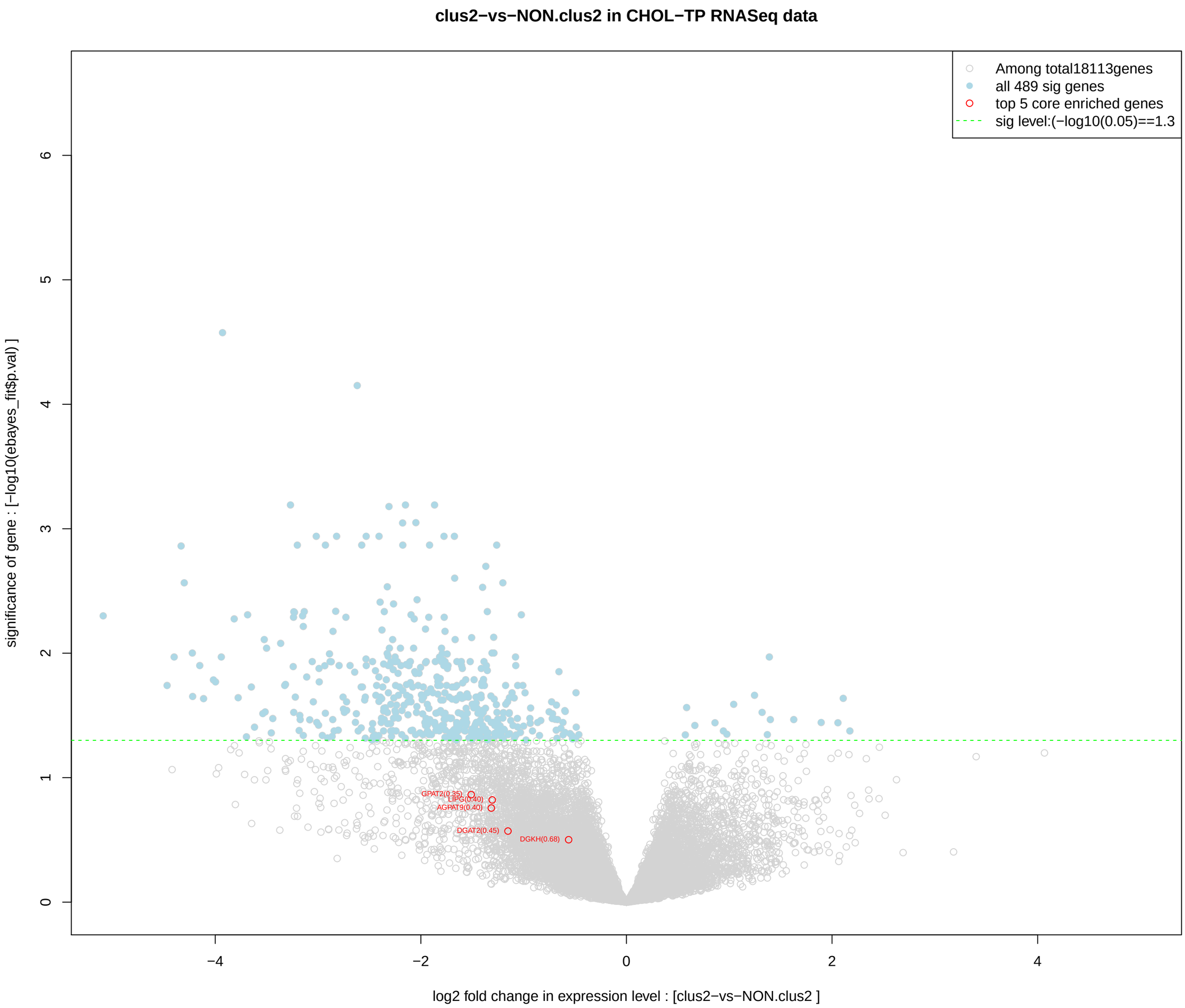
clus2−vs−NON.clus2 in CHOL−TP RNASeq data
Among total18113genes
all 489 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
GPAT2(0.35)
LIPG(0.40)
AGPAT9(0.40)
DGAT2(0.45)
DGKH(0.68)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
1
2
3
4
5
6
−4
−2
0
2
4