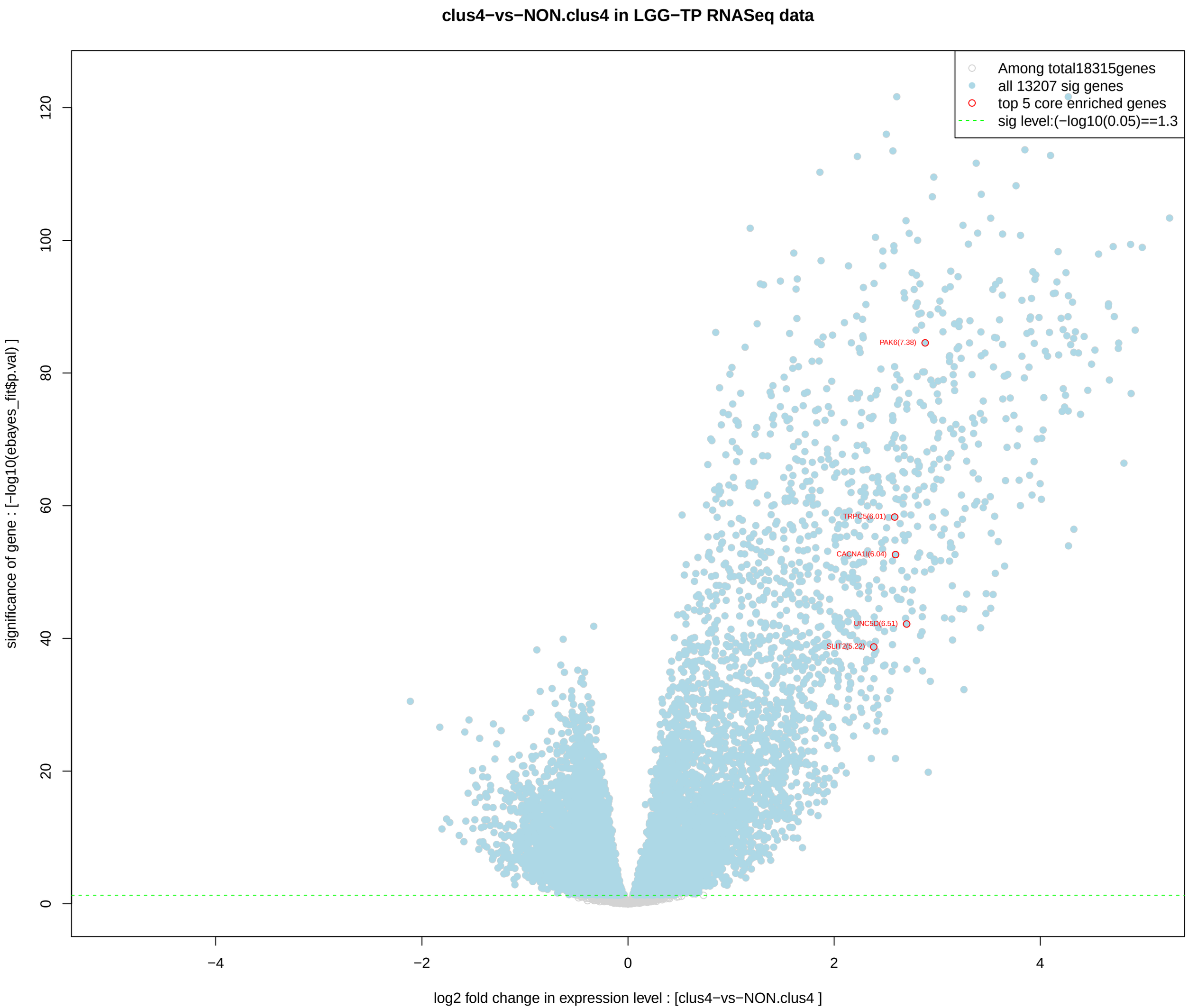
clus4−vs−NON.clus4 in LGG−TP RNASeq data
Among total18315genes
all 13207 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PAK6(7.38)
TRPC5(6.01)
CACNA1I(6.04)
UNC5D(6.51)
SLIT2(5.22)
significance of gene : [−log10(ebayes_fit$p.val)]
0
20
40
60
80
100
120
−4
−2
0
2
4
log2 fold change in expression level : [clus4−vs−NON.clus4]