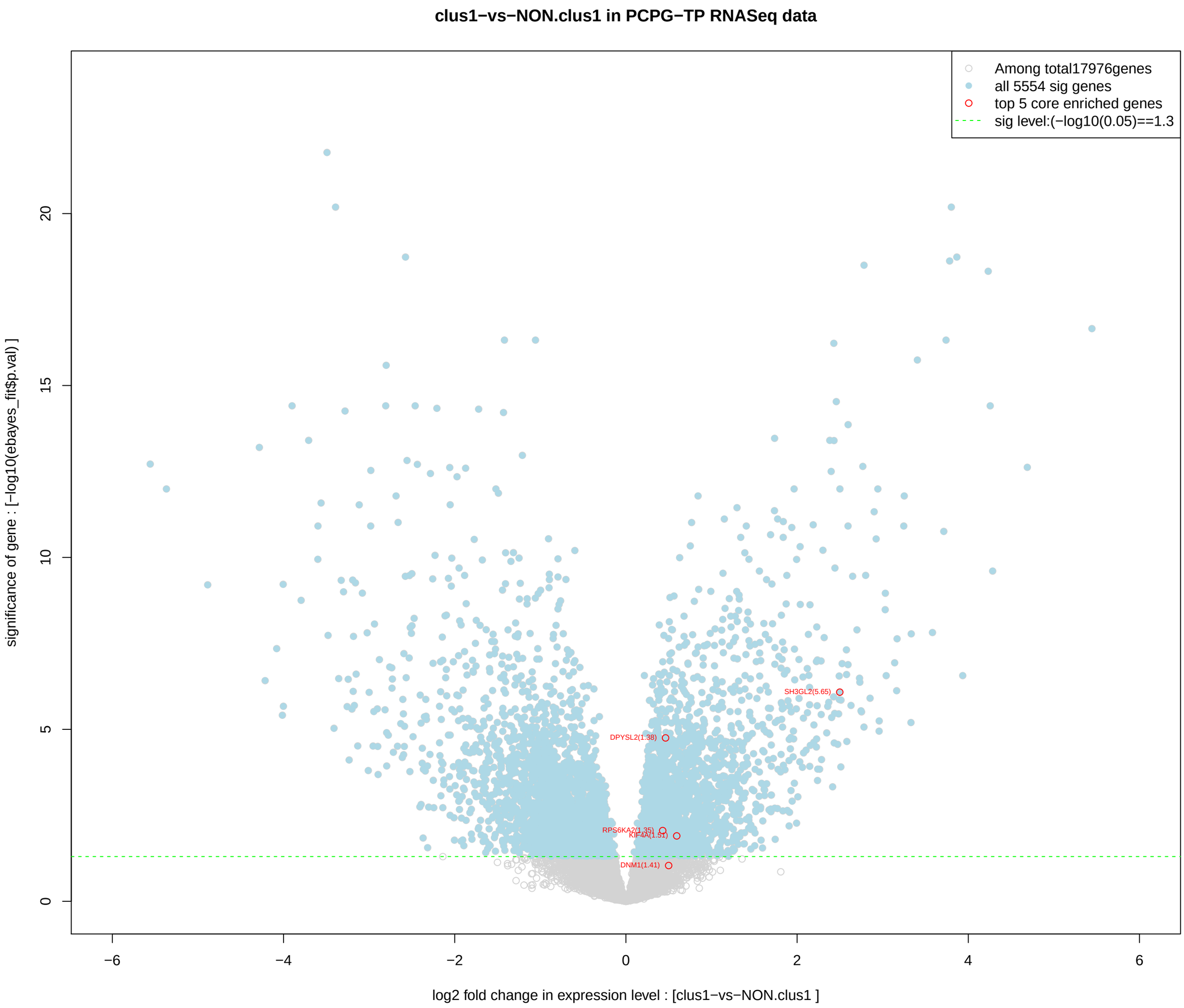
clus1−vs−NON.clus1 in PCPG−TP RNASeq data
Among total17976genes
all 5554 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
SH3GL2(5.65)
DPYSL2(1.38)
RPS6KA2(1.35)
KIF4A(1.51)
DNM1(1.41)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
5
10
15
20
−6
−4
−2
0
2
4
6