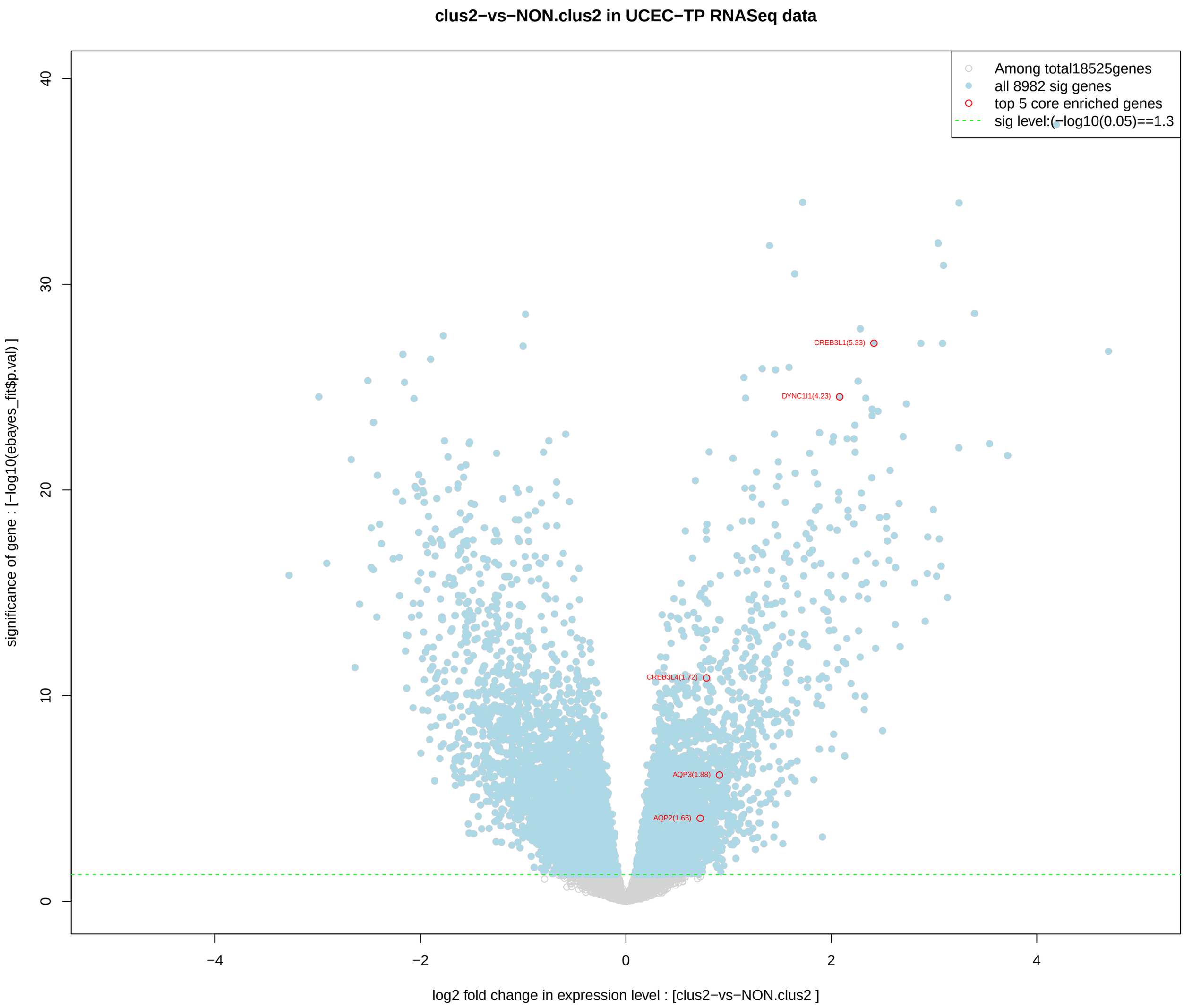
clus2−vs−NON.clus2 in UCEC−TP RNASeq data
Among total18525genes
all 8982 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CREB3L1(5.33)
DYNC1I1(4.23)
CREB3L4(1.72)
AQP3(1.88)
AQP2(1.65)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
10
20
30
40
−4
−2
0
2
4