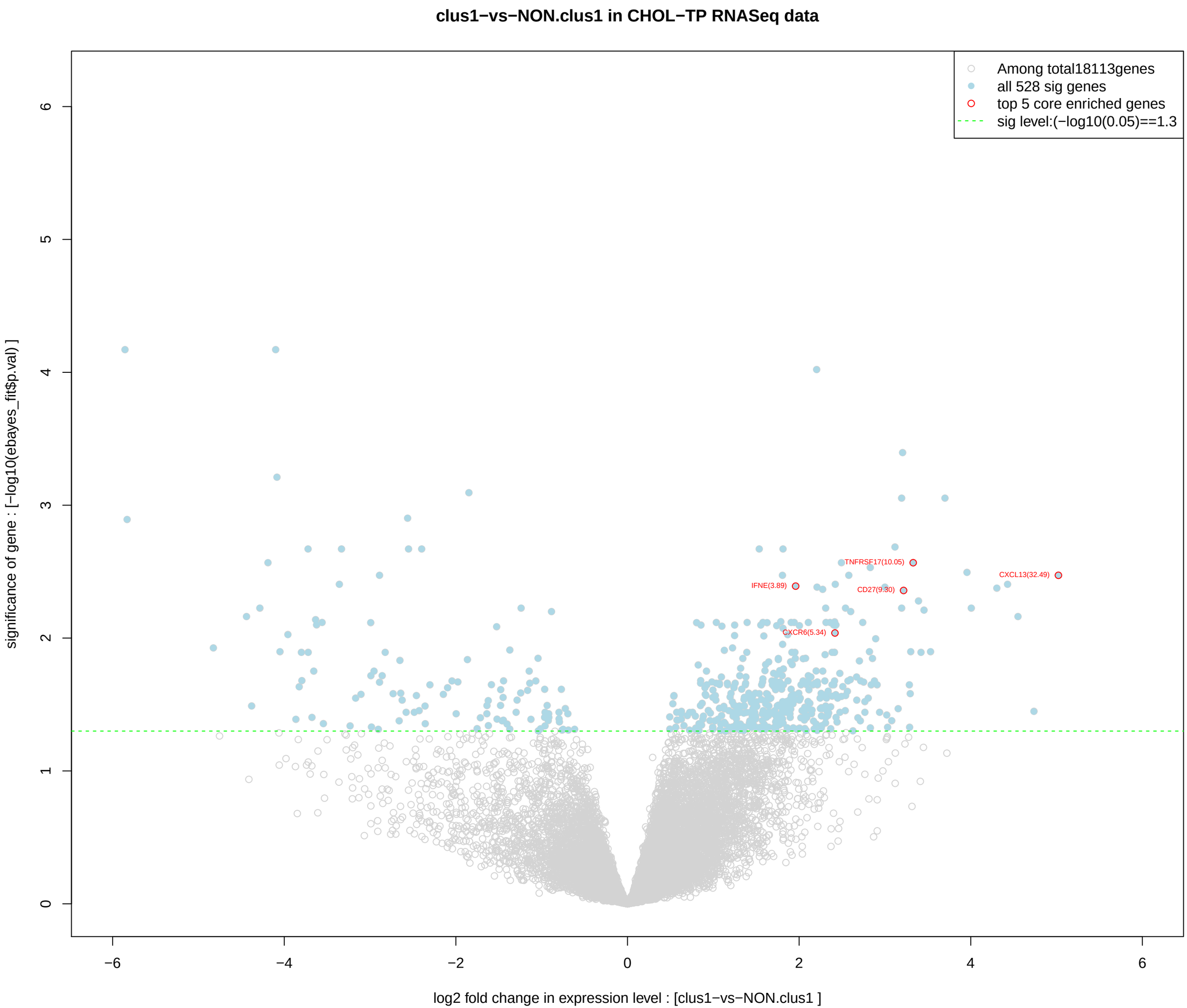
clus1−vs−NON.clus1 in CHOL−TP RNASeq data
Among total18113genes
all 528 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
TNFRSF17(10.05)
CXCL13(32.49)
IFNE(3.89)
CD27(9.30)
CXCR6(5.34)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
1
2
3
4
5
6
−6
−4
−2
0
2
4
6