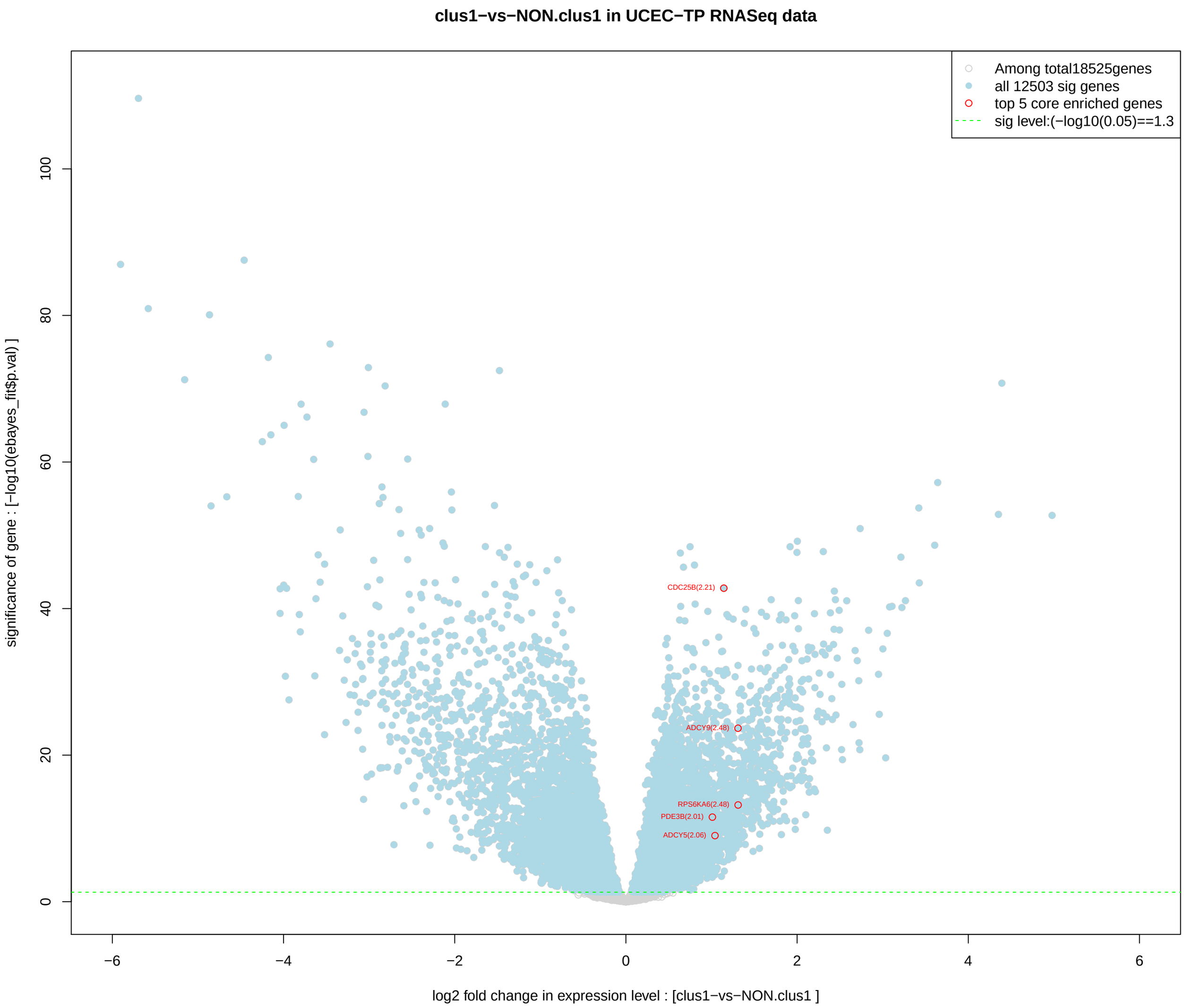
clus1−vs−NON.clus1 in UCEC−TP RNASeq data
Among total18525genes
all 12503 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CDC25B(2.21)
ADCY9(2.48)
RPS6KA6(2.48)
PDE3B(2.01)
ADCY5(2.06)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
20
40
60
80
100
−6
−4
−2
0
2
4
6