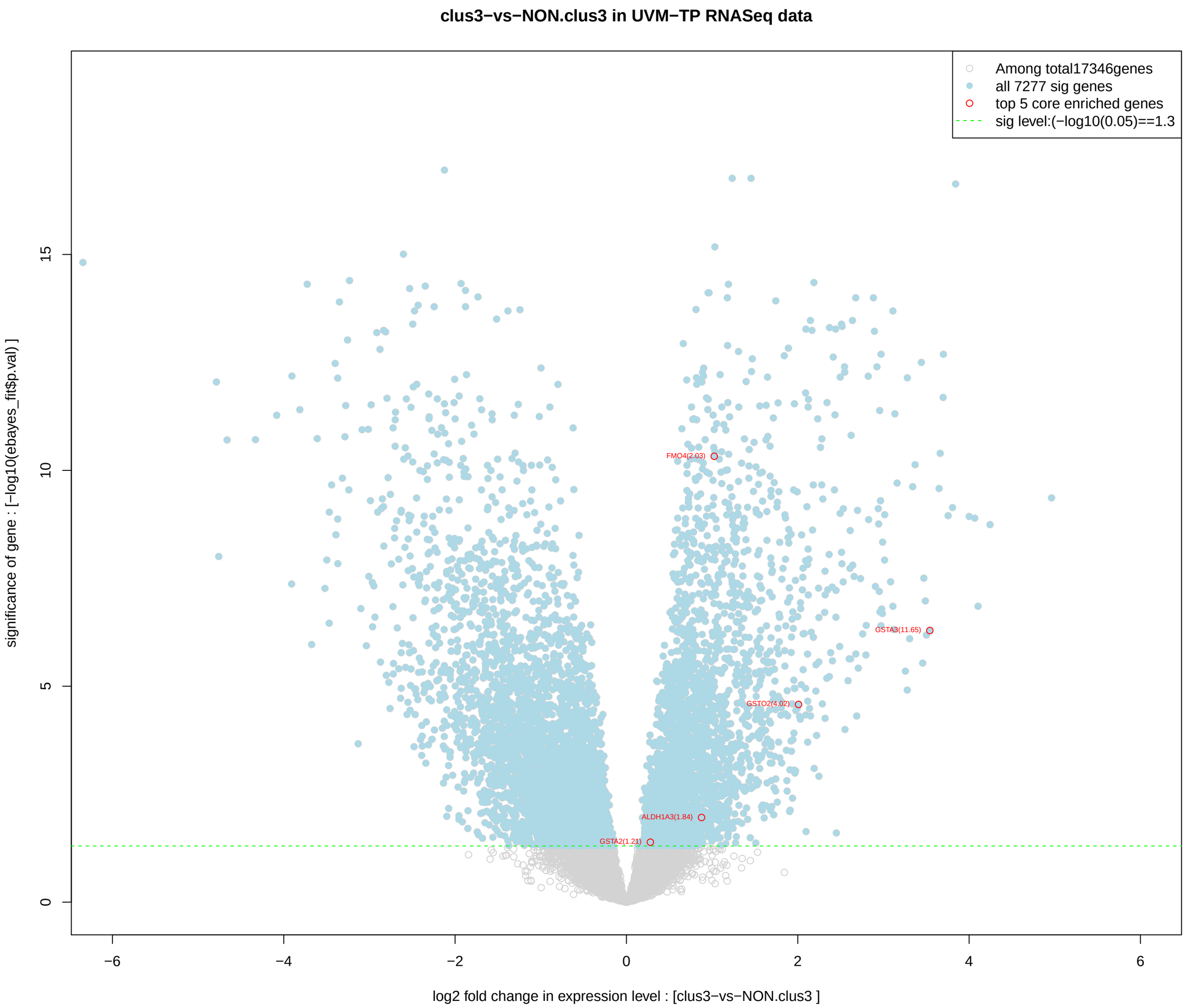
clus3−vs−NON.clus3 in UVM−TP RNASeq data
Among total17346genes
all 7277 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
FMO4(2.03)
GSTA3(11.65)
GSTO2(4.02)
ALDH1A3(1.84)
GSTA2(1.21)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
5
10
15
−6
−4
−2
0
2
4
6