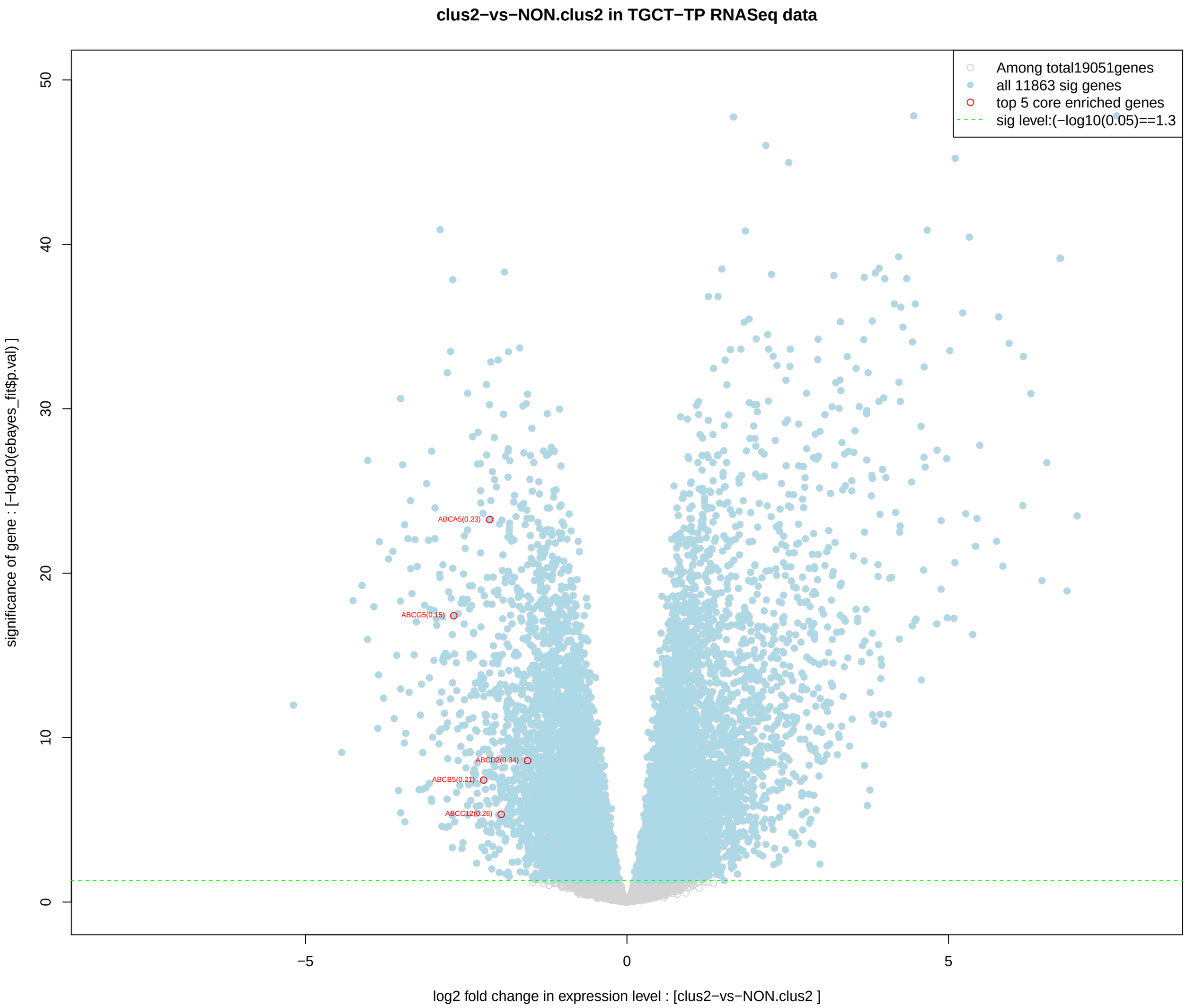
clus2−vs−NON.clus2 in TGCT−TP RNASeq data
Among total19051genes
all 11863 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ABCA5(0.23)
ABCG5(0.15)
ABCD2(0.34)
ABCB5(0.21)
ABCC12(0.26)
significance of gene : [−log10(ebayes_fit$p.val) ]
log2 fold change in expression level : [clus2−vs−NON.clus2 ]
0
10
20
30
40
50
−5
0
5