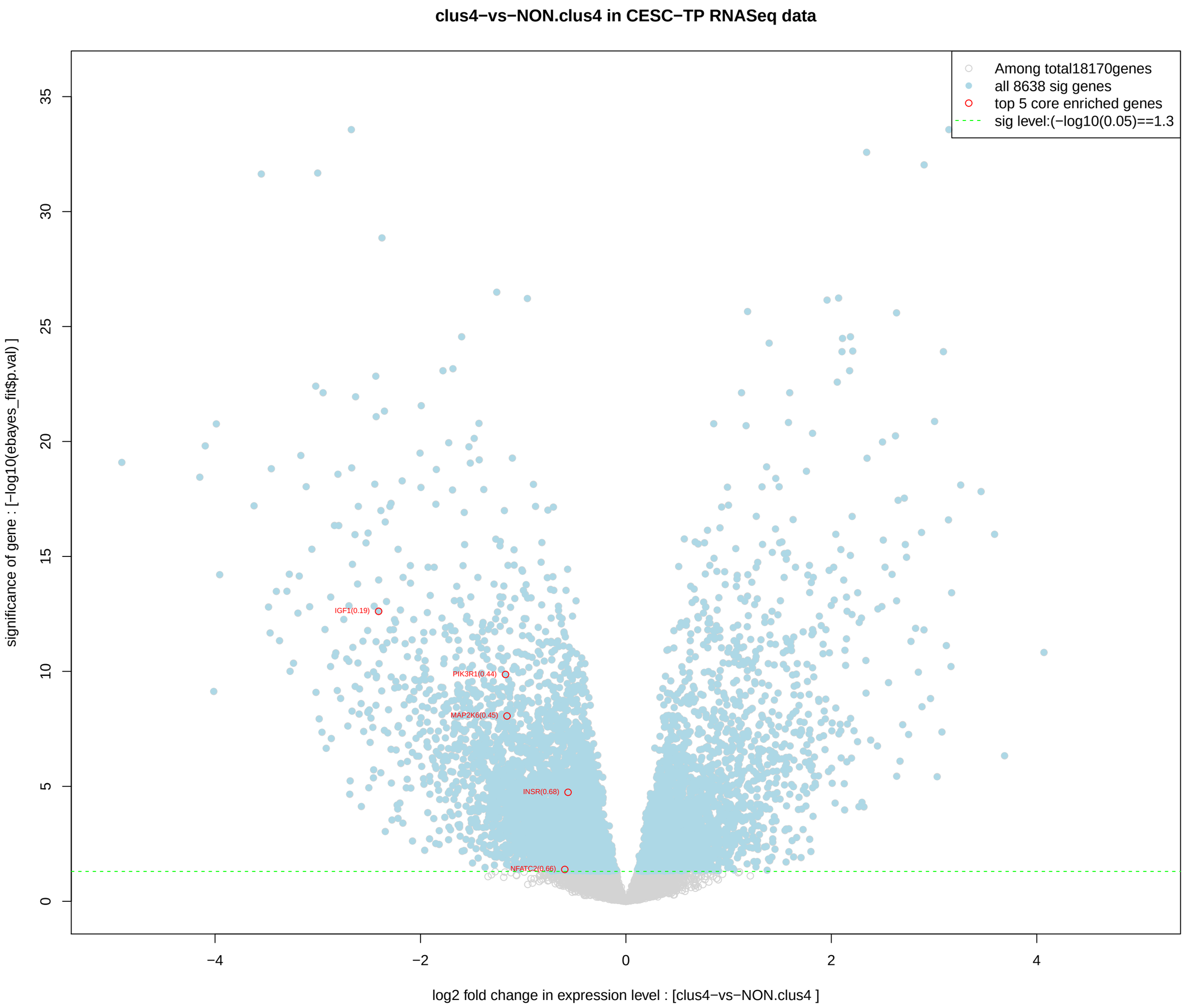
clus4−vs−NON.clus4 in CESC−TP RNASeq data
Among total18170genes
all 8638 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
IGF1(0.19)
PIK3R1(0.44)
MAP2K6(0.45)
INSR(0.68)
NFATC2(0.66)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0
5
10
15
20
25
30
35
−4
−2
0
2
4