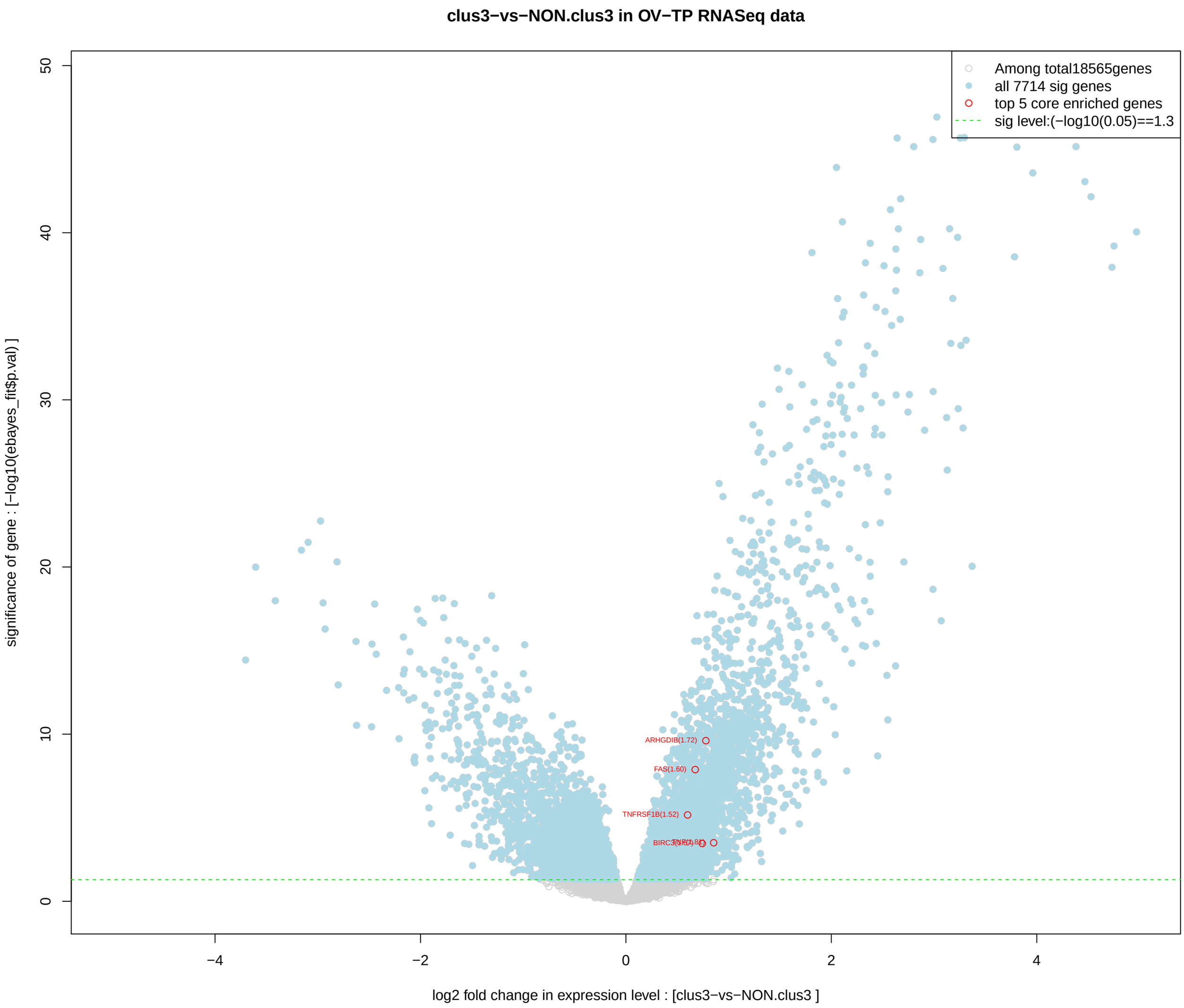
clus3−vs−NON.clus3 in OV−TP RNASeq data
Among total18565genes
all 7714 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ARHGDIB(1.72)
FAS(1.60)
TNFRSF1B(1.52)
BIRC3(1.60)
TNFR(1.81)
significance of gene : [−log10(ebayes_fit$p.val) ]
log2 fold change in expression level : [clus3−vs−NON.clus3 ]
0
10
20
30
40
50
−4
−2
0
2
4