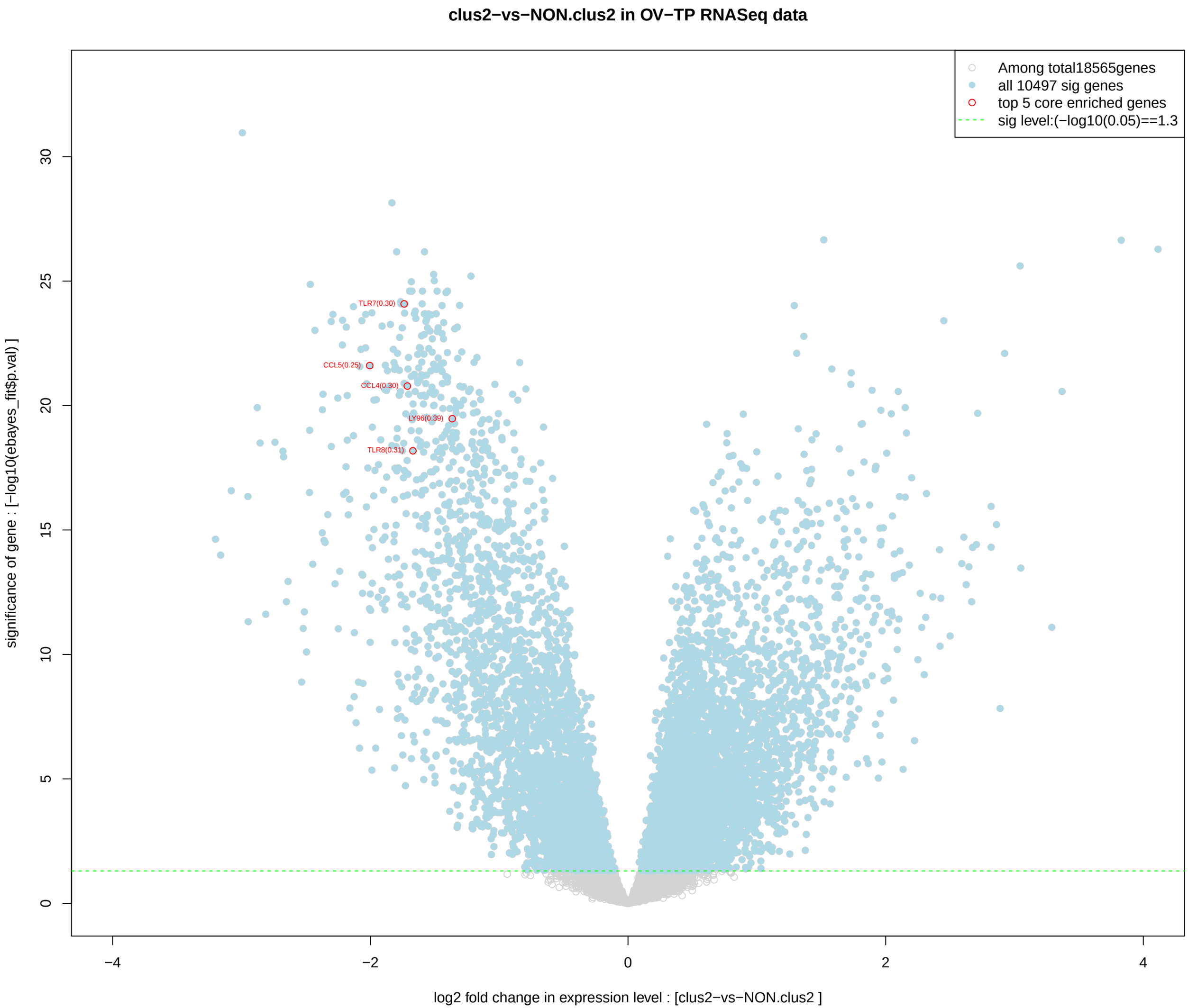
clus2−vs−NON.clus2 in OV−TP RNASeq data
Among total18565genes
all 10497 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
TLR7(0.30)
CCL5(0.25)
CCL4(0.30)
LY96(0.39)
TLR8(0.31)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
5
10
15
20
25
30
−4
−2
0
2
4