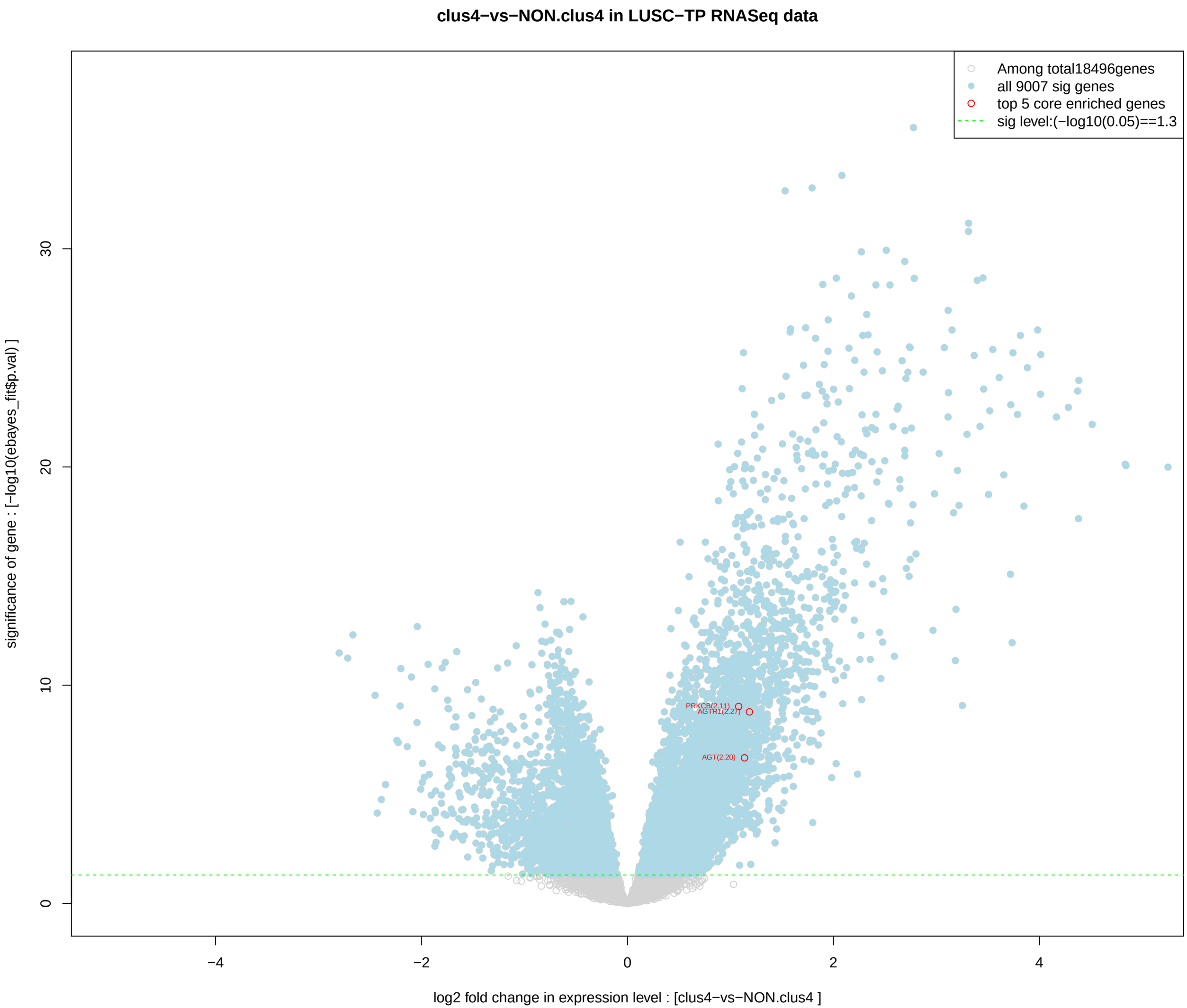
clus4−vs−NON.clus4 in LUSC−TP RNASeq data
Among total18496genes
all 9007 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PRKCB(2.11)
AGTR1(2.27)
AGT(2.20)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0
10
20
30
−4
−2
0
2
4