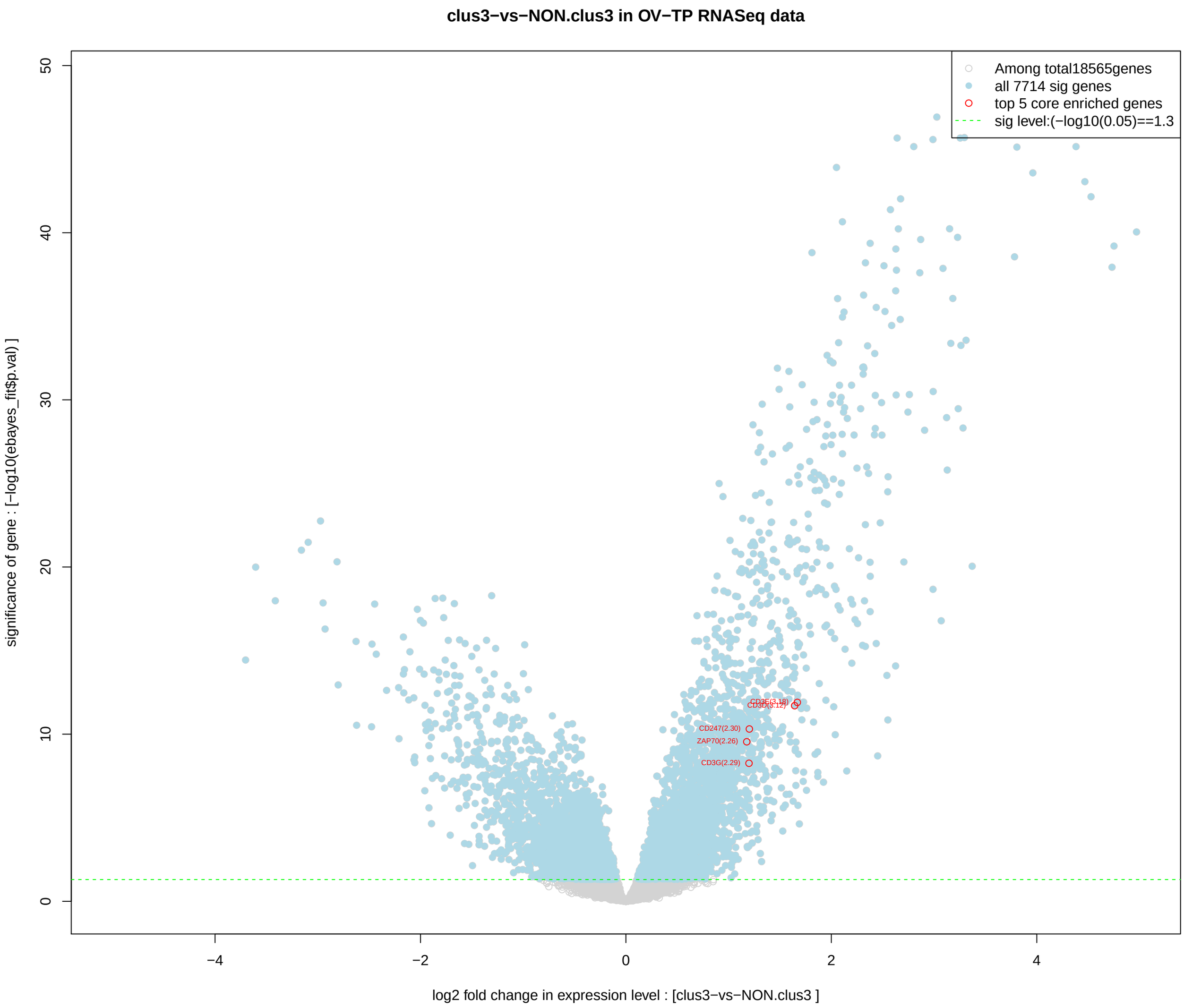
clus3−vs−NON.clus3 in OV−TP RNASeq data
Among total18565genes
all 7714 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CD3E(3.18)
CD3D(3.12)
CD247(2.30)
ZAP70(2.26)
CD3G(2.29)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
10
20
30
40
50
−4
−2
0
2
4