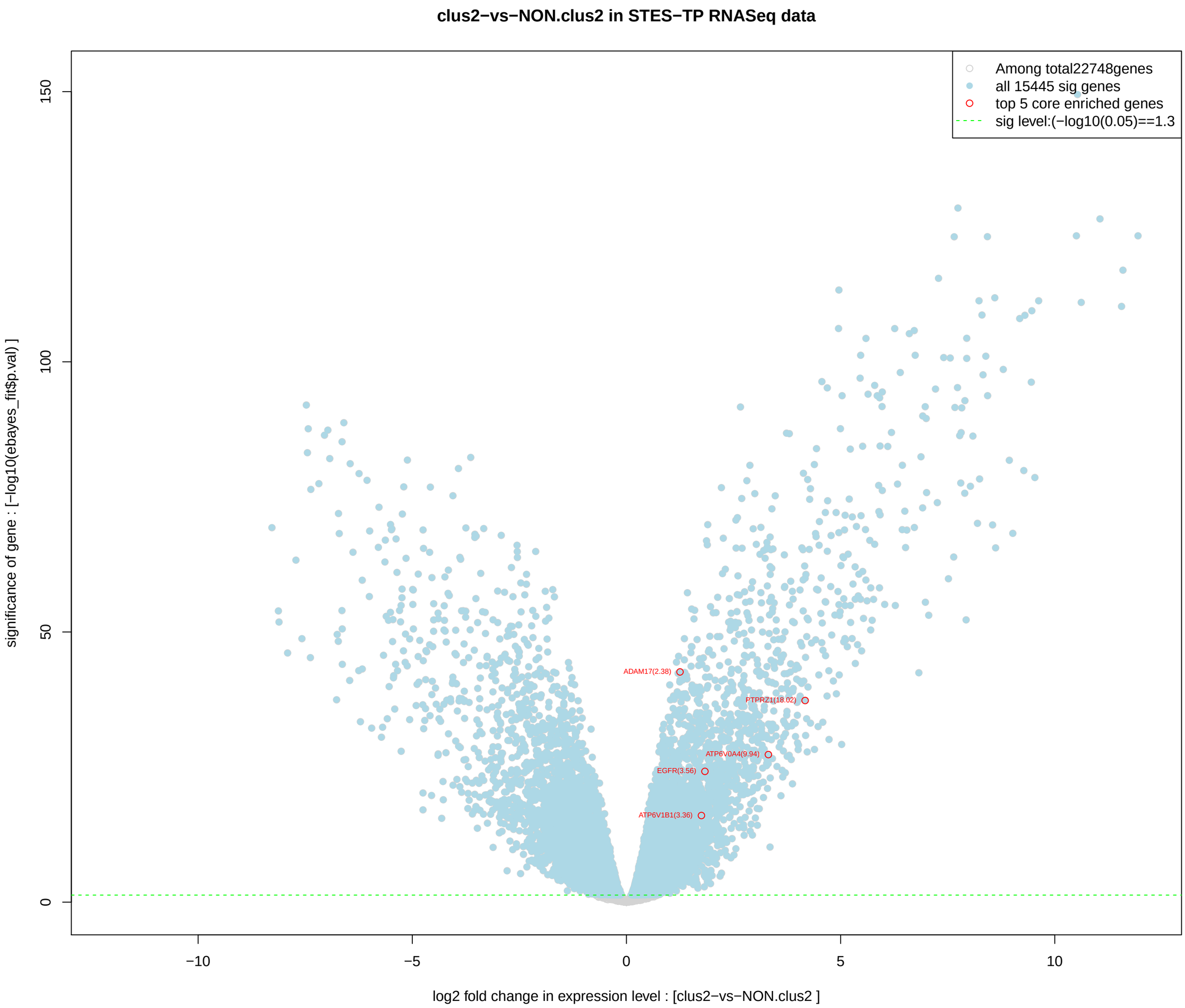
clus2−vs−NON.clus2 in STES−TP RNASeq data
Among total22748genes
all 15445 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ADAM17(2.38)
PTPRZ1(18.02)
ATP6V0A4(9.94)
EGFR(3.56)
ATP6V1B1(3.36)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
50
100
150
−10
−5
0
5
10