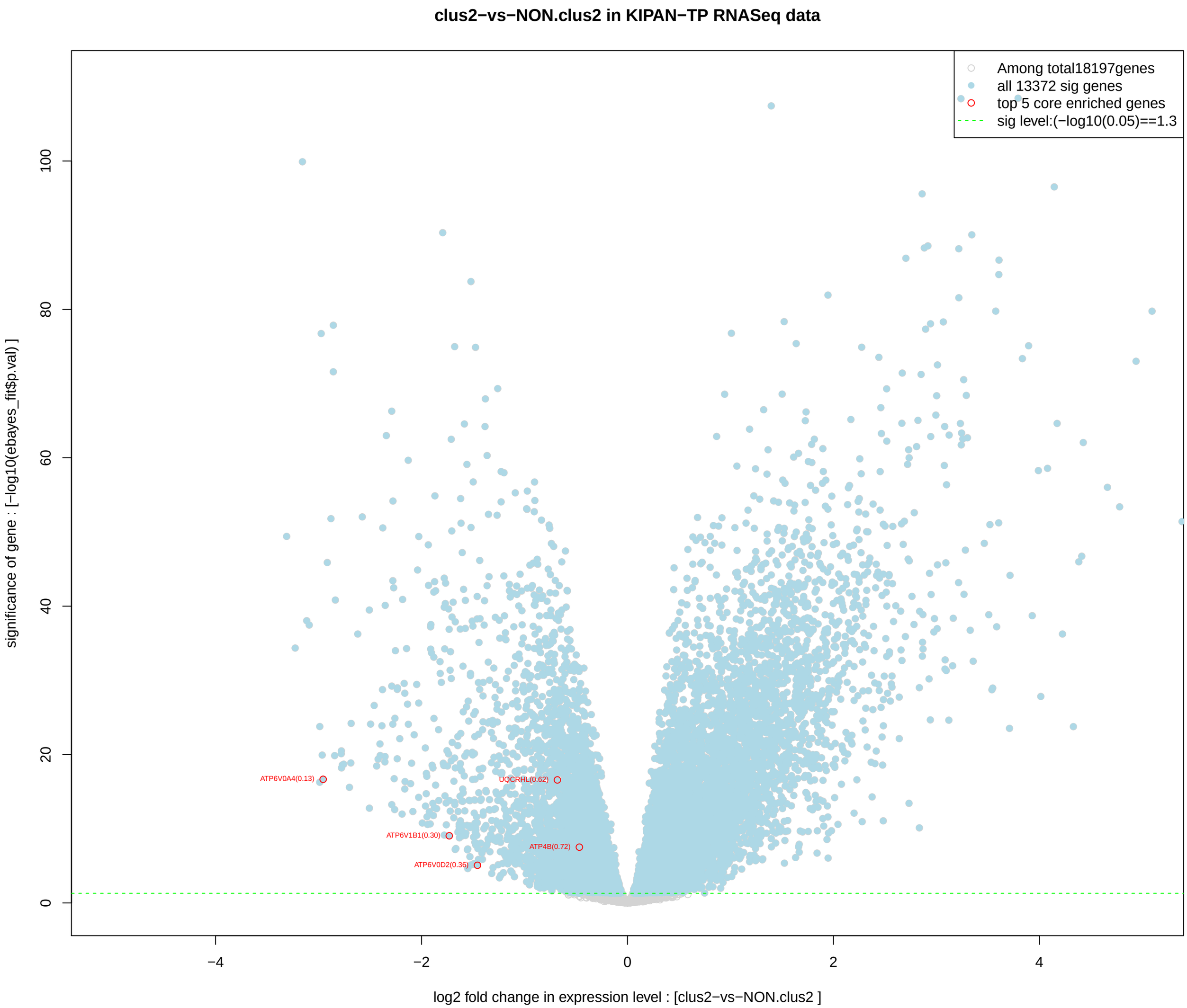
clus2−vs−NON.clus2 in KIPAN−TP RNASeq data
Among total18197genes
all 13372 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ATP6V0A4(0.13)
UQCRHL(0.62)
ATP6V1B1(0.30)
ATP4B(0.72)
ATP6V0D2(0.36)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
20
40
60
80
100
−4
−2
0
2
4