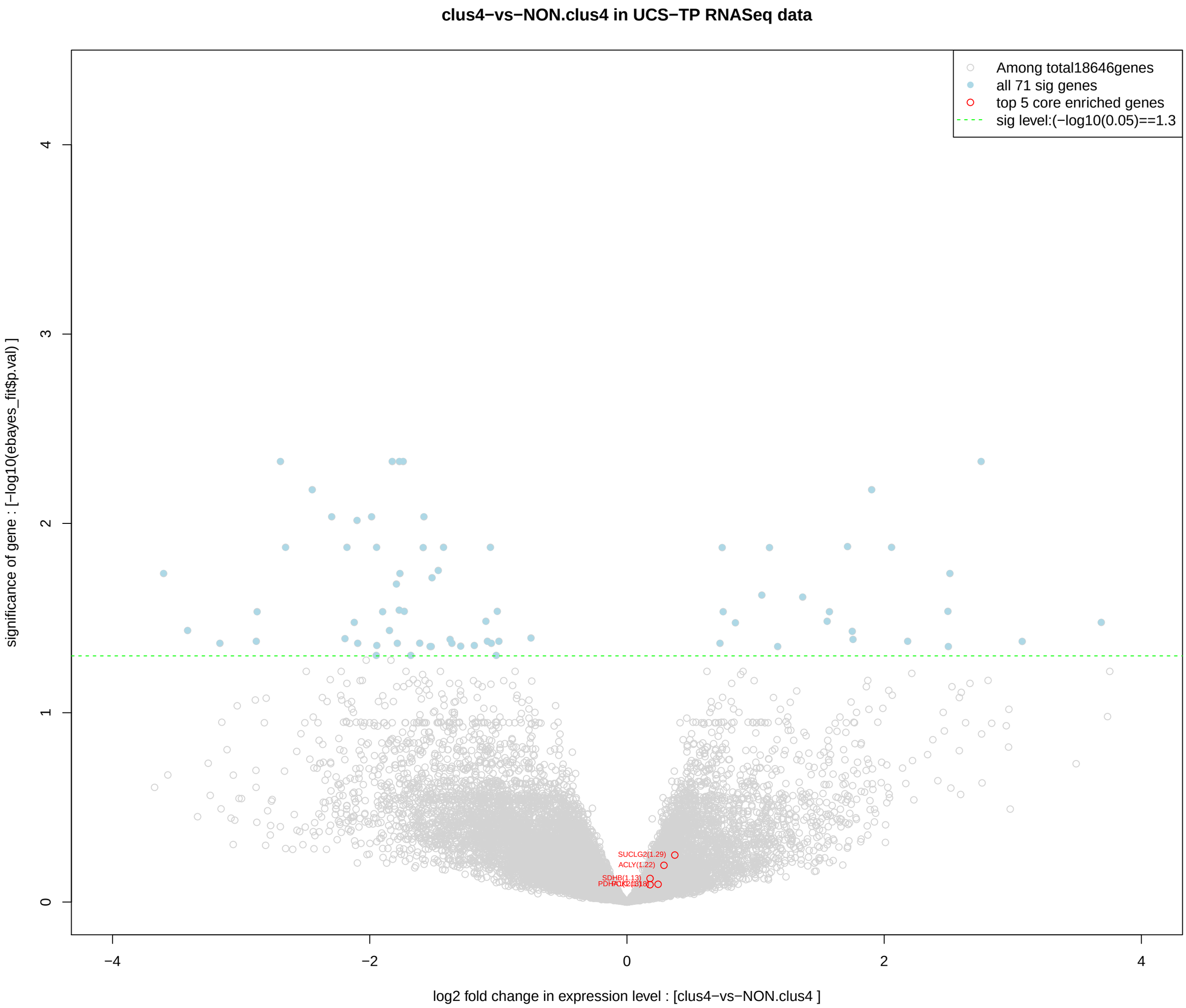
clus4−vs−NON.clus4 in UCS−TP RNASeq data
Among total18646genes
all 71 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
SUCLG2(1.29)
ACLY(1.22)
SDHB(1.13)
PDHA1(1.18)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
−4
−2
0
2
4
0
1
2
3
4