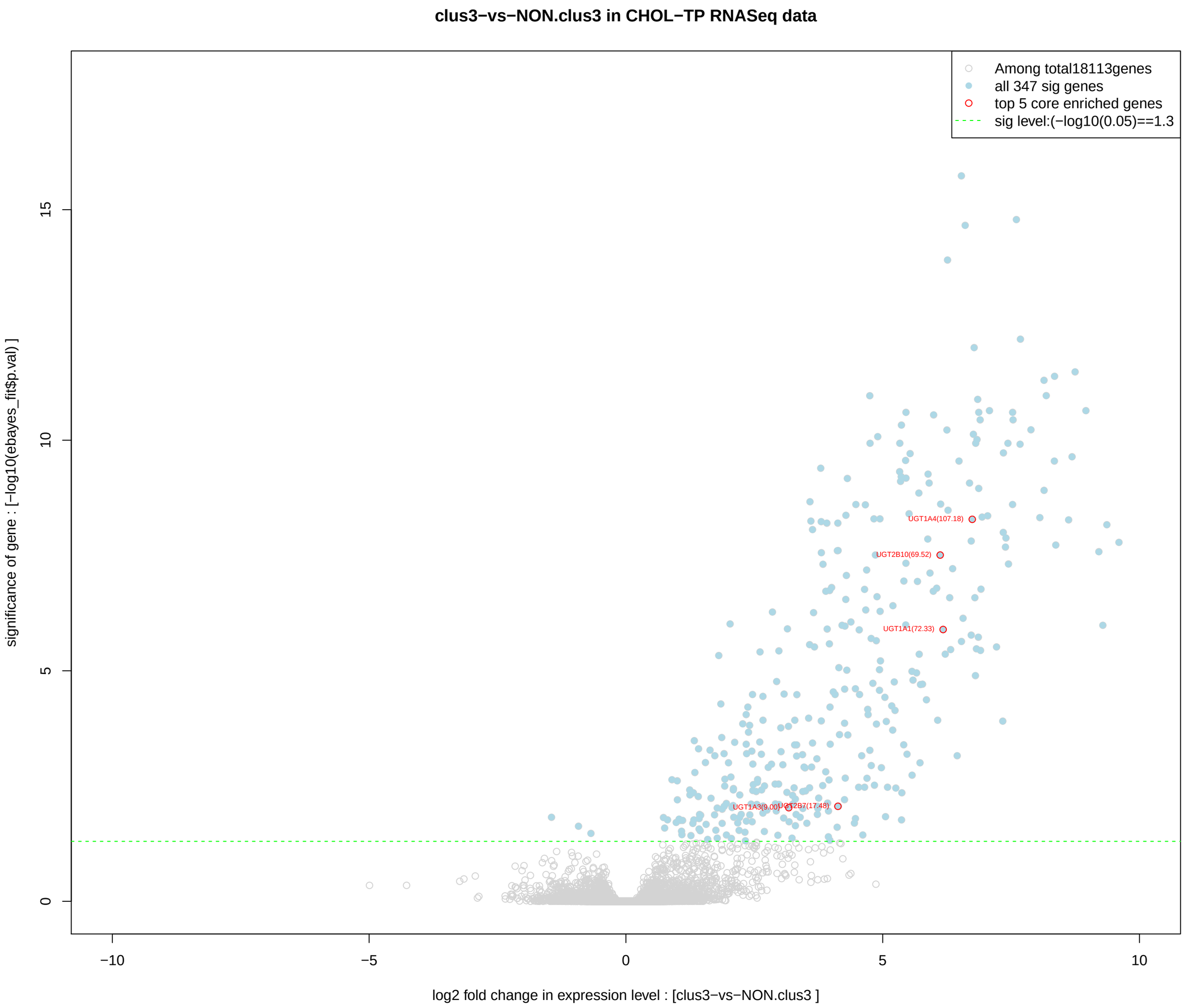
clus3−vs−NON.clus3 in CHOL−TP RNASeq data
Among total18113genes
all 347 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
UGT1A4(107.18)
UGT2B10(69.52)
UGT1A1(72.33)
UGT1A3(9.00)
UGT2B7(17.48)
significance of gene : [−log10(ebayes_fit$p.val) ]
log2 fold change in expression level : [clus3−vs−NON.clus3 ]
0
5
10
15
−10
−5
0
5
10