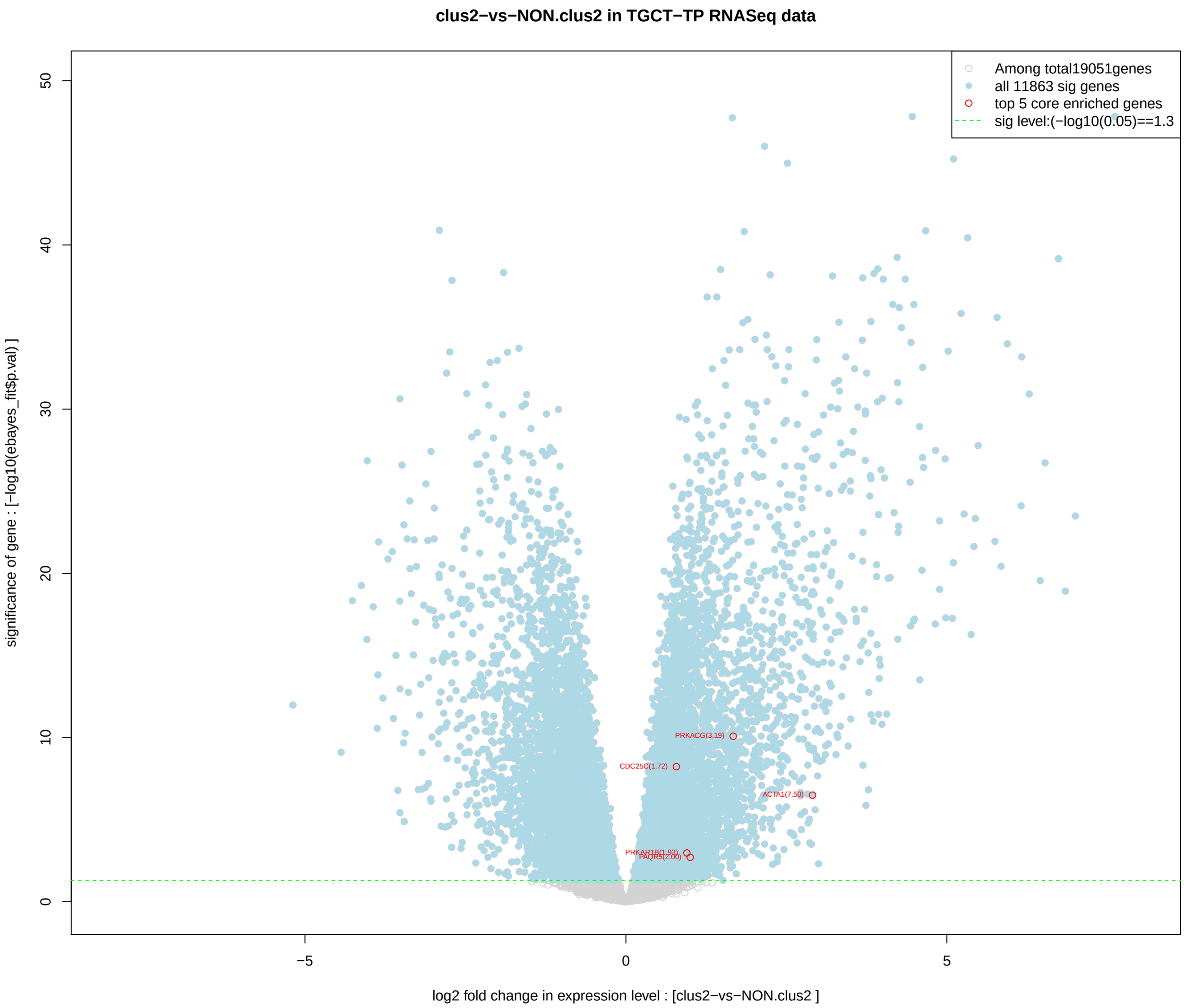
clus2−vs−NON.clus2 in TGCT−TP RNASeq data
Among total19051genes
all 11863 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PRKACG(3.19)
CDC25C(1.72)
ACTA1(7.50)
PRKAR1B(1.93)
PAQR5(2.00)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
10
20
30
40
50
−5
0
5