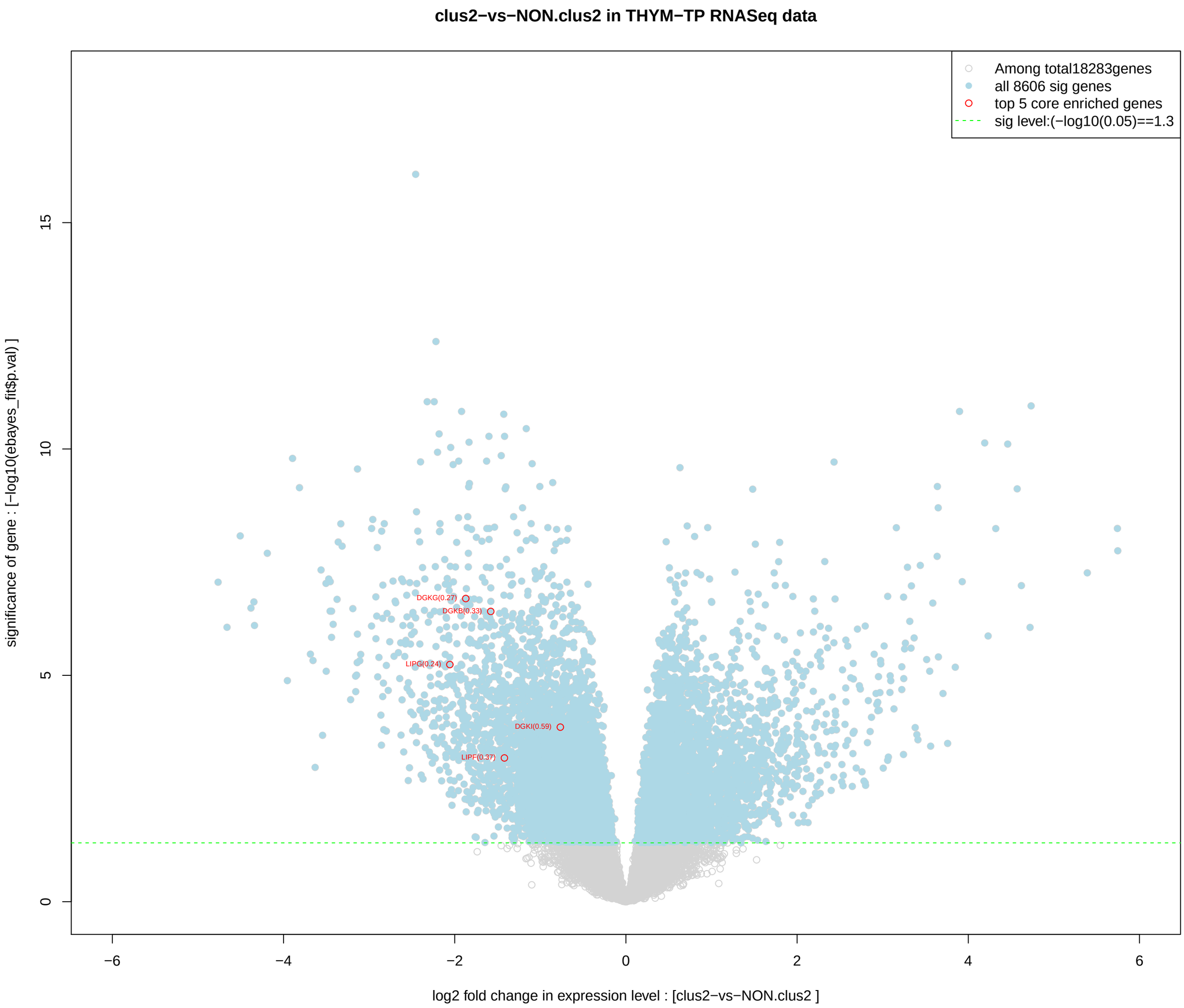
clus2−vs−NON.clus2 in THYM−TP RNASeq data
Among total18283genes
all 8606 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
DGKG(0.27)
DGKB(0.33)
LIPG(0.24)
DGKI(0.59)
LIPF(0.37)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
5
10
15
−6
−4
−2
0
2
4
6