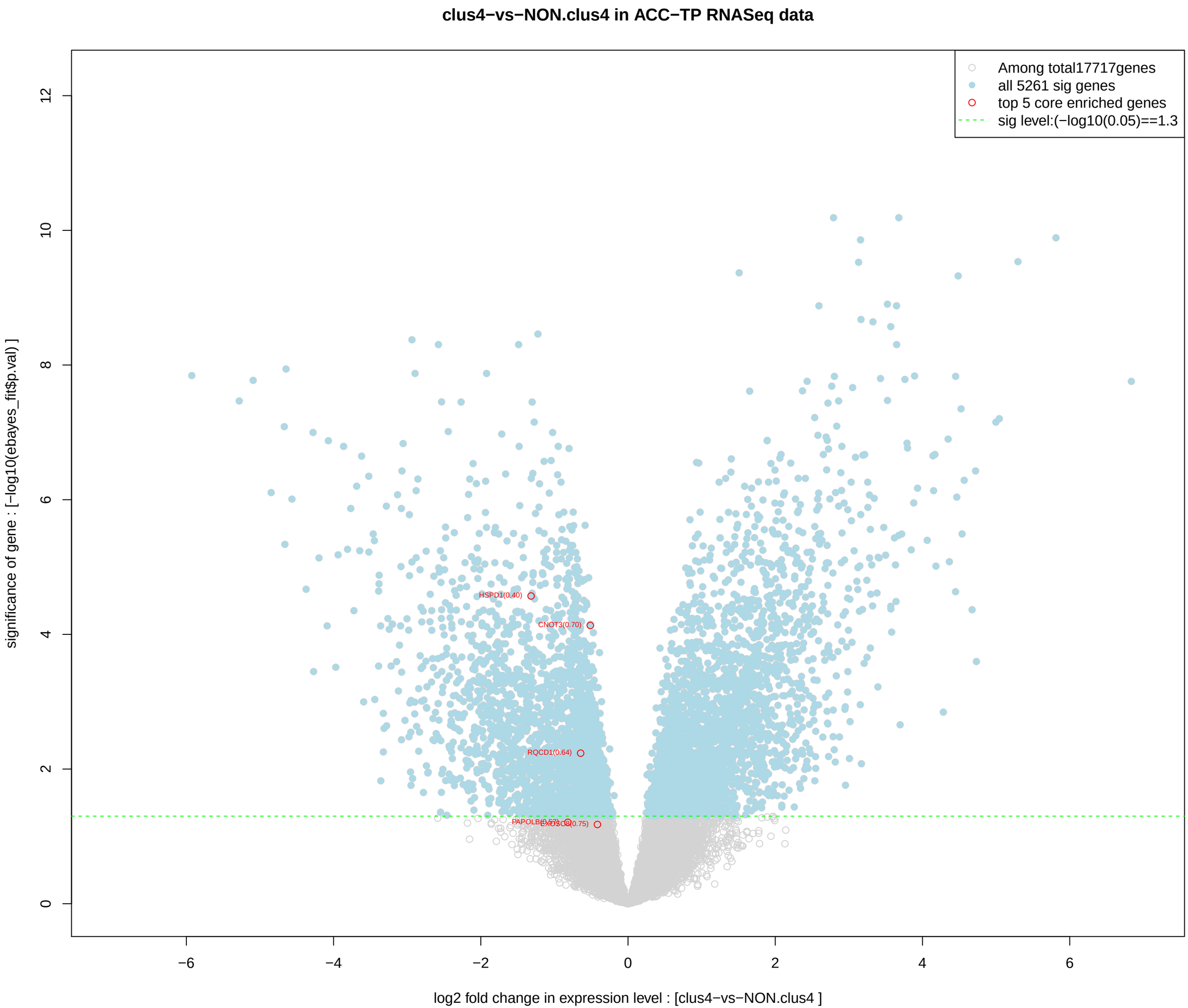
clus4−vs−NON.clus4 in ACC−TP RNASeq data
Among total17717genes
all 5261 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
HSPD1(0.40)
CNOT3(0.70)
RQCD1(0.64)
PAPOLB(0.73)
EXOSC8(0.75)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0
2
4
6
8
10
12
−6
−4
−2
0
2
4
6