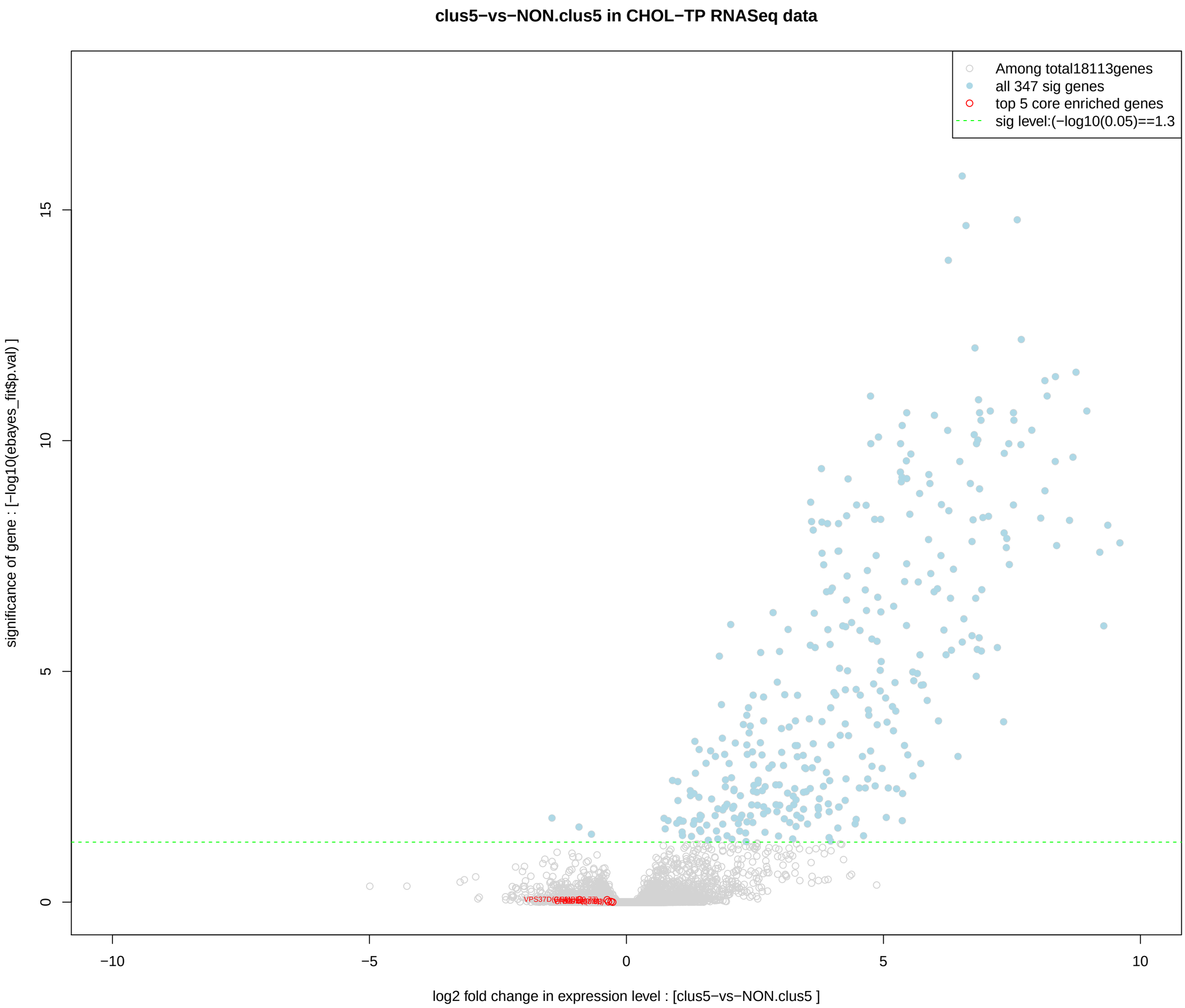
clus5−vs−NON.clus5 in CHOL−TP RNASeq data
Among total18113genes
all 347 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus5−vs−NON.clus5]
0
5
10
15
−10
−5
0
5
10