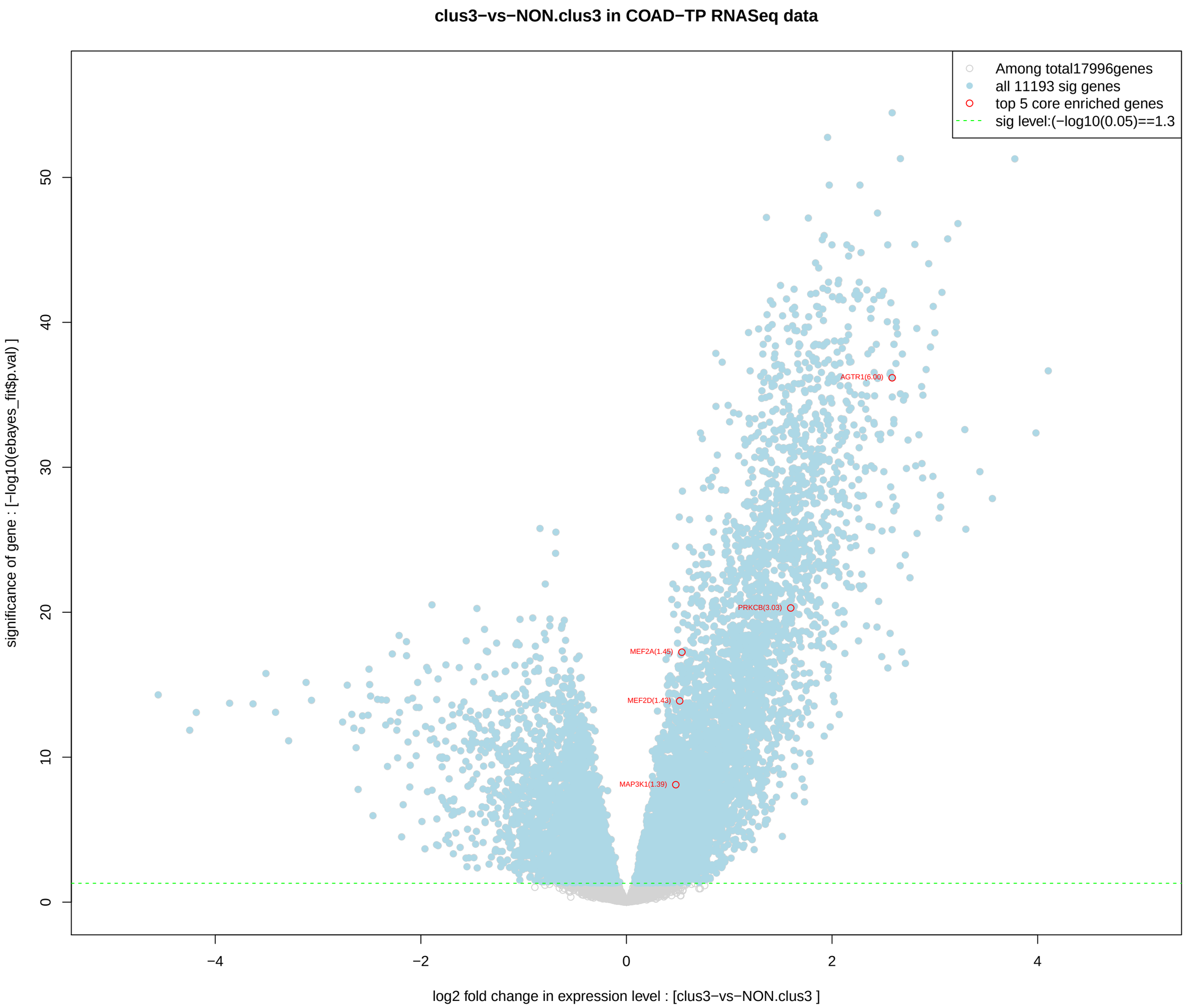
clus3−vs−NON.clus3 in COAD−TP RNASeq data
Among total17996genes
all 11193 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
AGTR1(6.00)
PRKCB(3.03)
MEF2A(1.45)
MEF2D(1.43)
MAP3K1(1.39)
significance of gene : [−log10(ebayes_fit$p.val)]
0
10
20
30
40
50
−4
−2
0
2
4
log2 fold change in expression level : [clus3−vs−NON.clus3]