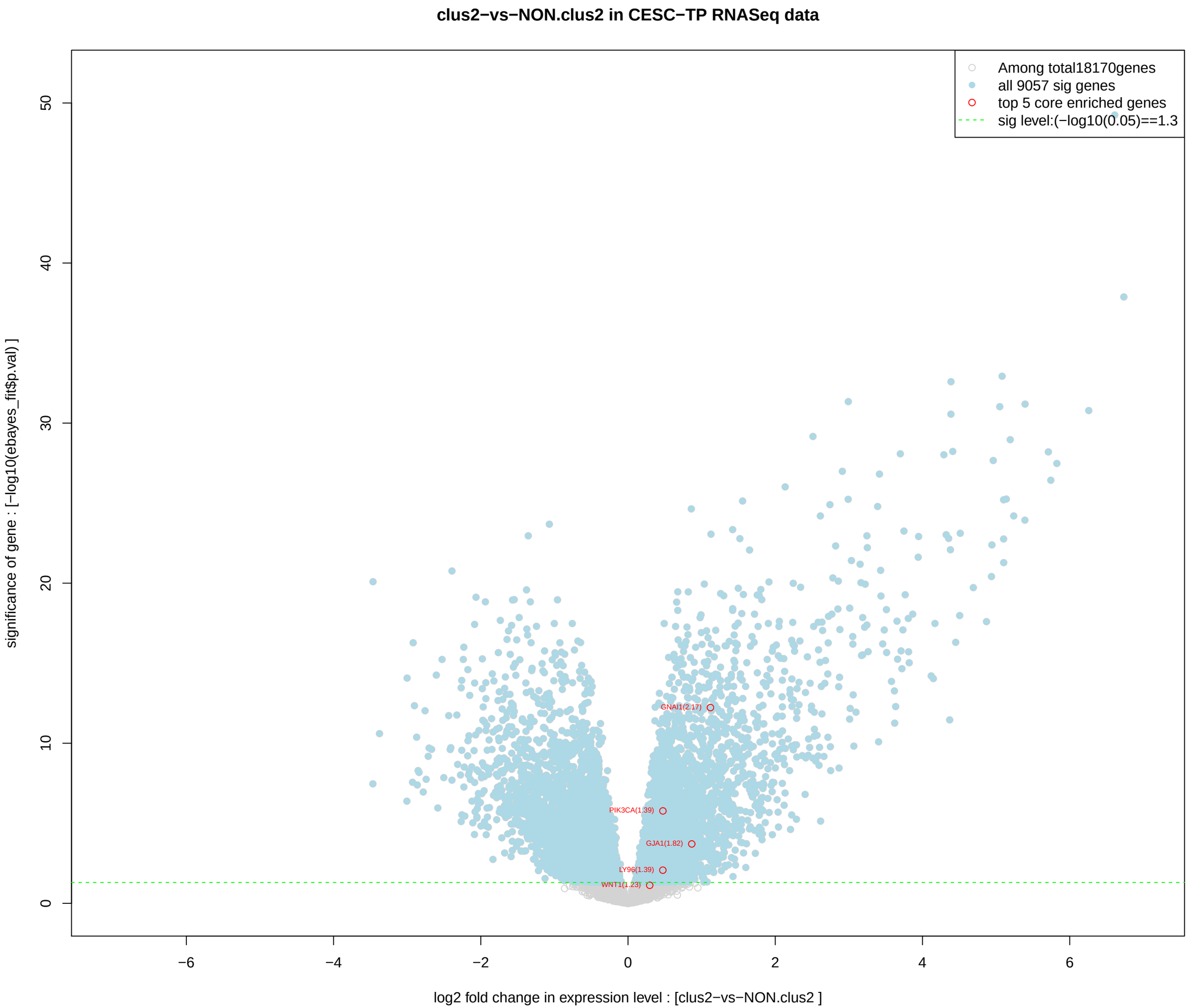
clus2−vs−NON.clus2 in CESC−TP RNASeq data
Among total18170genes
all 9057 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
GNAI1(2.17)
PIK3CA(1.39)
GJA1(1.82)
LY96(1.39)
WNT1(1.23)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
10
20
30
40
50
−6
−4
−2
0
2
4
6