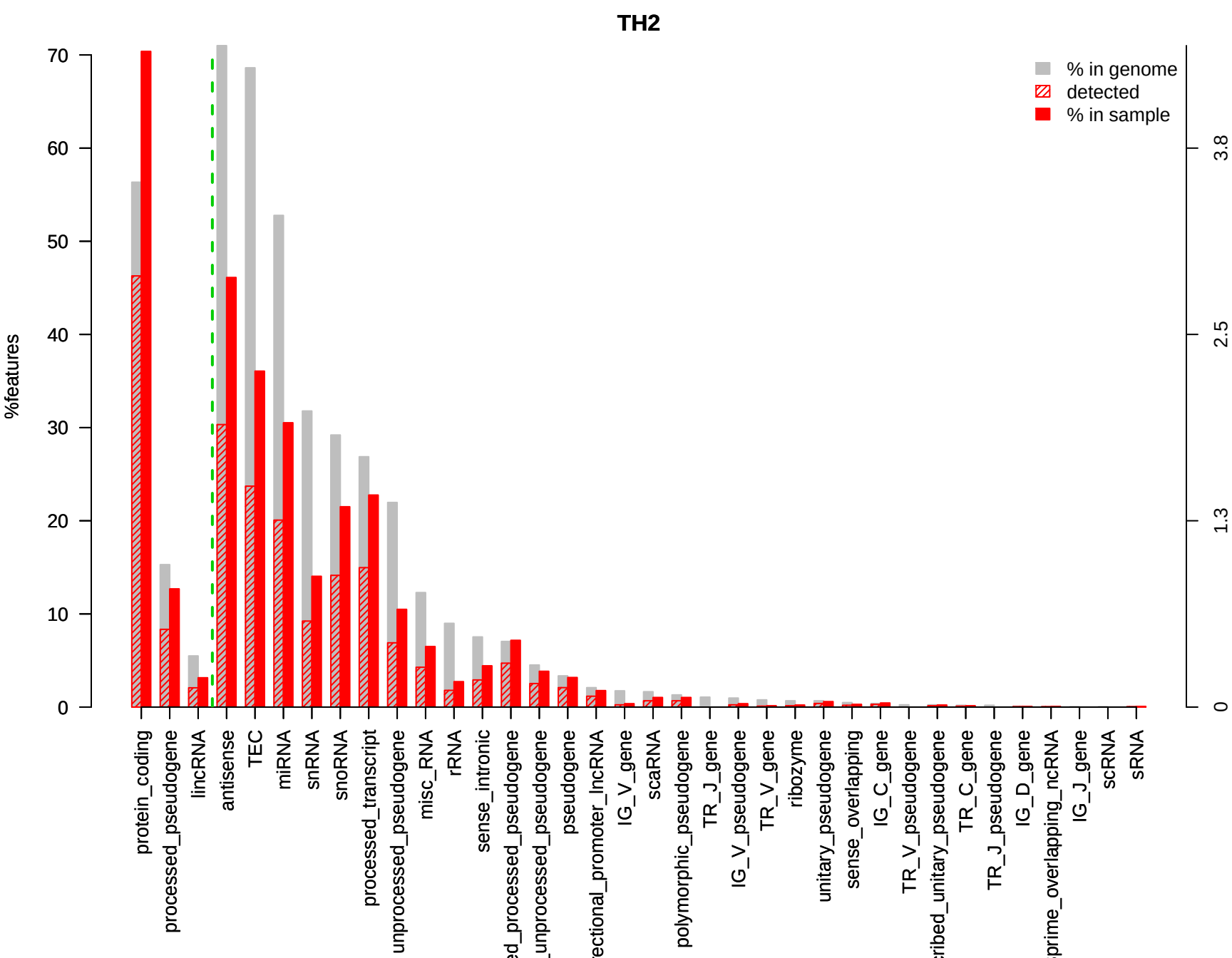
TH2
%features
% in genome
detected
% in sample
0
10
20
30
40
50
60
70
0
1.3
2.5
3.8
protein_coding
processed_pseudogene
lincRNA
antisense
TEC
miRNA
snRNA
snoRNA
processed_transcript
unprocessed_pseudogene
misc_RNA
rRNA
sense_intronic
ed_processed_pseudogene
unprocessed_pseudogene
pseudogene
ectional_promoter_lncRNA
IG_V_gene
scaRNA
polymorphic_pseudogene
TR_J_gene
IG_V_pseudogene
TR_V_gene
ribozyme
unitary_pseudogene
sense_overlapping
IG_C_gene
TR_V_pseudogene
ribed_unitary_pseudogene
TR_C_gene
TR_J_pseudogene
IG_D_gene
prime_overlapping_ncRNA
IG_J_gene
scRNA
sRNA