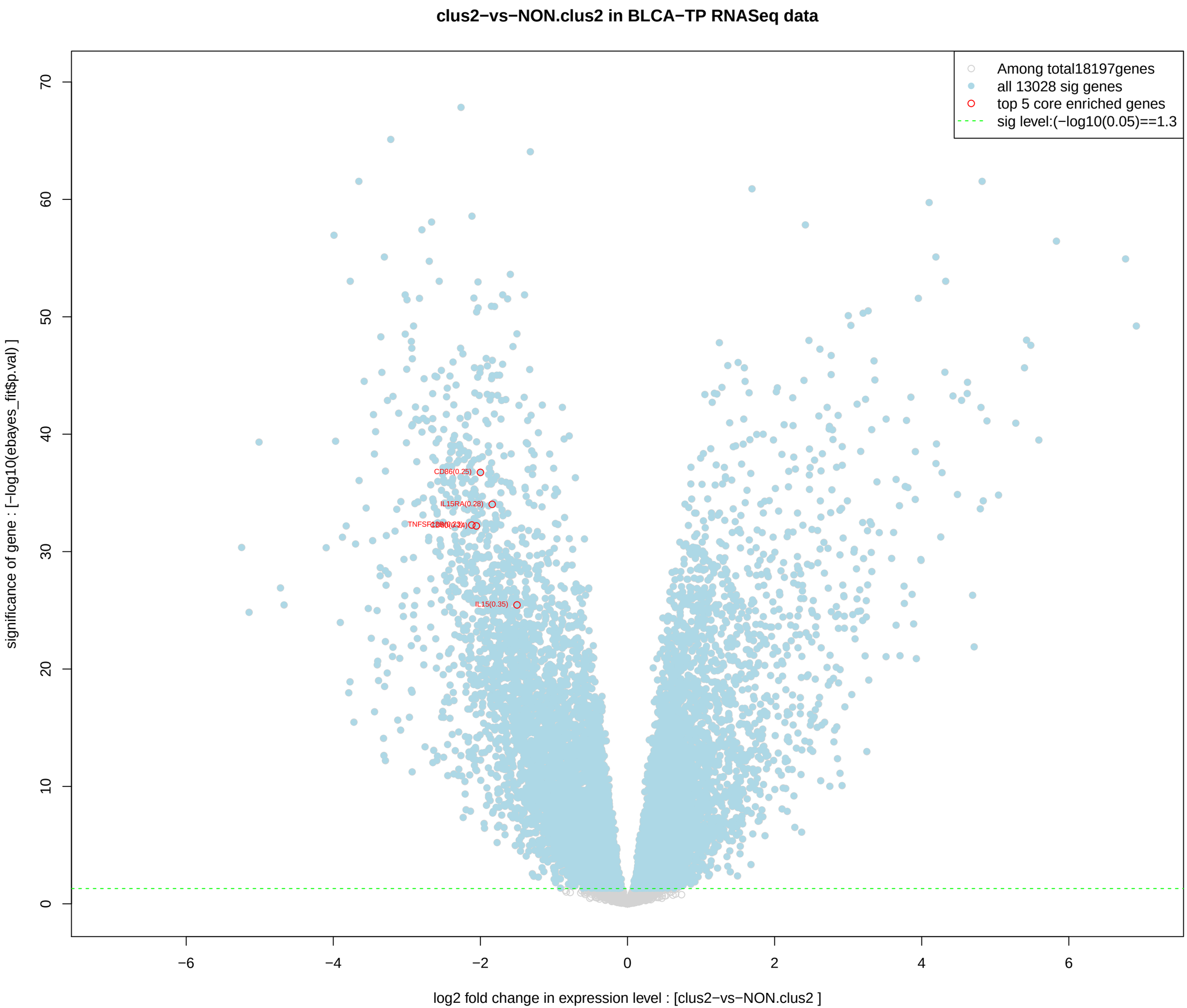
clus2−vs−NON.clus2 in BLCA−TP RNASeq data
Among total18197genes
all 13028 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CD86(0.25)
IL15RA(0.28)
TNFSF13B(0.22)
CD8(0.24)
IL15(0.35)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
10
20
30
40
50
60
70
−6
−4
−2
0
2
4
6