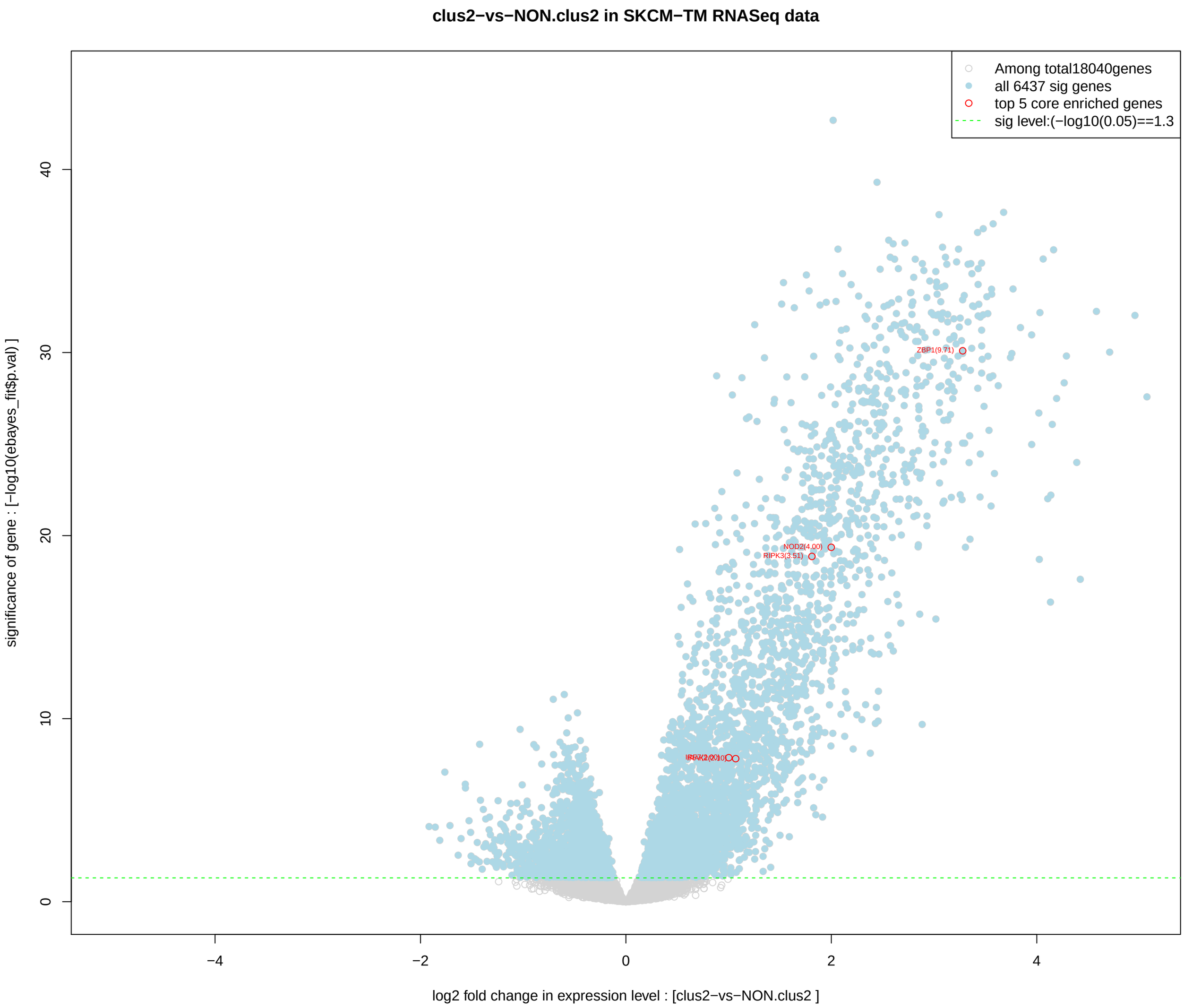
clus2−vs−NON.clus2 in SKCM−TM RNASeq data
Among total18040genes
all 6437 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ZBP1(9.71)
NOD2(4.00)
RIPK3(3.51)
IRF7(2.00)
IRAK2(2.10)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
10
20
30
40
−4
−2
0
2
4