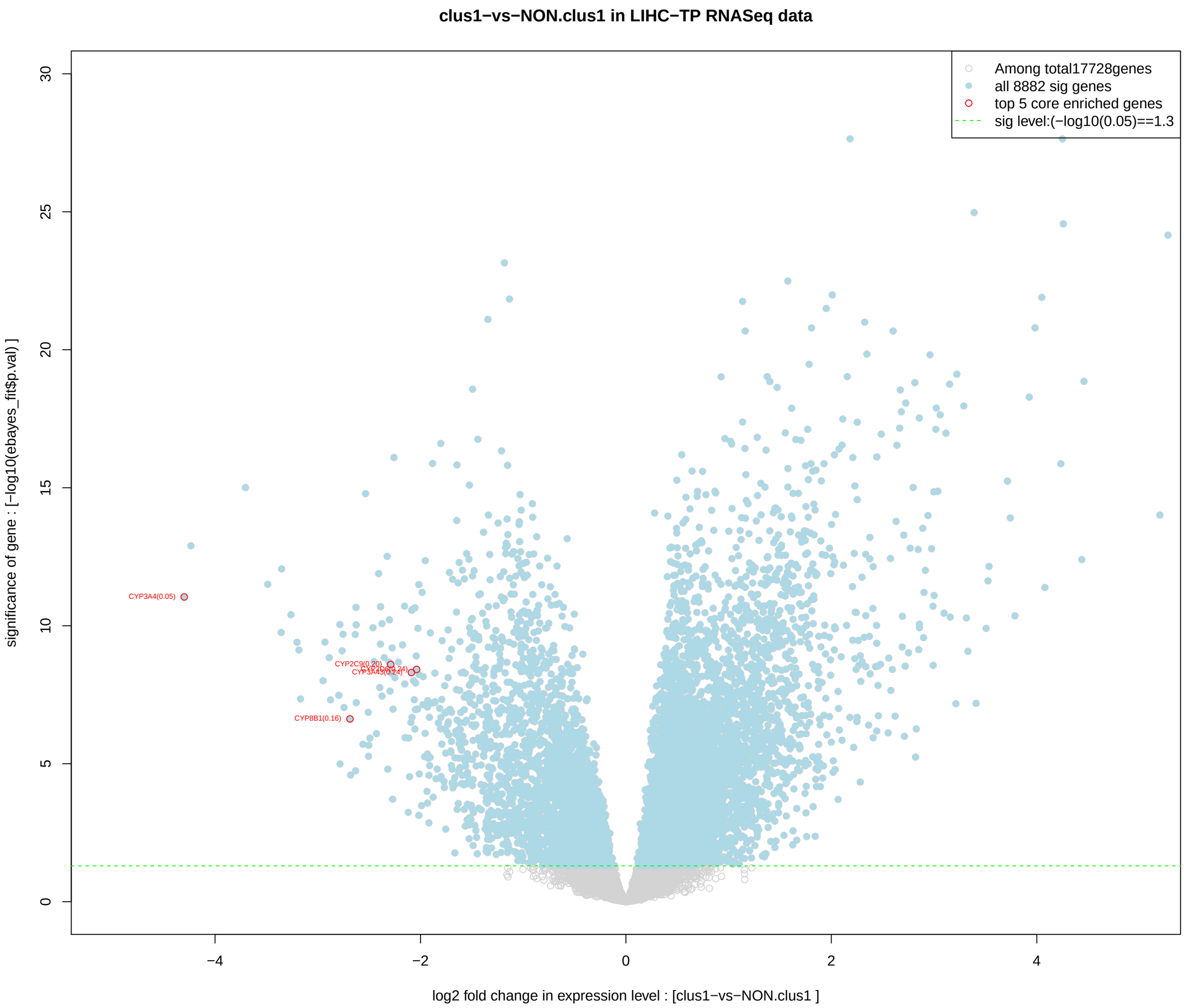
clus1−vs−NON.clus1 in LIHC−TP RNASeq data
Among total17728genes
all 8882 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CYP3A4(0.05)
CYP2C9(0.20)
CYP2D6(0.24)
CYP3A43(0.24)
CYP8B1(0.16)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
5
10
15
20
25
30
−4
−2
0
2
4