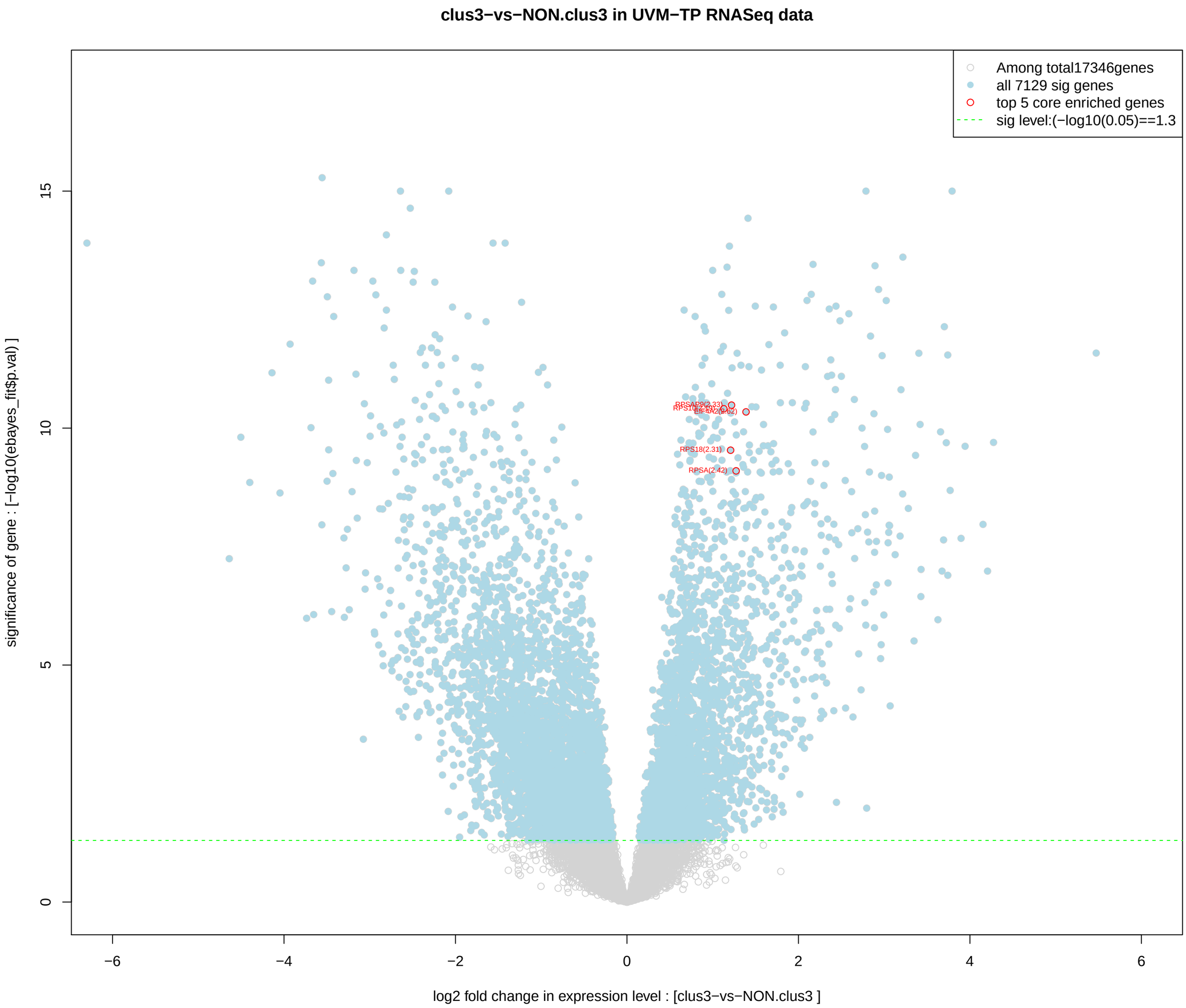
clus3−vs−NON.clus3 in UVM−TP RNASeq data
Among total17346genes
all 7129 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
RPSA(2.42)
RPS18(2.31)
Eif4A2(2.62)
RPS10(2.33)
RPSAP9(2.33)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
5
10
15
−6
−4
−2
0
2
4
6