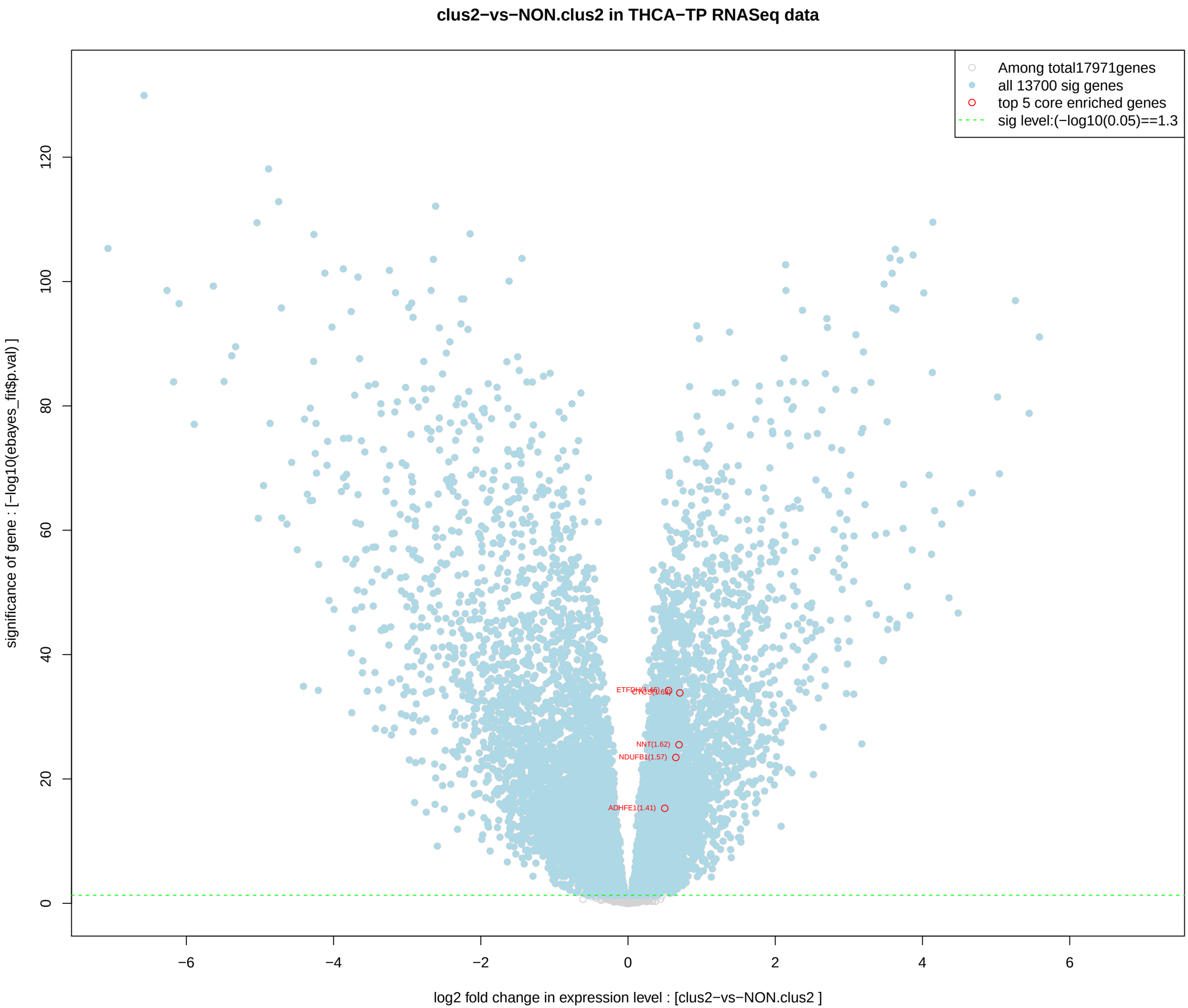
clus2−vs−NON.clus2 in THCA−TP RNASeq data
Among total17971genes
all 13700 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ETFDH(1.46)
CYCS(1.66)
NNT(1.62)
NDUFB1(1.57)
ADHFE1(1.41)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
20
40
60
80
100
120
−6
−4
−2
0
2
4
6