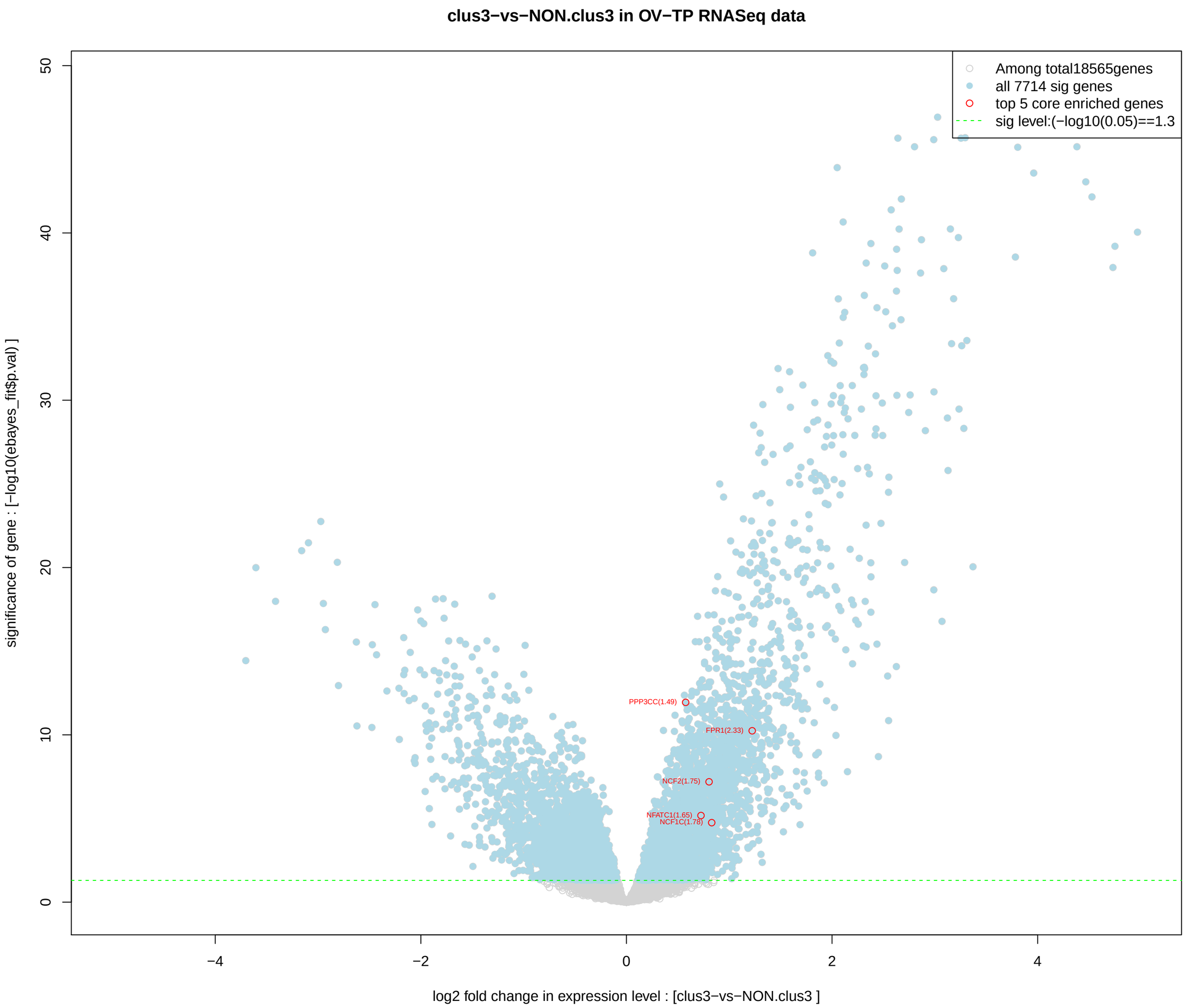
clus3−vs−NON.clus3 in OV−TP RNASeq data
Among total18565genes
all 7714 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PPP3CC(1.49)
FPR1(2.33)
NCF2(1.75)
NFATC1(1.65)
NCF1C(1.78)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
10
20
30
40
50
−4
−2
0
2
4