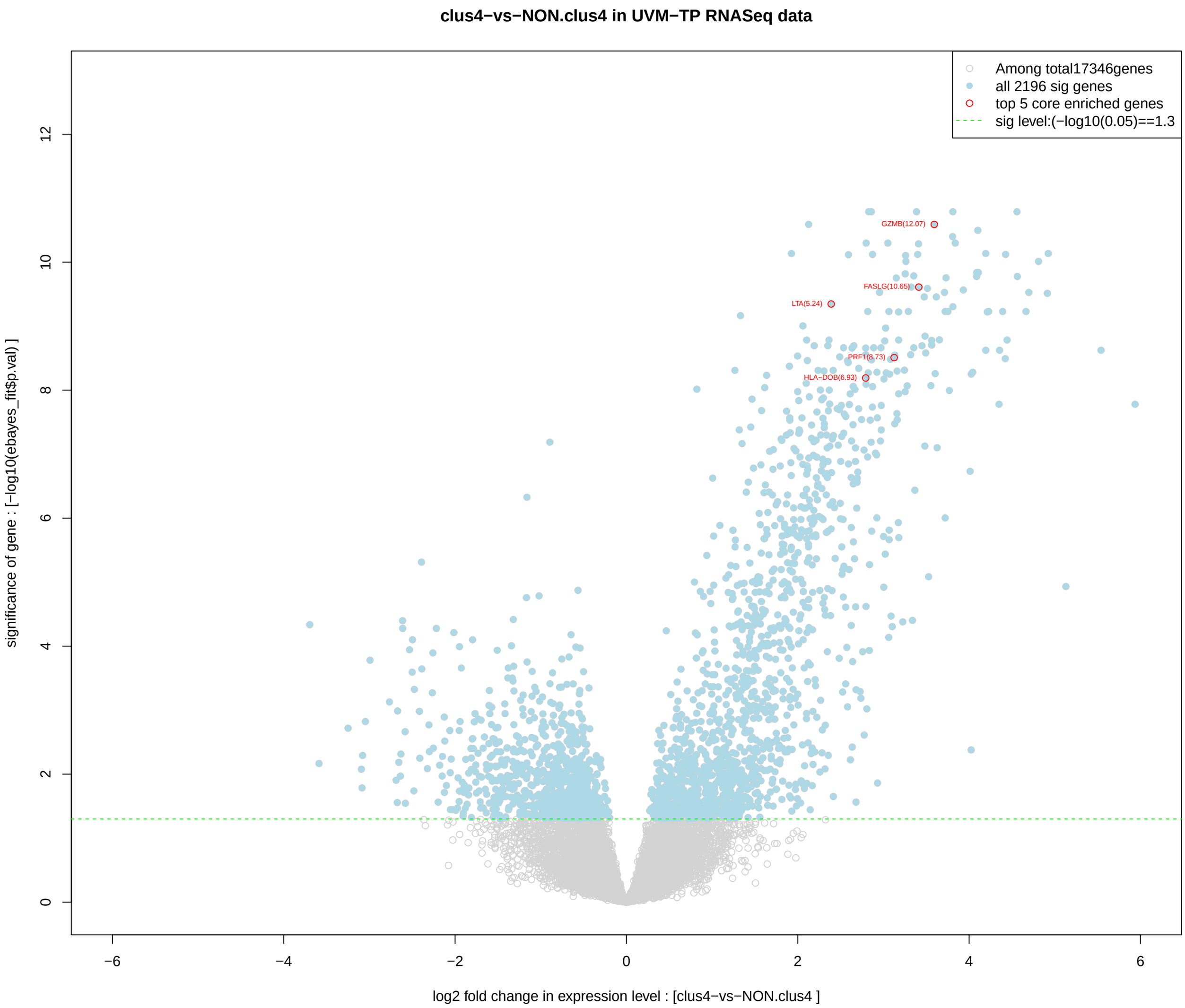
clus4−vs−NON.clus4 in UVM−TP RNASeq data
Among total17346genes
all 2196 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
GZMB(12.07)
FASLG(10.65)
LTA(5.24)
PRF1(8.73)
HLA−DOB(6.93)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0
2
4
6
8
10
12
−6
−4
−2
0
2
4
6