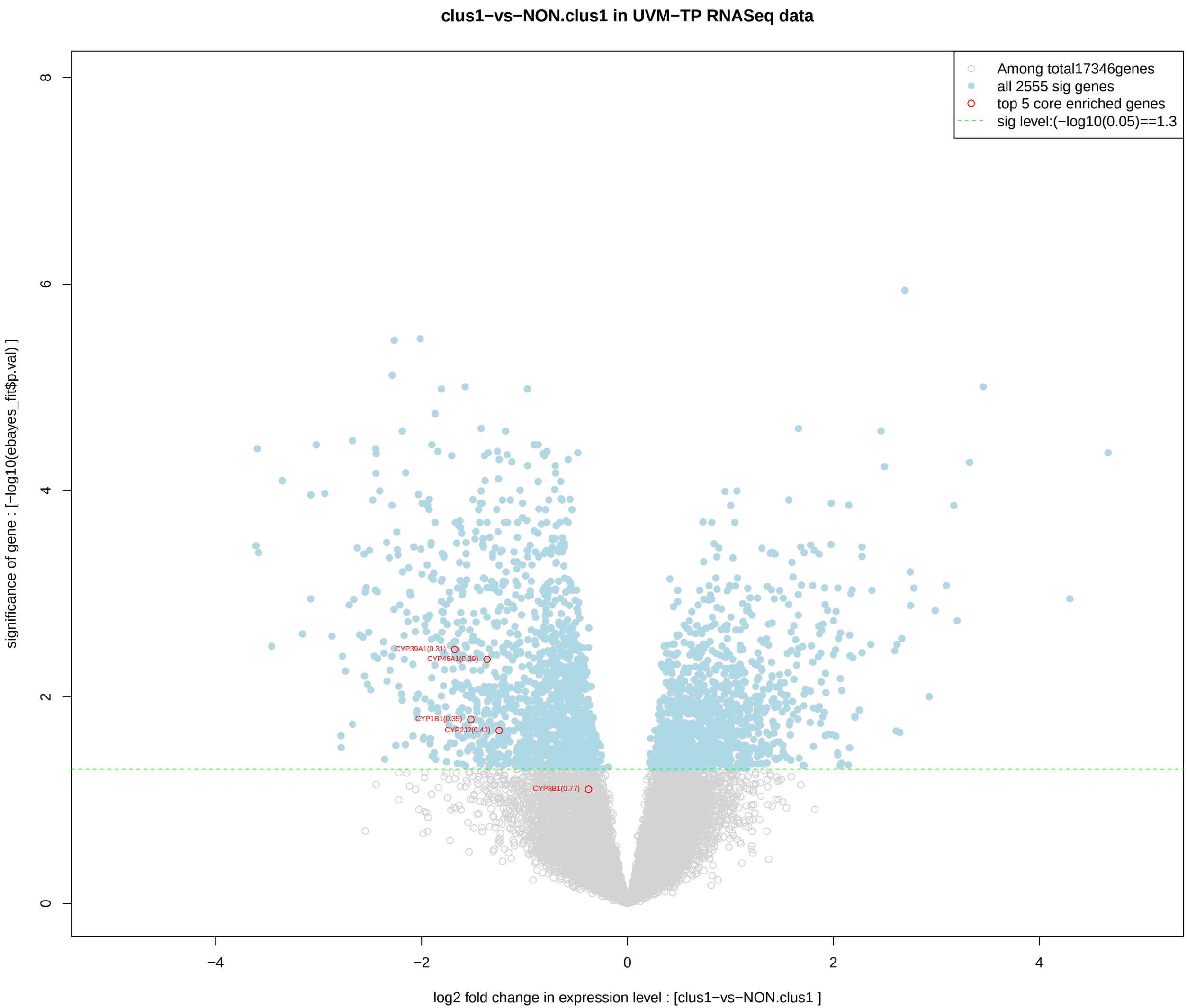
clus1−vs−NON.clus1 in UVM−TP RNASeq data
Among total17346genes
all 2555 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CYP39A1(0.31)
CYP46A1(0.39)
CYP1B1(0.35)
CYP2J2(0.42)
CYP8B1(0.77)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
2
4
6
8
−4
−2
0
2
4