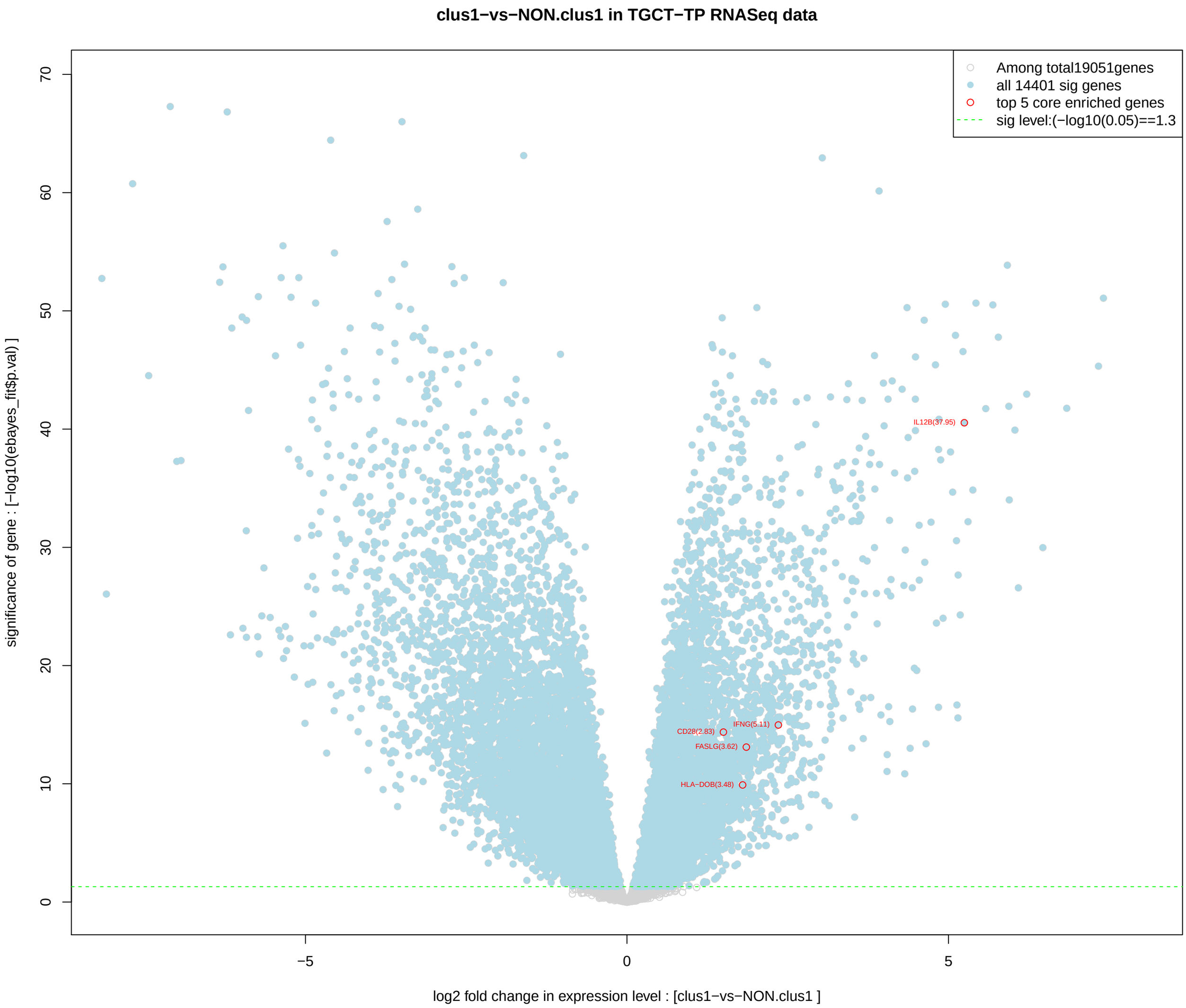
clus1−vs−NON.clus1 in TGCT−TP RNASeq data
Among total19051genes
all 14401 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
IL12B(37.95)
IFNG(5.11)
CD28(2.83)
FASLG(3.62)
HLA−DOB(3.48)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
10
20
30
40
50
60
70
−5
0
5