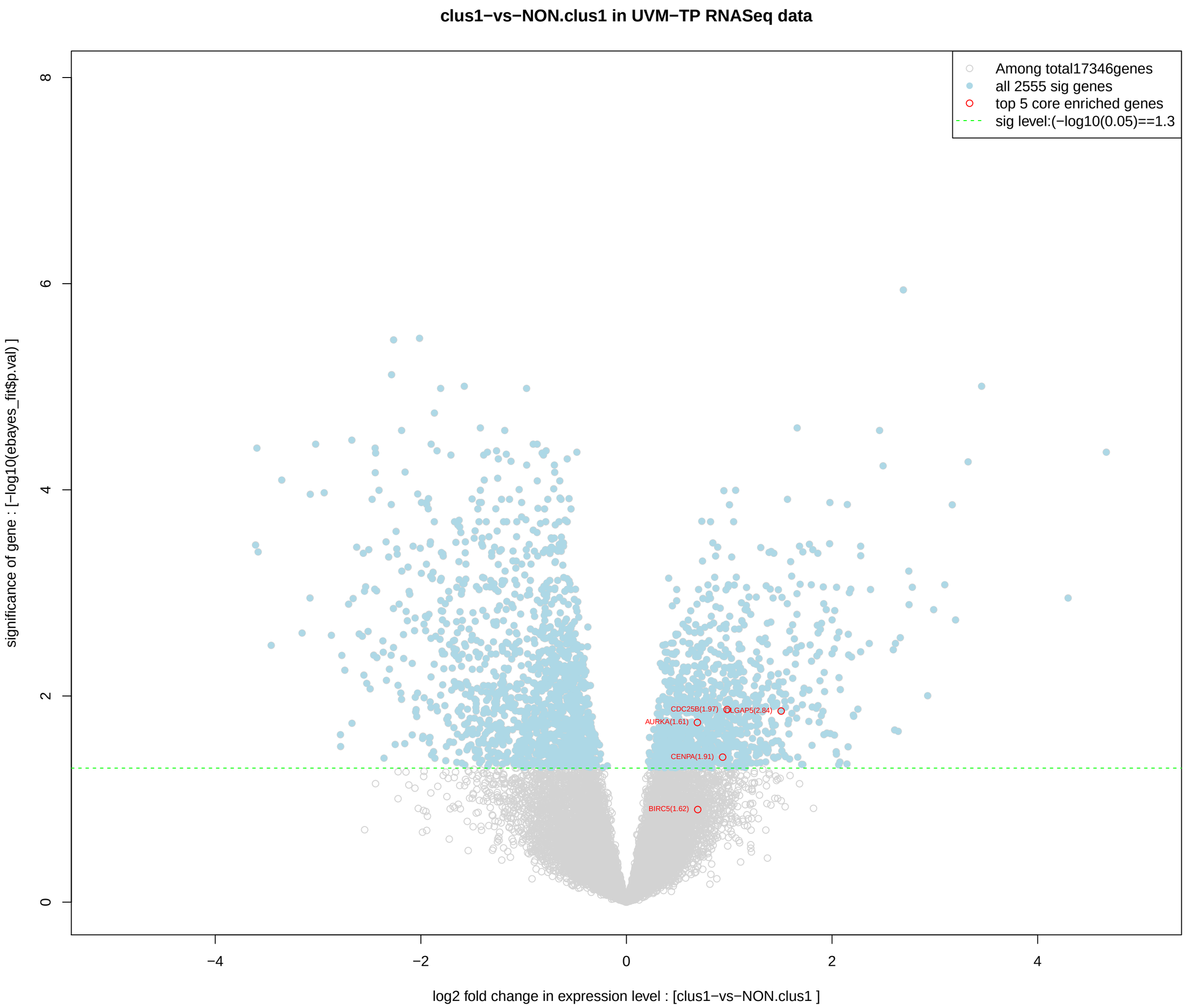
clus1−vs−NON.clus1 in UVM−TP RNASeq data
Among total17346genes
all 2555 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CDC25B(1.97)
DLGAP5(2.84)
AURKA(1.61)
CENPA(1.91)
BIRC5(1.62)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
2
4
6
8
−4
−2
0
2
4