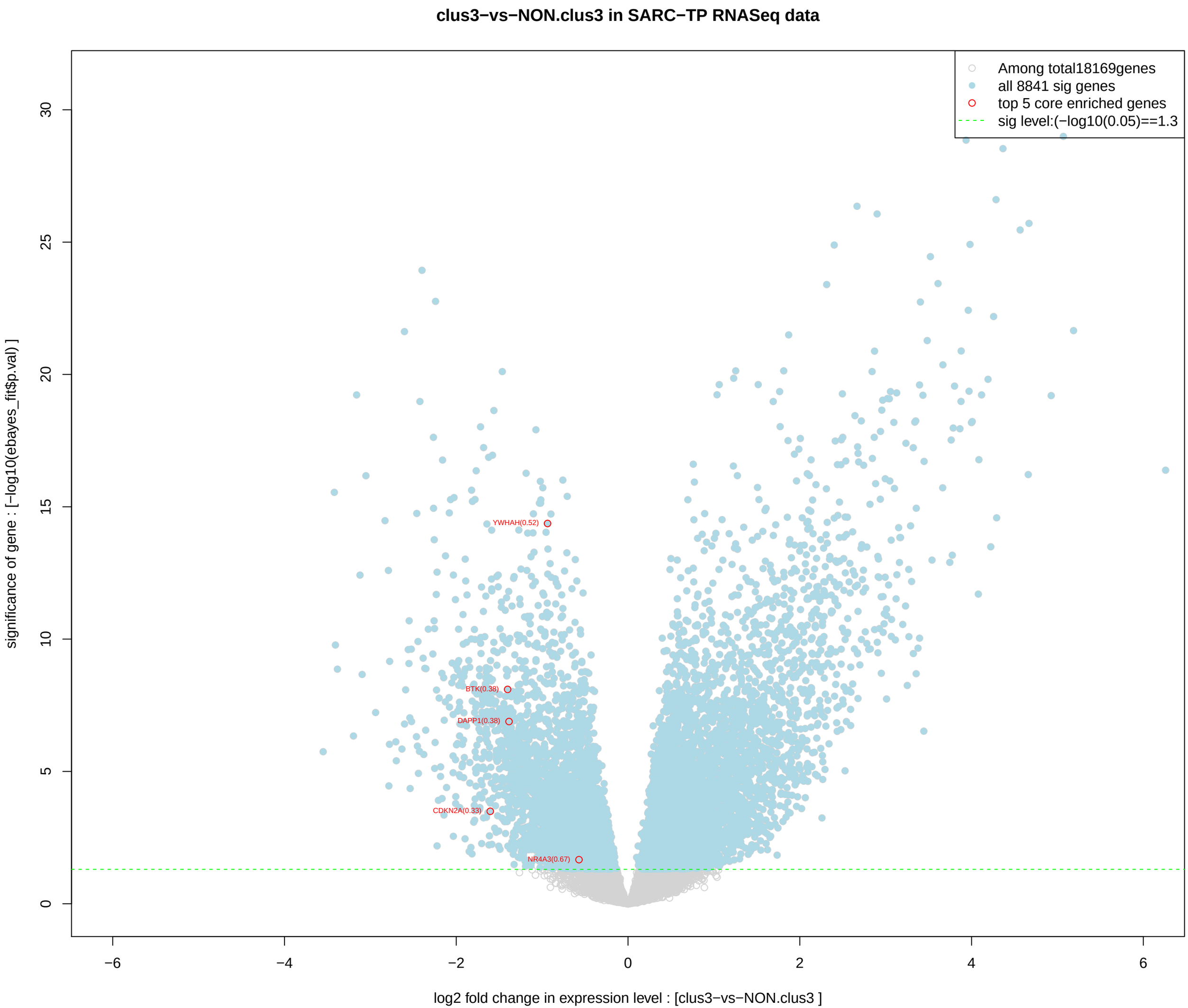
clus3−vs−NON.clus3 in SARC−TP RNASeq data
Among total18169genes
all 8841 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
YWHAH(0.52)
BTK(0.38)
DAPP1(0.38)
CDKN2A(0.33)
NR4A3(0.67)
significance of gene : [−log10(ebayes_fit$p.val) ]
log2 fold change in expression level : [clus3−vs−NON.clus3 ]
0
5
10
15
20
25
30
−6
−4
−2
0
2
4
6