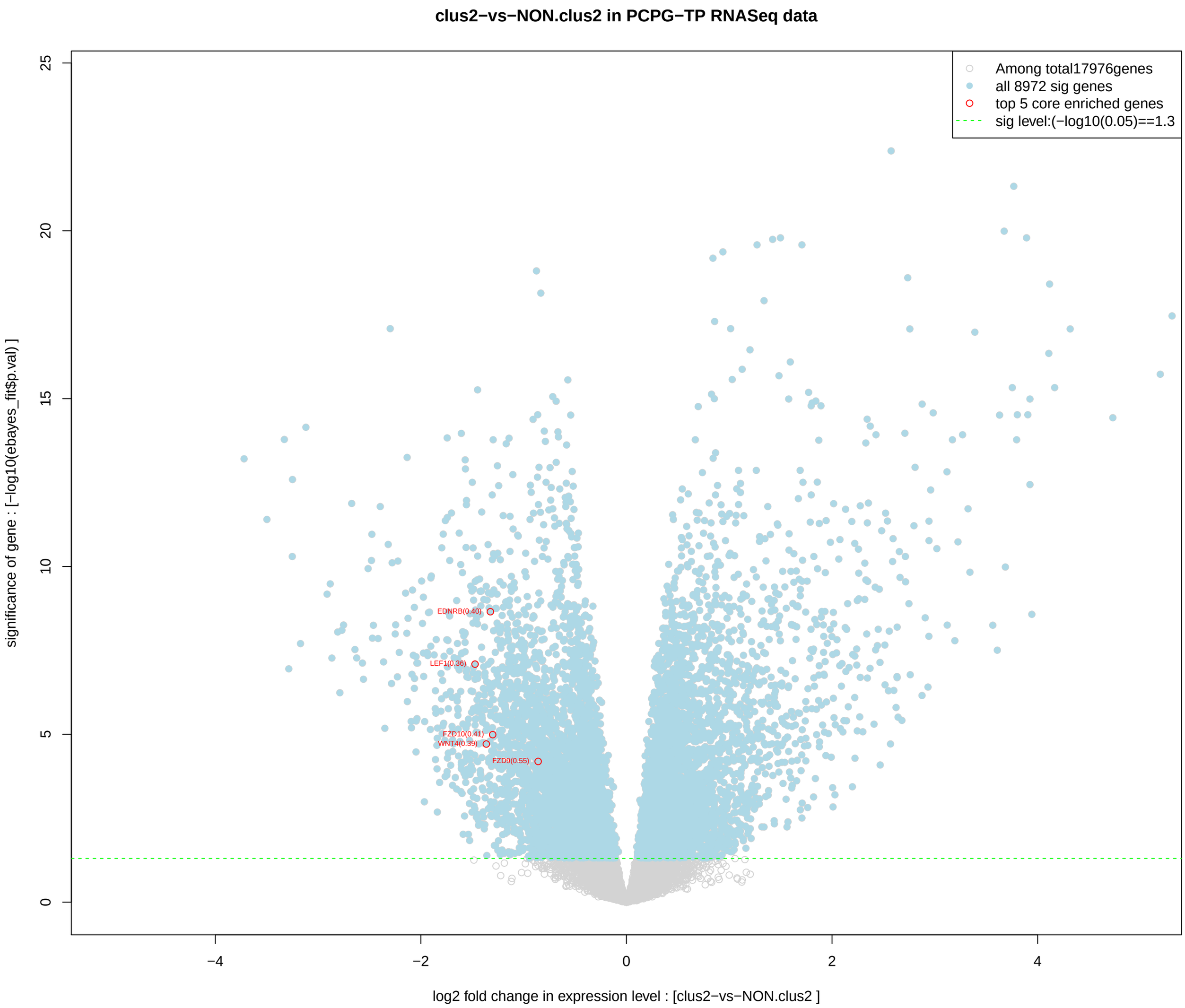
clus2−vs−NON.clus2 in PCPG−TP RNASeq data
Among total17976genes
all 8972 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
EDNRB(0.40)
LEF1(0.36)
FZD10(0.41)
WNT4(0.39)
FZD9(0.55)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
5
10
15
20
25
−4
−2
0
2
4