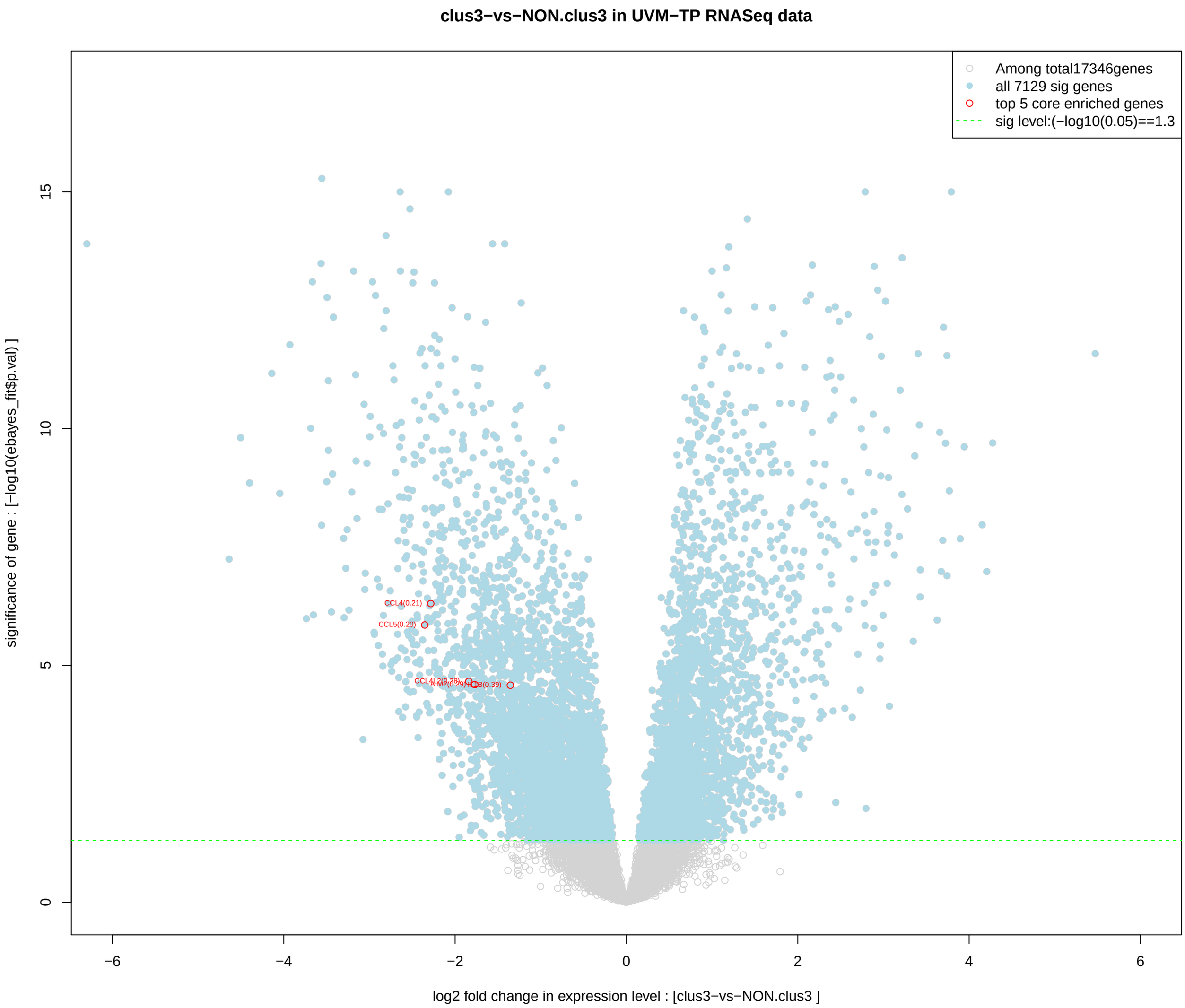
clus3−vs−NON.clus3 in UVM−TP RNASeq data
Among total17346genes
all 7129 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CCL4(0.21)
CCL5(0.20)
CCL4L2(0.28)
AIM2(0.29)
HCB(0.39)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
5
10
15
−6
−4
−2
0
2
4
6