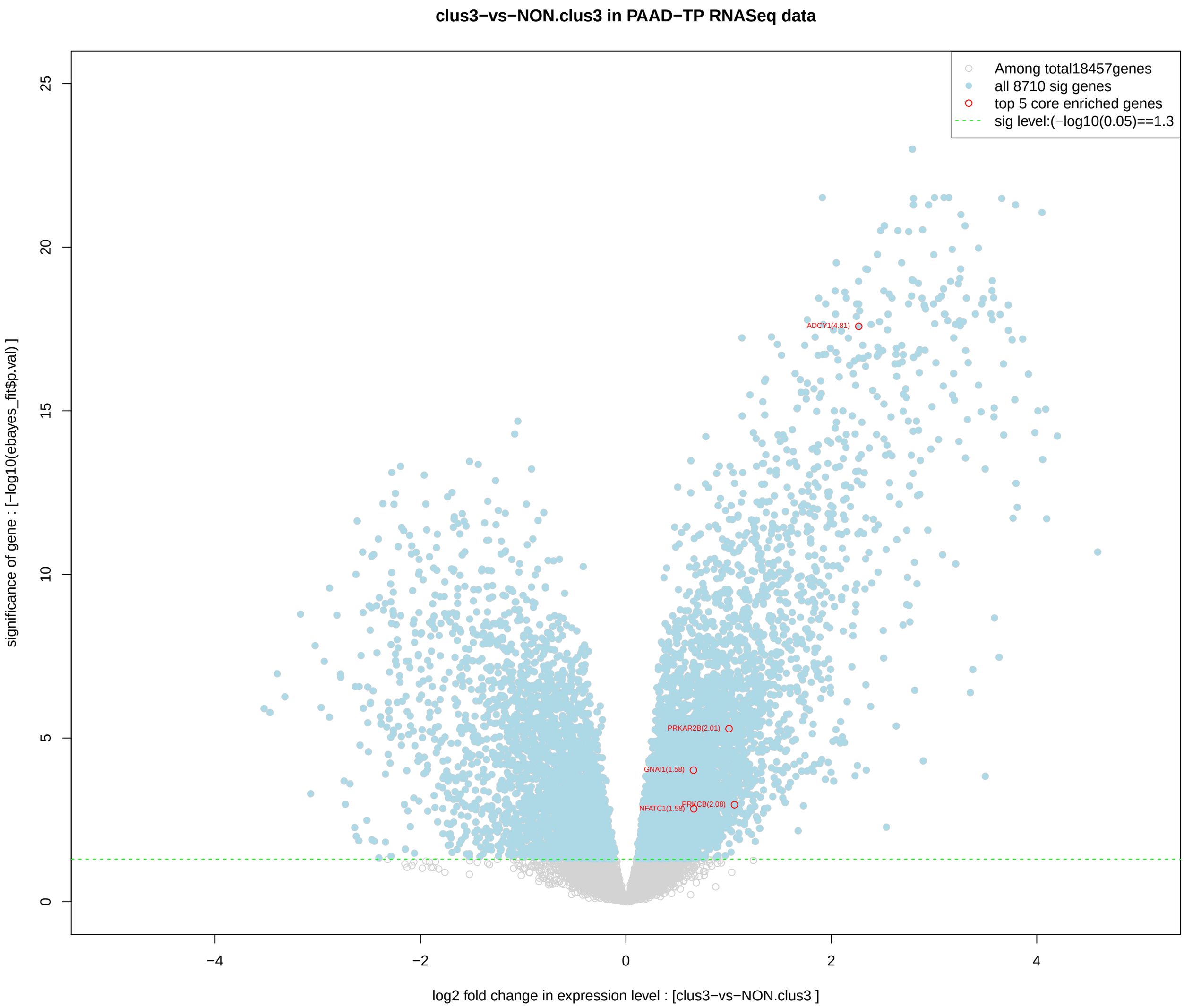
clus3−vs−NON.clus3 in PAAD−TP RNASeq data
Among total18457genes
all 8710 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ADCY1(4.81)
PRKAR2B(2.01)
GNAI1(1.58)
PRKCB(2.08)
NFATC1(1.58)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
5
10
15
20
25
−4
−2
0
2
4