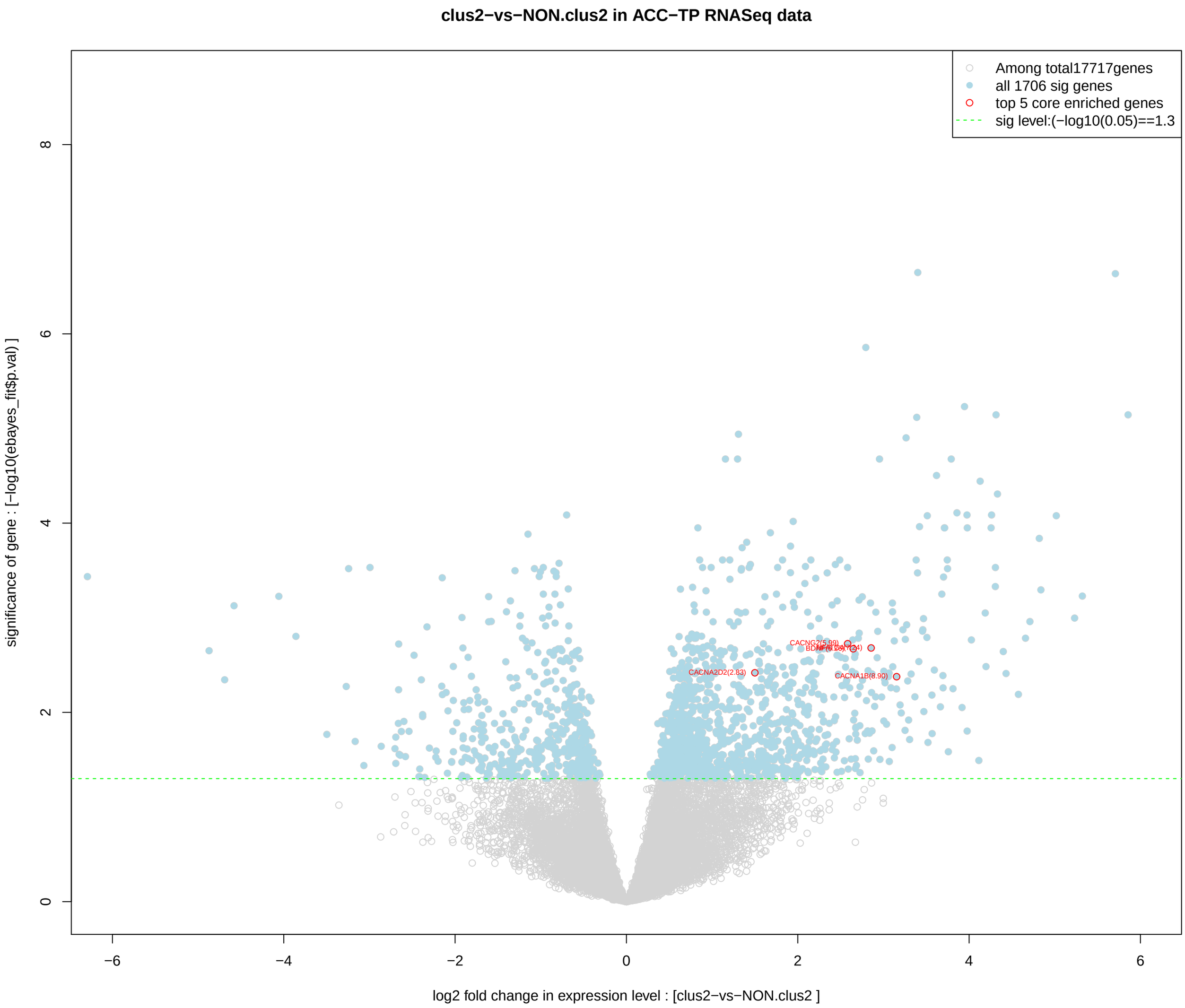
clus2−vs−NON.clus2 in ACC−TP RNASeq data
Among total17717genes
all 1706 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CACNG2(5.99)
BDNF(6.26)
CACNA2D2(2.83)
CACNA1B(8.90)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
2
4
6
8
−6
−4
−2
0
2
4
6