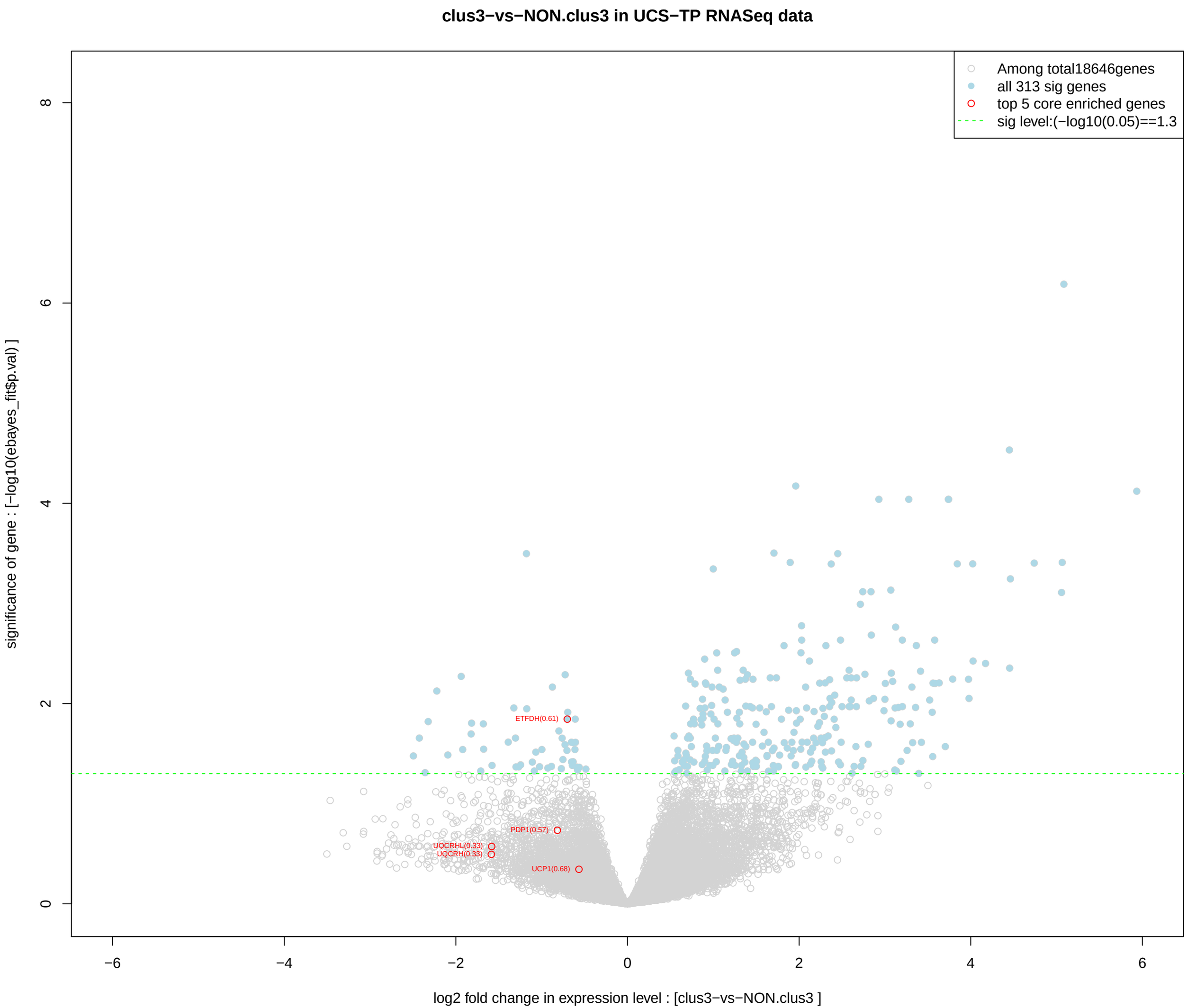
clus3−vs−NON.clus3 in UCS−TP RNASeq data
Among total18646genes
all 313 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ETFDH(0.61)
PDP1(0.57)
UQCRHL(0.33)
UQCRH(0.33)
UCP1(0.68)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
2
4
6
8
−6
−4
−2
0
2
4
6