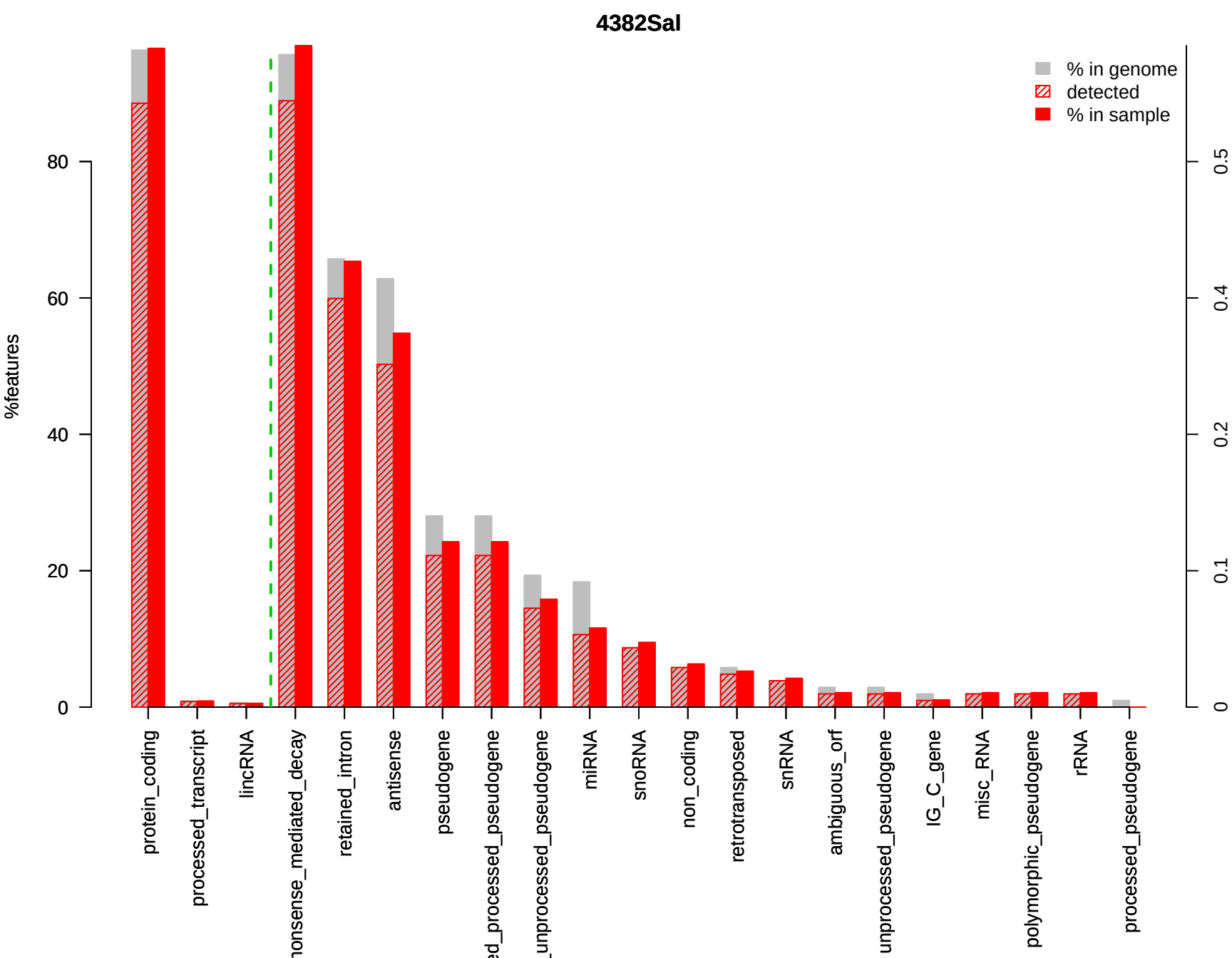
4382Sal
% in genome
detected
% in sample
%features
0
20
40
60
80
0
0.1
0.2
0.4
0.5
protein_coding
processed_transcript
lincRNA
nonsense_mediated_decay
retained_intron
antisense
pseudogene
ed_processed_pseudogene
unprocessed_pseudogene
miRNA
snoRNA
non_coding
retrotransposed
snRNA
ambiguous_orf
unprocessed_pseudogene
IG_C_gene
misc_RNA
polymorphic_pseudogene
rRNA
processed_pseudogene