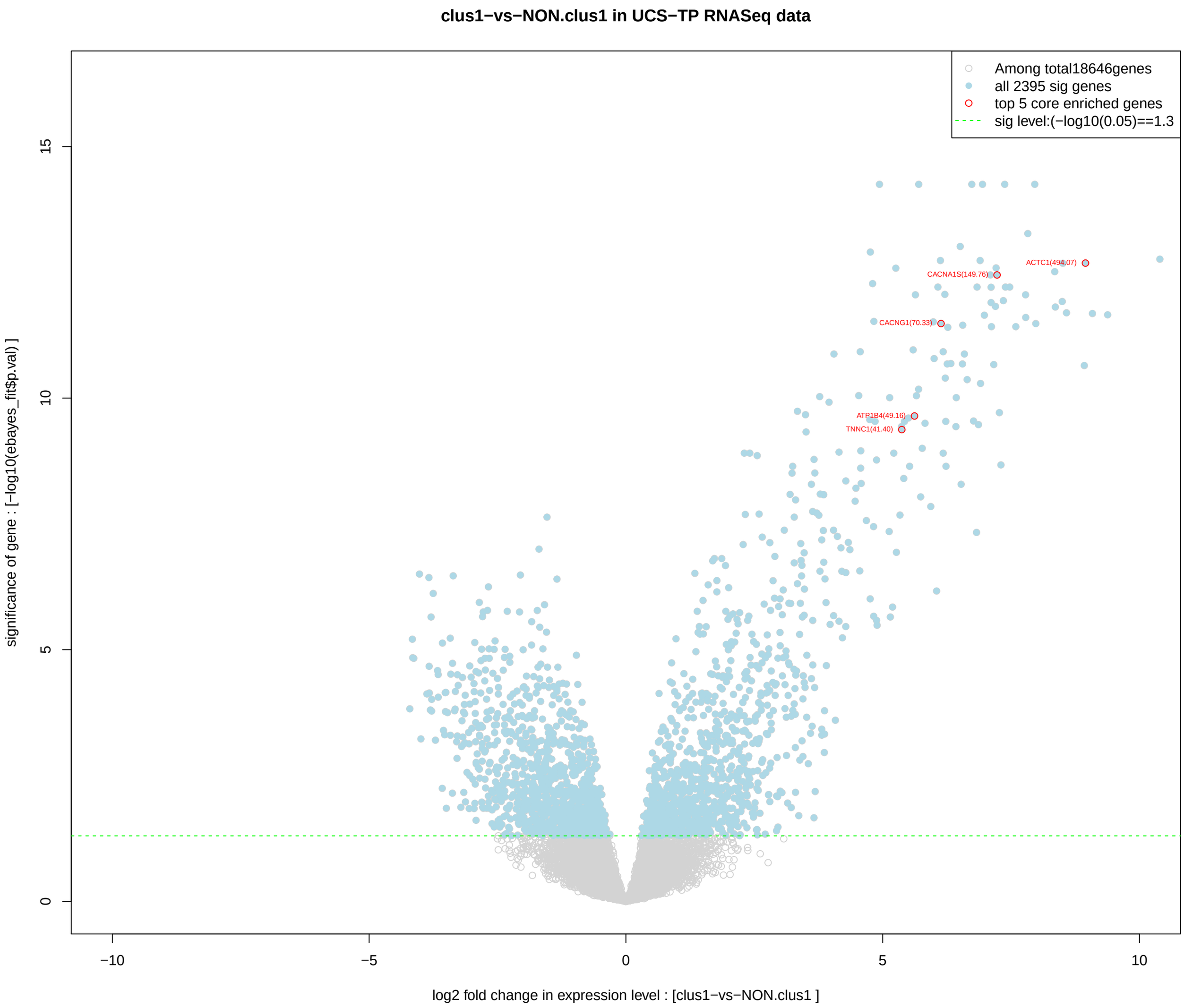
clus1−vs−NON.clus1 in UCS−TP RNASeq data
Among total18646genes
all 2395 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ACTC1(494.07)
CACNA1S(149.76)
CACNG1(70.33)
ATP1B4(49.16)
TNNC1(41.40)
significance of gene : [−log10(ebayes_fit$p.val)]
0
5
10
15
−10
−5
0
5
10
log2 fold change in expression level : [clus1−vs−NON.clus1]