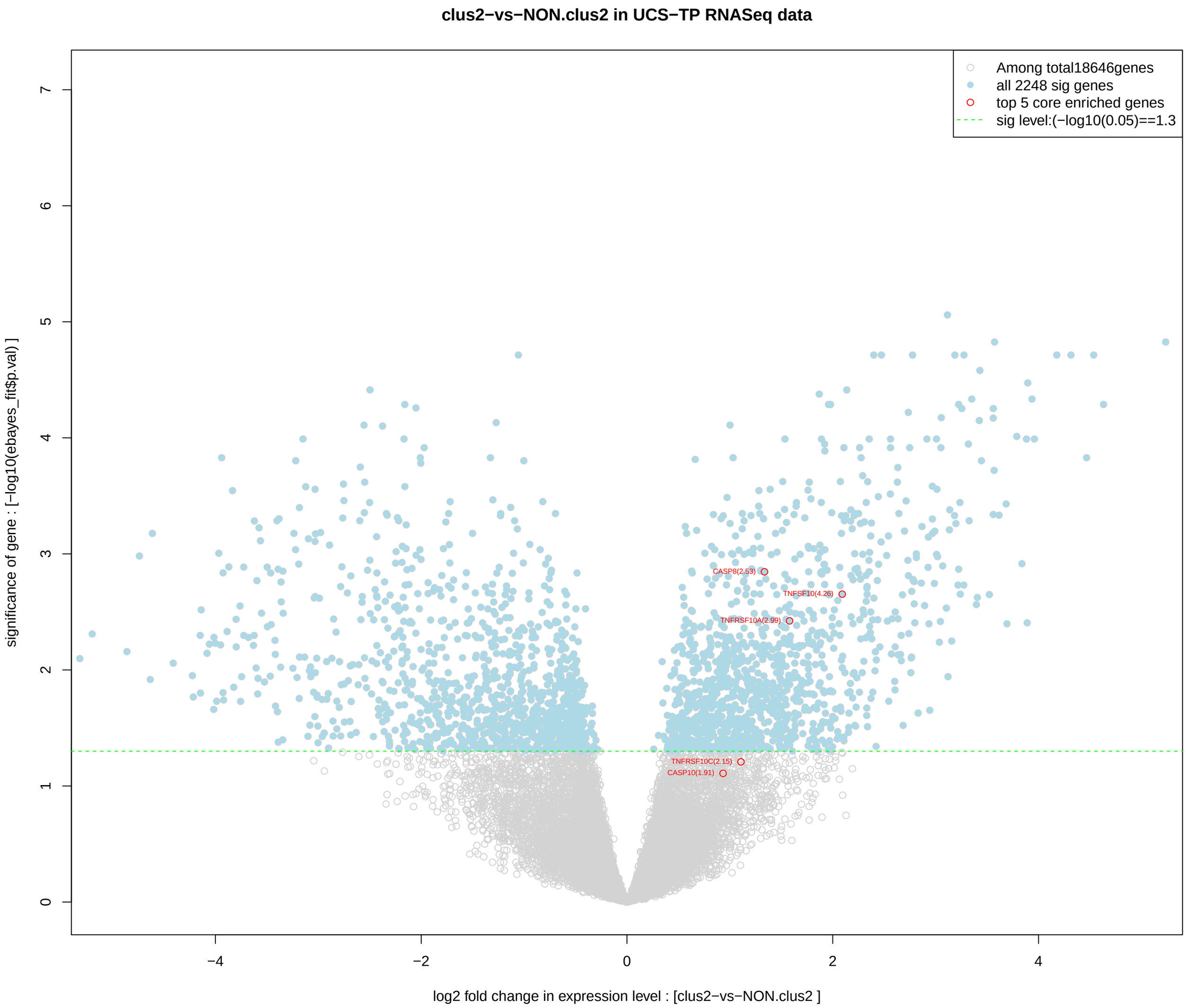
clus2−vs−NON.clus2 in UCS−TP RNASeq data
Among total18646genes
all 2248 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CASP8(2.53)
TNFSF10(4.26)
TNFRSF10A(2.99)
TNFRSF10C(2.15)
CASP10(1.91)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
1
2
3
4
5
6
7
−4
−2
0
2
4