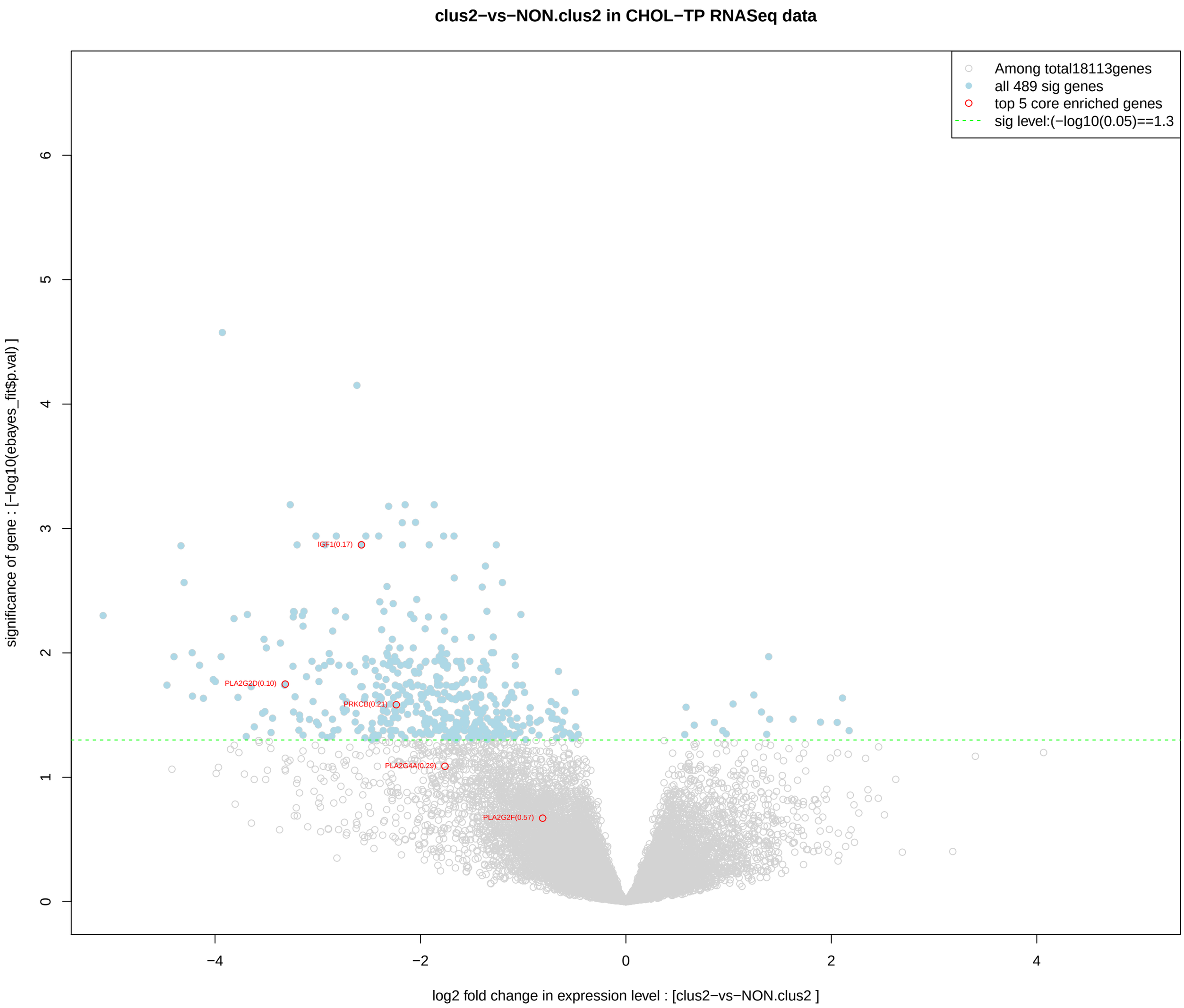
clus2−vs−NON.clus2 in CHOL−TP RNASeq data
Among total18113genes
all 489 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
IGF1(0.17)
PLA2G2D(0.10)
PRKCB(0.21)
PLA2G4A(0.29)
PLA2G2F(0.57)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
1
2
3
4
5
6
−4
−2
0
2
4