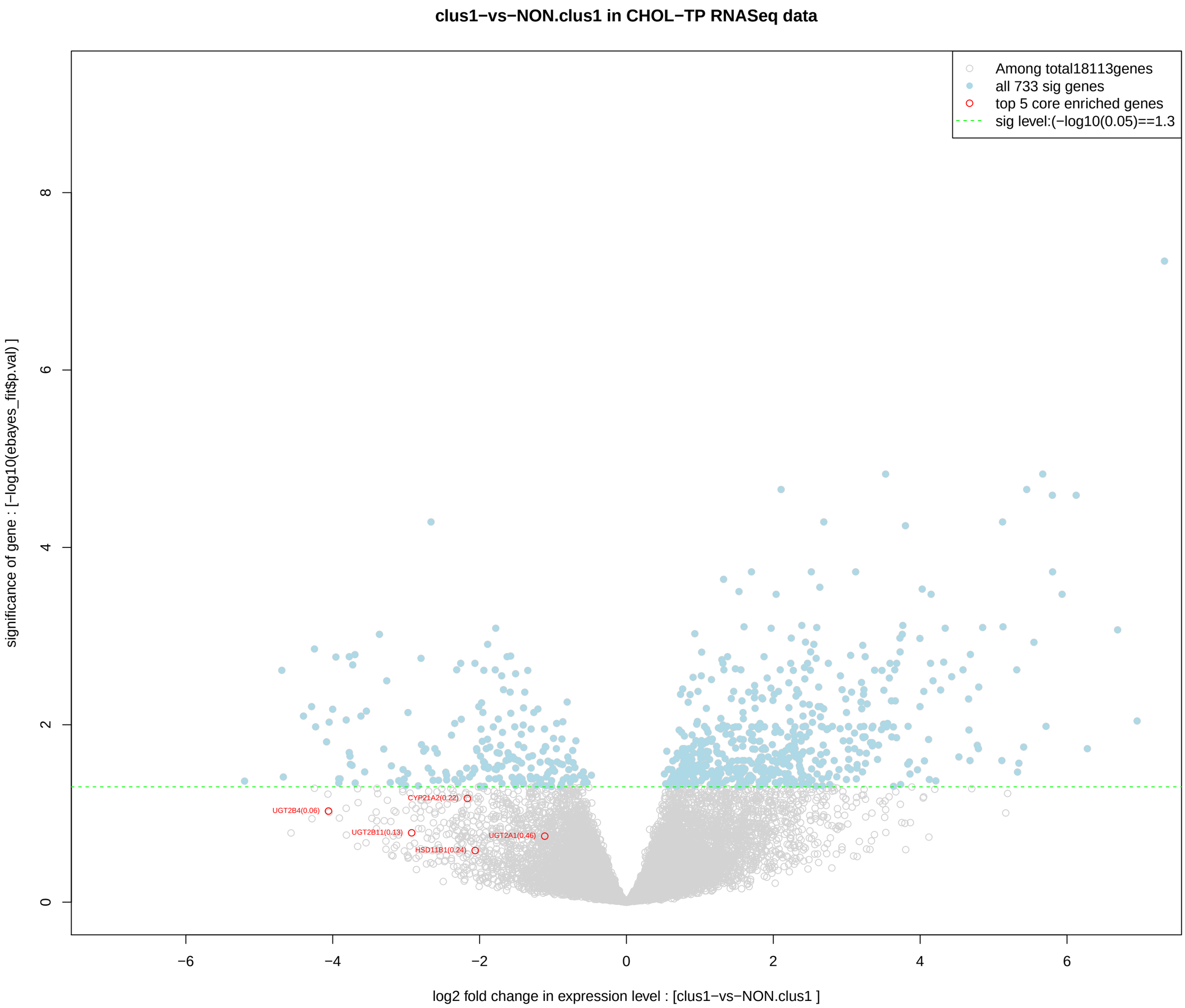
clus1−vs−NON.clus1 in CHOL−TP RNASeq data
Among total18113genes
all 733 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CYP21A2(0.22)
UGT2B4(0.06)
UGT2B11(0.13)
UGT2A1(0.46)
HSD11B1(0.24)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
2
4
6
8
−6
−4
−2
0
2
4
6