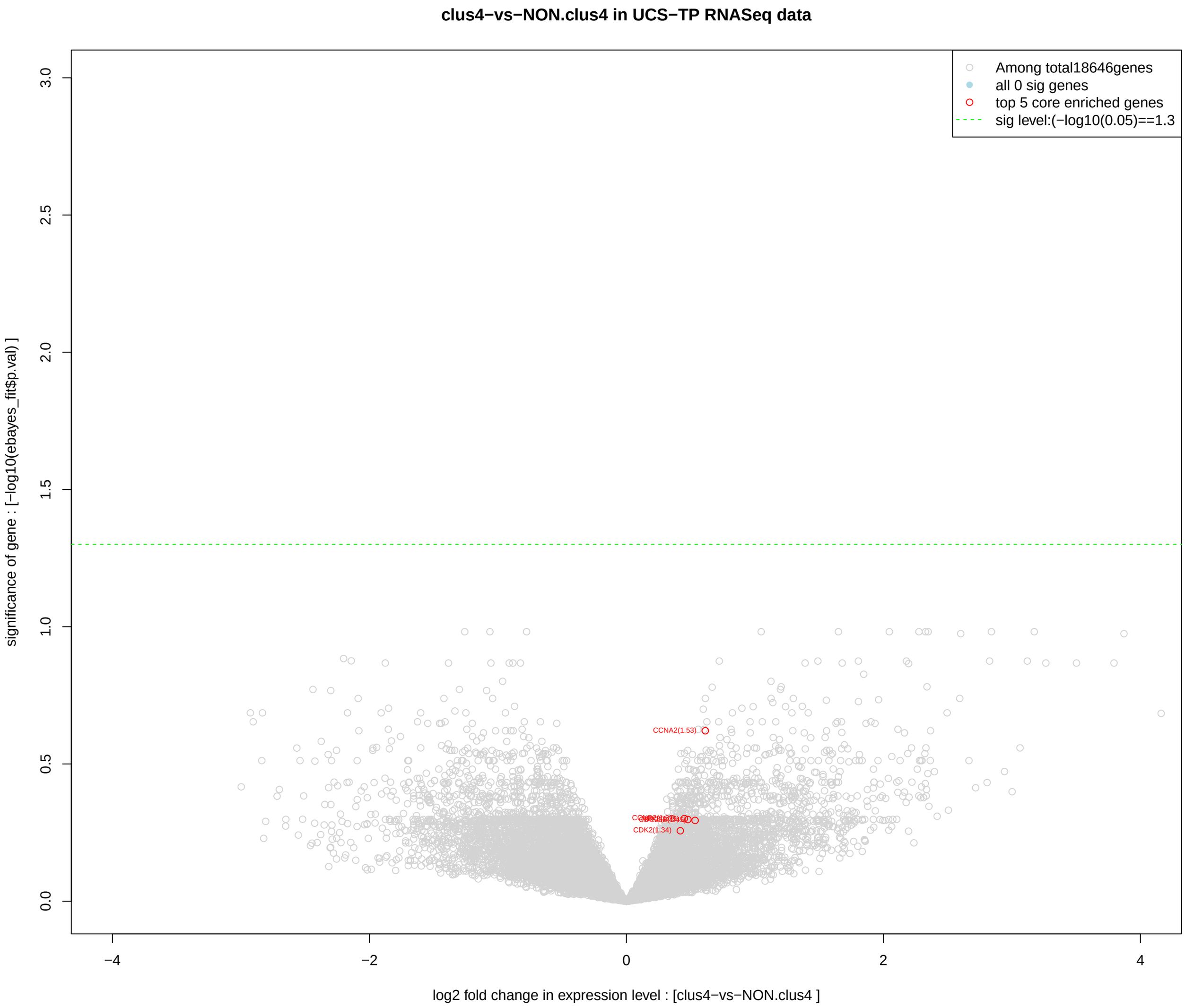
clus4−vs−NON.clus4 in UCS−TP RNASeq data
Among total18646genes
all 0 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CCNA2(1.53)
CDK2(1.34)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0.0
0.5
1.0
1.5
2.0
2.5
3.0
−4
−2
0
2
4