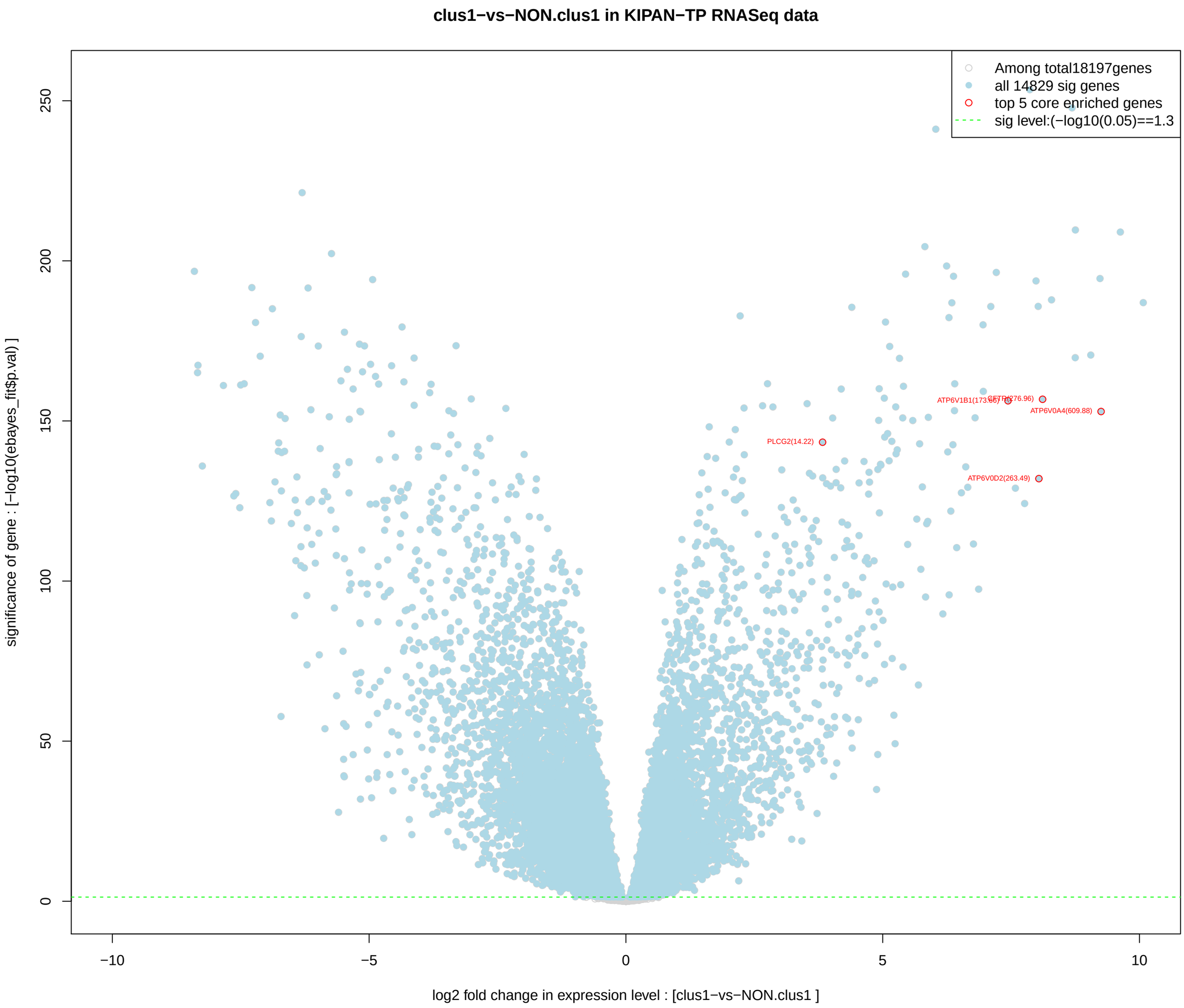
clus1−vs−NON.clus1 in KIPAN−TP RNASeq data
Among total18197genes
all 14829 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ATP6V1B1(173.65)
CFTR(276.96)
ATP6V0A4(609.88)
PLCG2(14.22)
ATP6V0D2(263.49)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
50
100
150
200
250
−10
−5
0
5
10