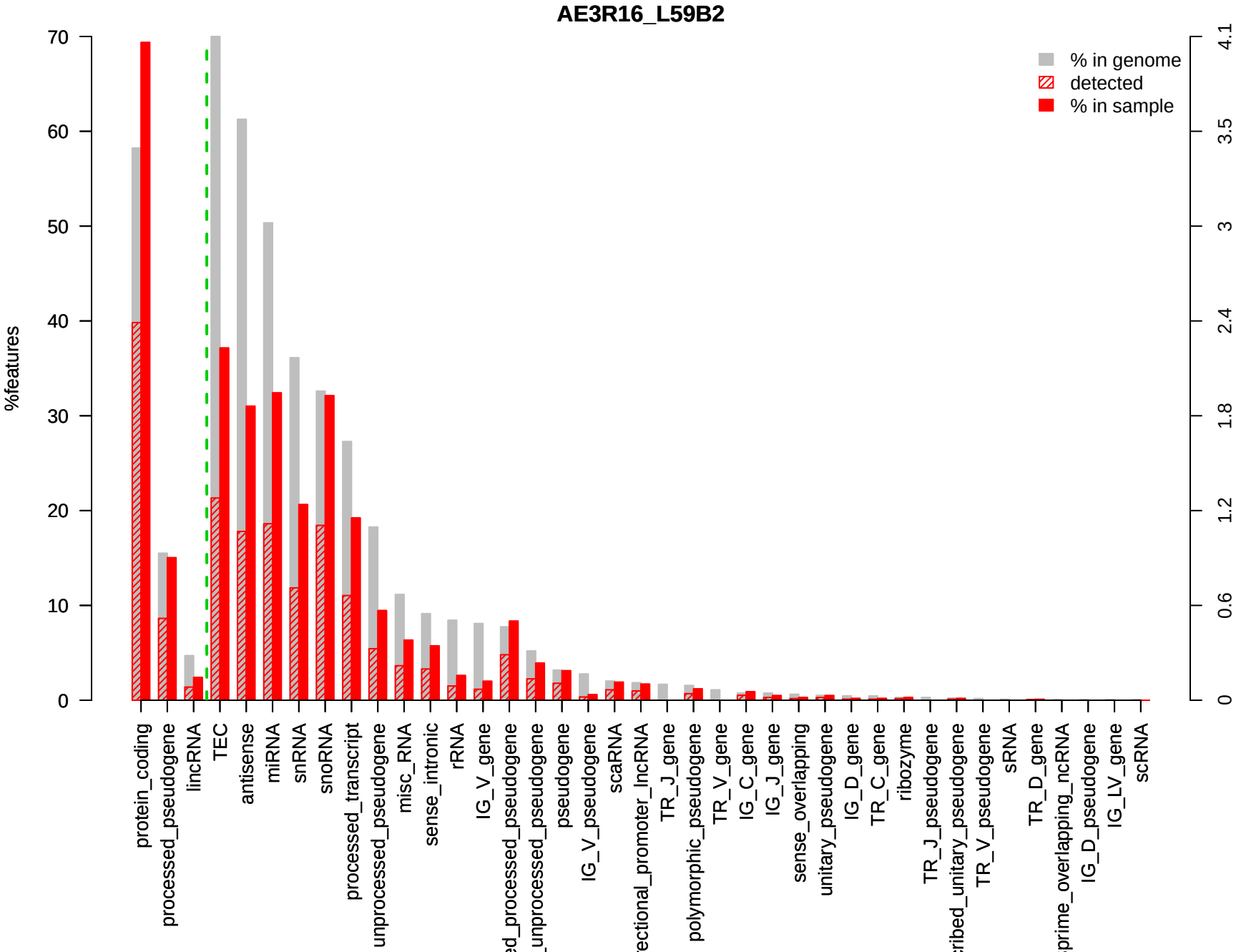
AE3R16_L59B2
% in genome
detected
% in sample
%features
0
10
20
30
40
50
60
70
0
0.6
1.2
1.8
2.4
3
3.5
4.1
protein_coding
processed_pseudogene
lncRNA
TEC
antisense
miRNA
snRNA
snoRNA
processed_transcript
unprocessed_pseudogene
misc_RNA
sense_intronic
rRNA
IG_V_gene
ed_processed_pseudogene
unprocessed_pseudogene
pseudogene
IG_V_pseudogene
scaRNA
ectional_promoter_lncRNA
TR_J_gene
polymorphic_pseudogene
TR_V_gene
IG_C_gene
IG_J_gene
sense_overlapping
unitary_pseudogene
IG_D_gene
TR_C_gene
ribozyme
TR_J_pseudogene
ribed_unitary_pseudogene
TR_V_pseudogene
sRNA
TR_D_gene
prime_overlapping_ncRNA
IG_D_pseudogene
IG_LV_gene
scRNA