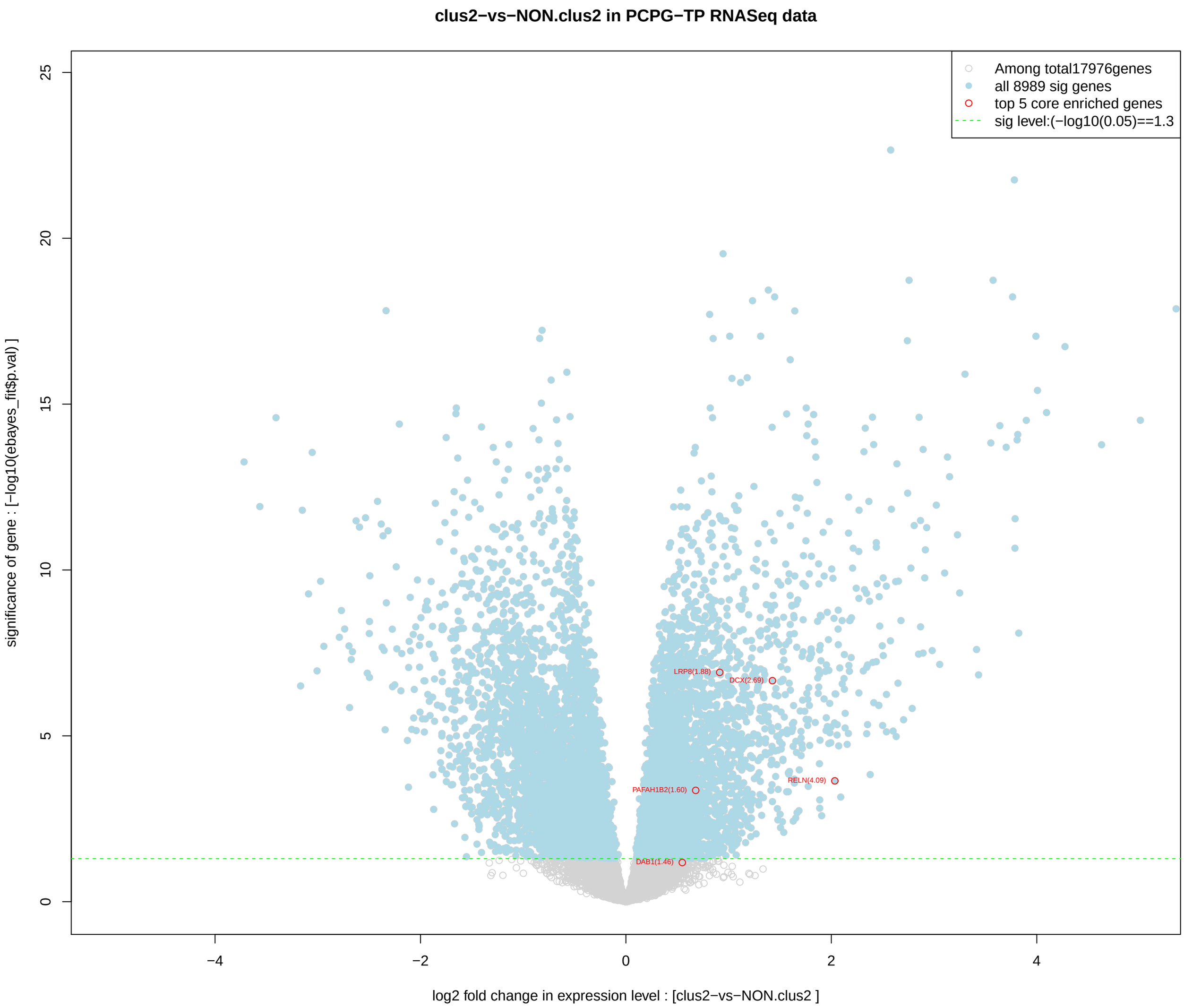
clus2−vs−NON.clus2 in PCPG−TP RNASeq data
Among total17976genes
all 8989 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
LRP8(1.88)
DCX(2.69)
RELN(4.09)
PAFAH1B2(1.60)
DAB1(1.46)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
5
10
15
20
25
−4
−2
0
2
4