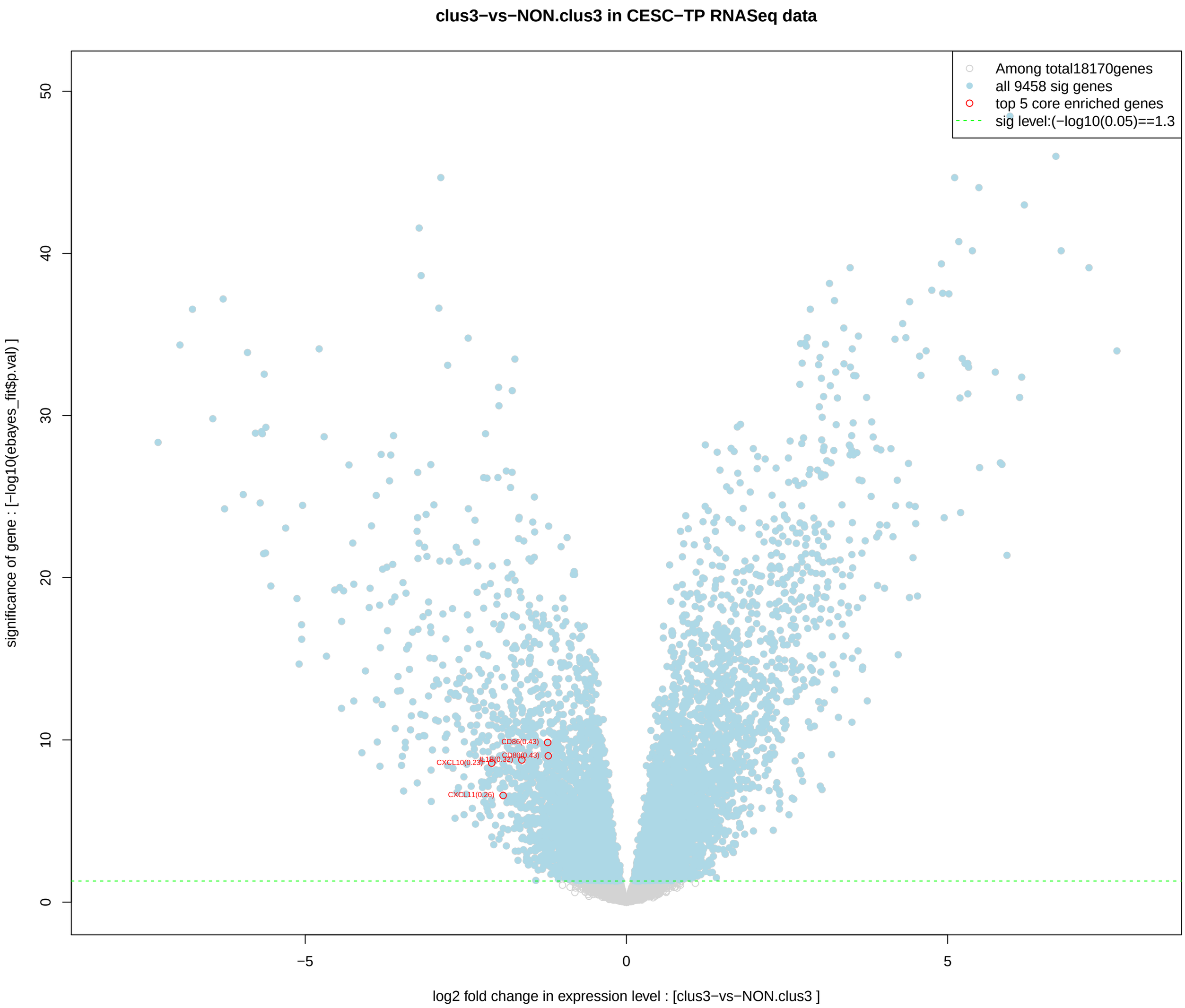
clus3−vs−NON.clus3 in CESC−TP RNASeq data
Among total18170genes
all 9458 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CD86(0.43)
CD80(0.43)
IL1B(0.32)
CXCL10(0.23)
CXCL11(0.26)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
10
20
30
40
50
−5
0
5