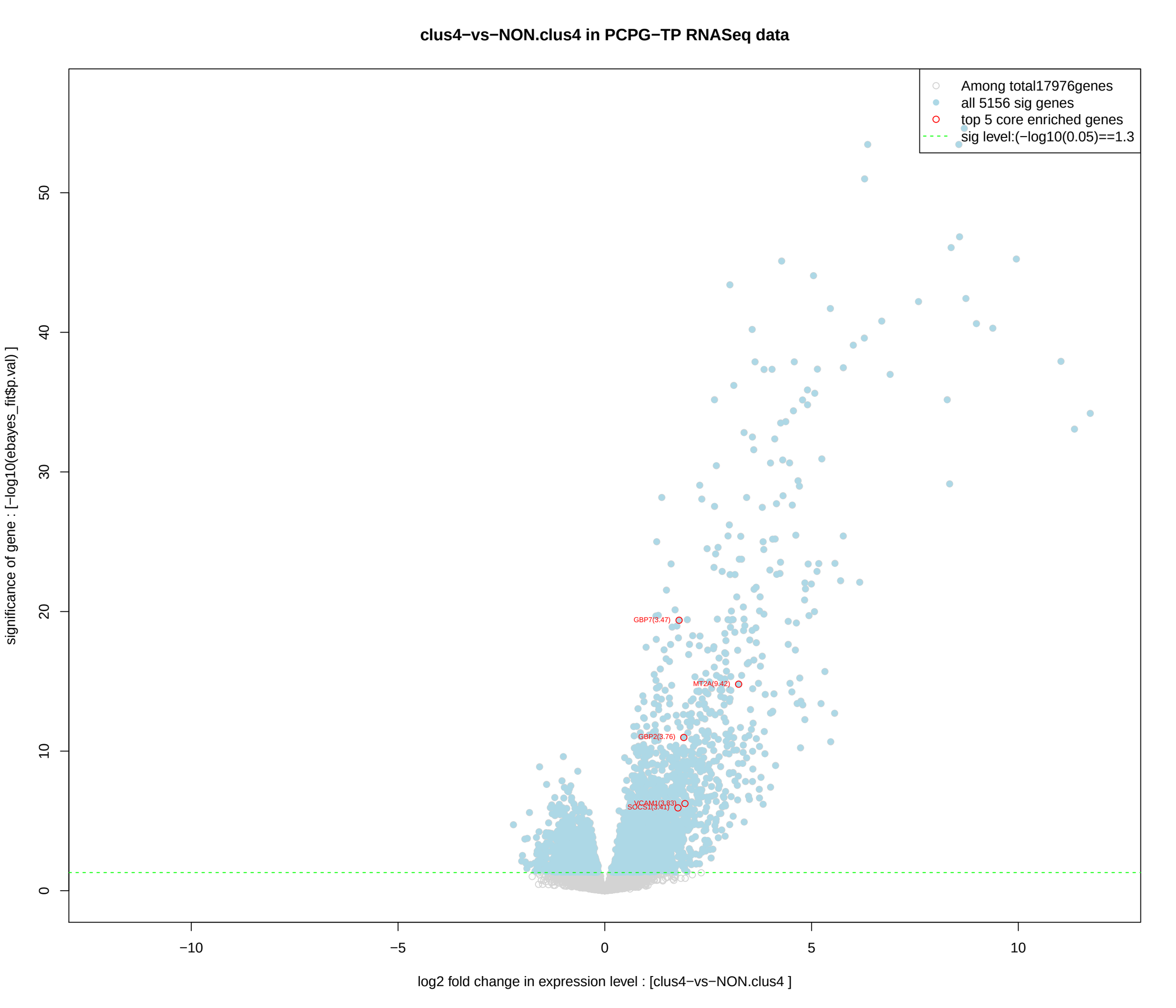
clus4−vs−NON.clus4 in PCPG−TP RNASeq data
Among total17976genes
all 5156 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
GBP7(3.47)
MT2A(9.42)
GBP2(3.76)
VCAM1(3.83)
SOCS1(3.41)
significance of gene : [−log10(ebayes_fit$p.val)]
0
10
20
30
40
50
−10
−5
0
5
10
log2 fold change in expression level : [clus4−vs−NON.clus4]