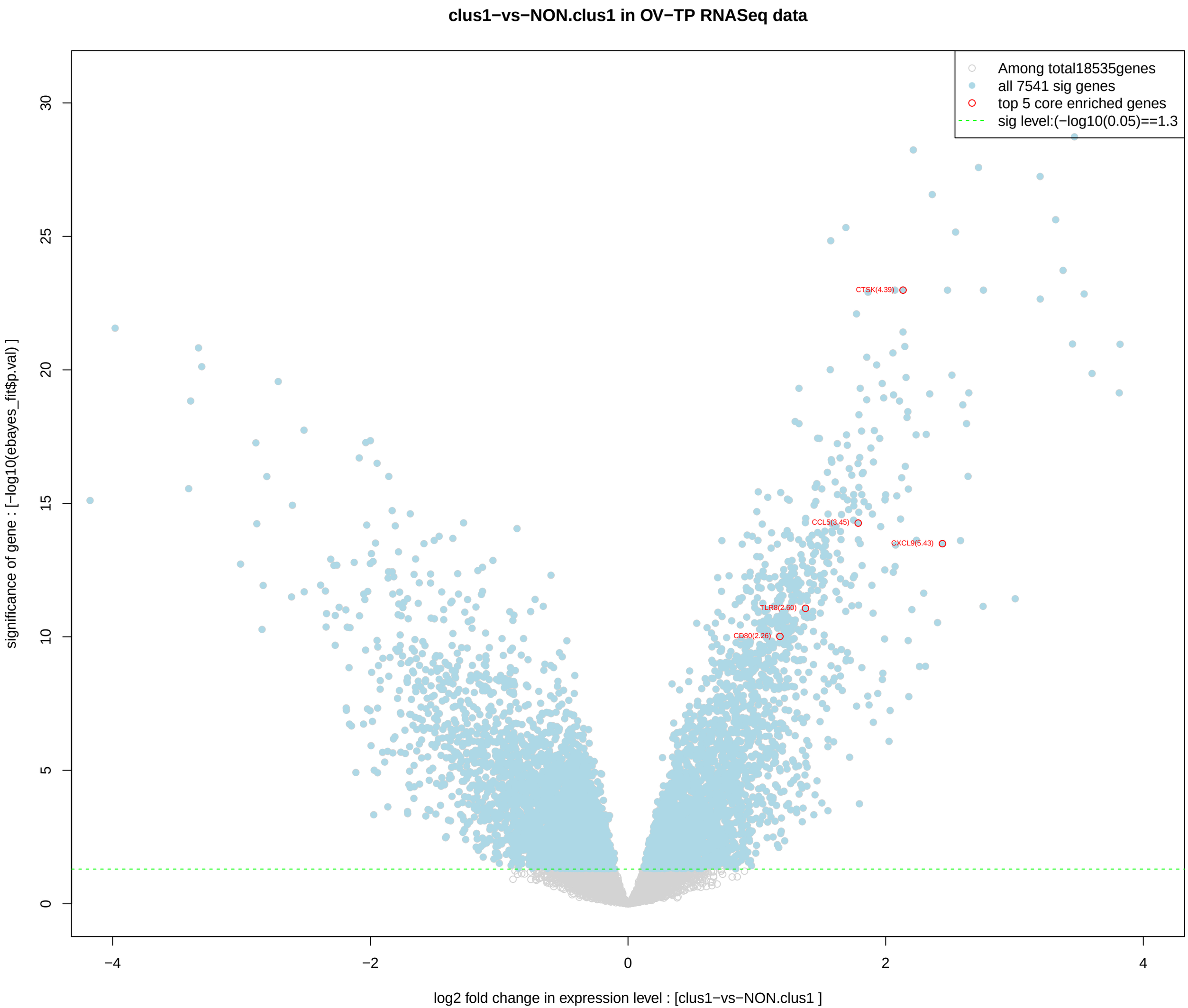
clus1−vs−NON.clus1 in OV−TP RNASeq data
Among total18535genes
all 7541 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CTSK(4.39)
CCL5(3.45)
CXCL9(5.43)
TLR8(2.60)
CD80(2.26)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
5
10
15
20
25
30
−4
−2
0
2
4