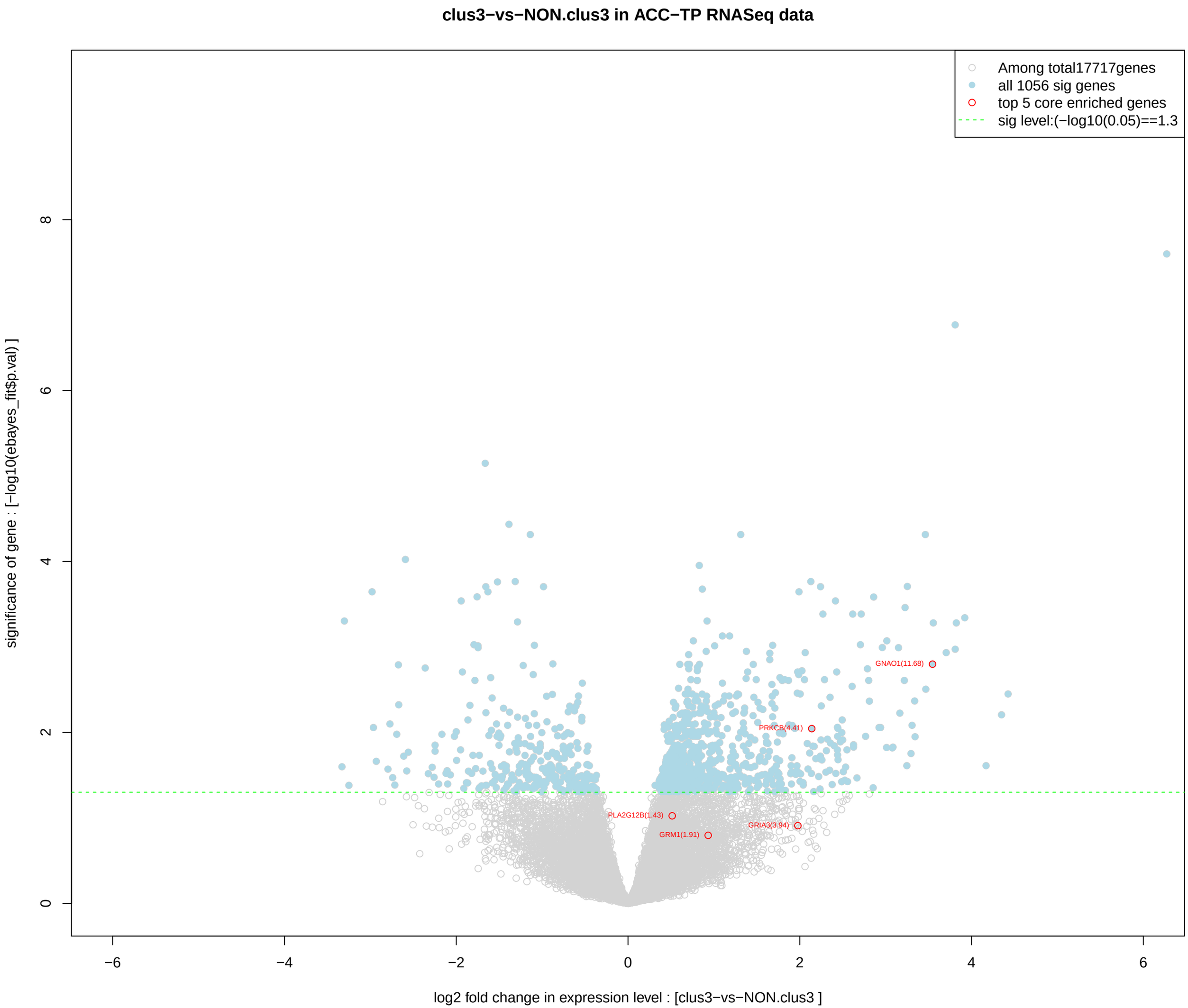
clus3−vs−NON.clus3 in ACC−TP RNASeq data
Among total17717genes
all 1056 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
GNAO1(11.68)
PRKCB(4.41)
PLA2G12B(1.43)
GRM1(1.91)
GRIA3(3.94)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
2
4
6
8
−6
−4
−2
0
2
4
6