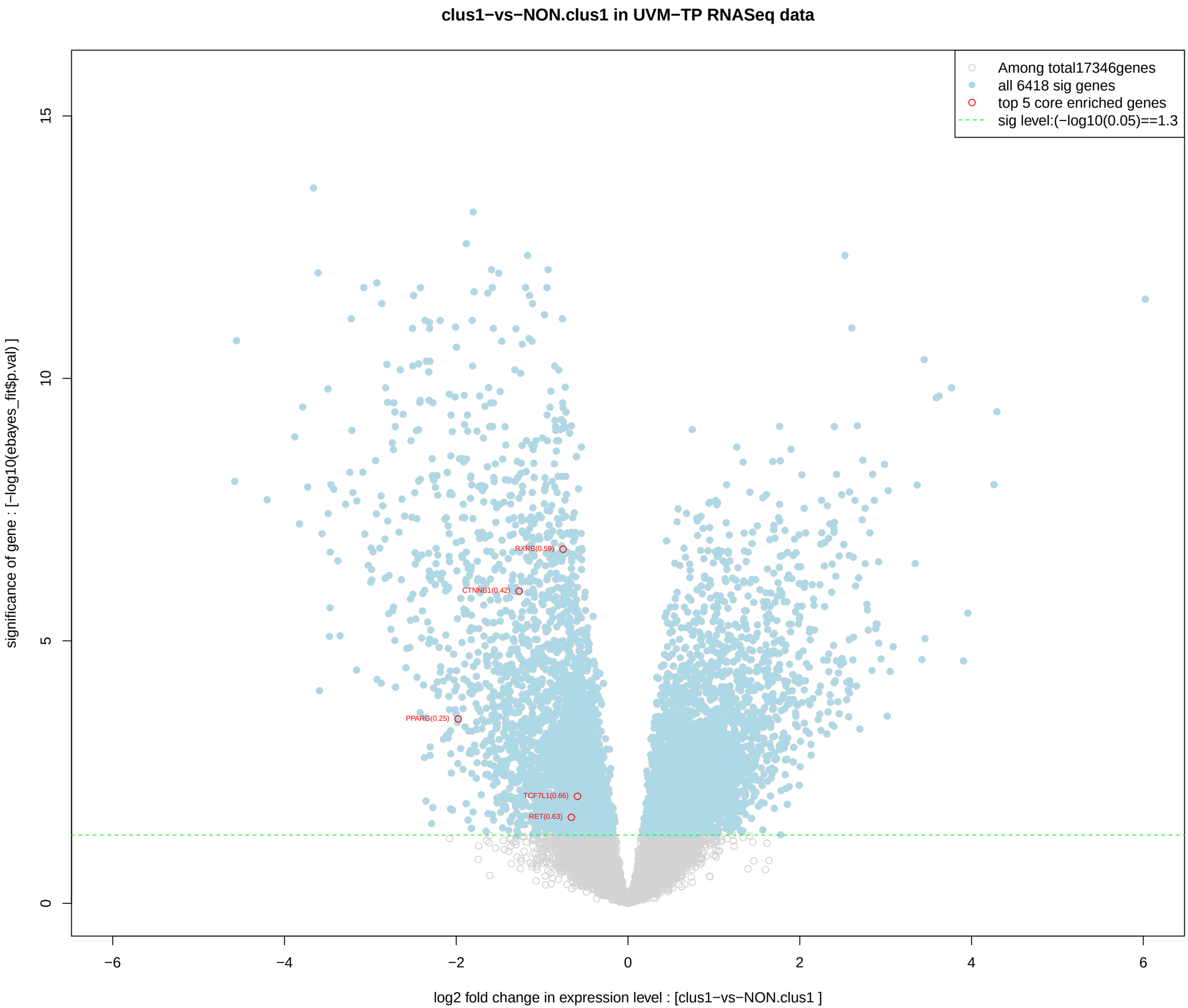
clus1−vs−NON.clus1 in UVM−TP RNASeq data
Among total17346genes
all 6418 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
RXRB(0.59)
CTNNB1(0.42)
PPARG(0.25)
TCF7L1(0.66)
RET(0.63)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
5
10
15
−6
−4
−2
0
2
4
6