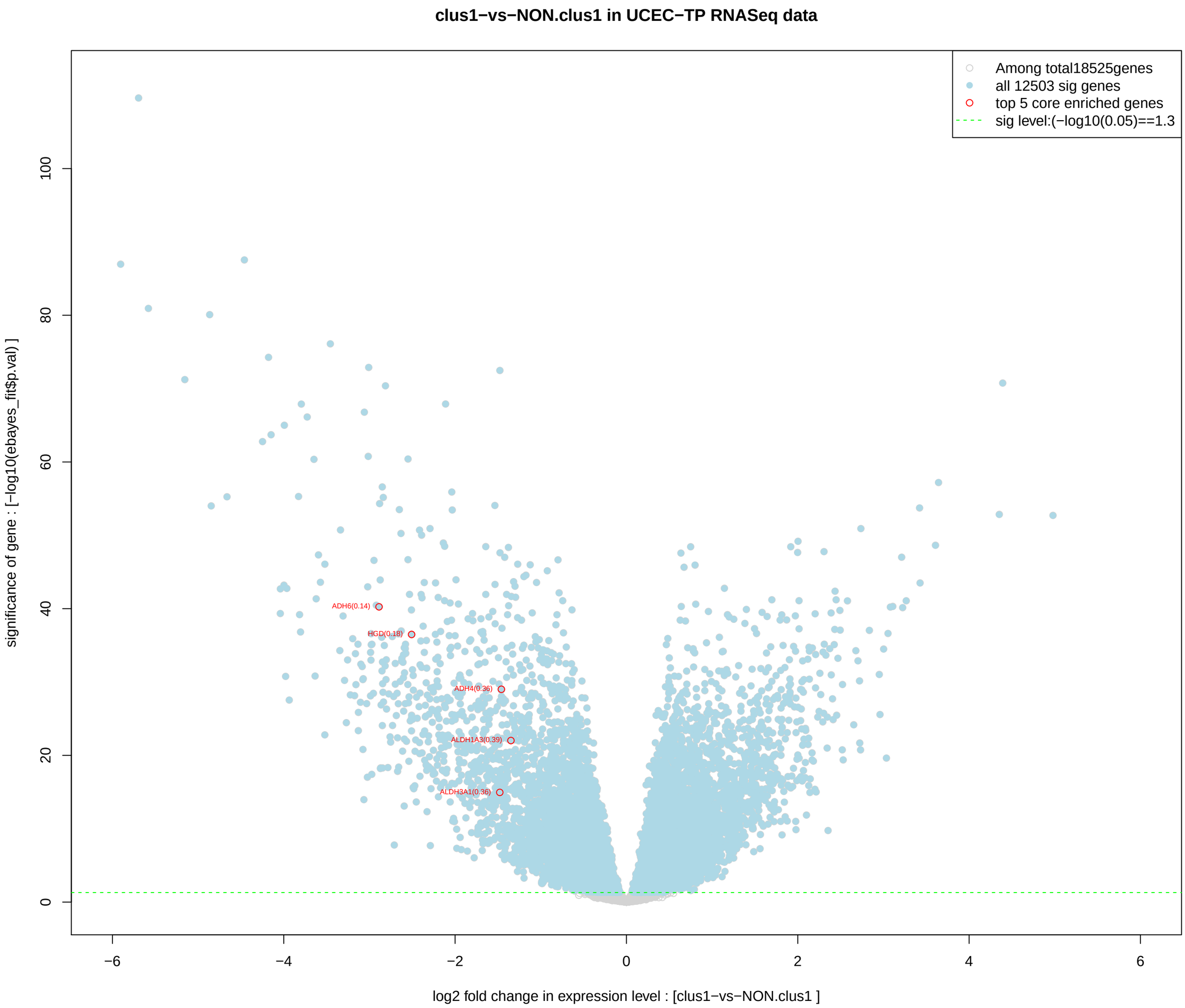
clus1−vs−NON.clus1 in UCEC−TP RNASeq data
Among total18525genes
all 12503 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ADH6(0.14)
HGD(0.18)
ADH4(0.36)
ALDH1A3(0.39)
ALDH3A1(0.36)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
20
40
60
80
100
−6
−4
−2
0
2
4
6