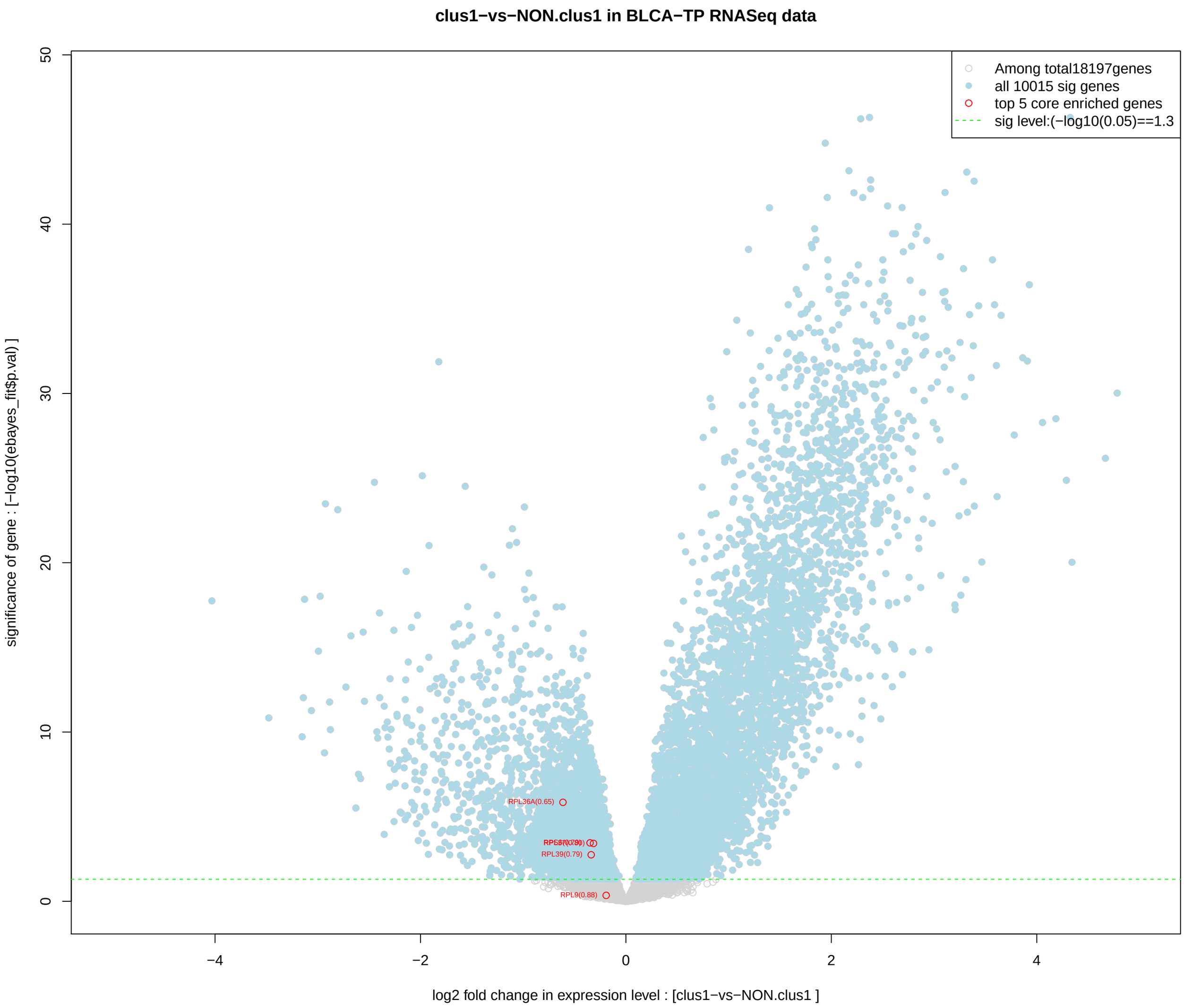
clus1−vs−NON.clus1 in BLCA−TP RNASeq data
Among total18197genes
all 10015 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
RPL36A(0.65)
RPL39(0.79)
RPL9(0.88)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
10
20
30
40
50
−4
−2
0
2
4