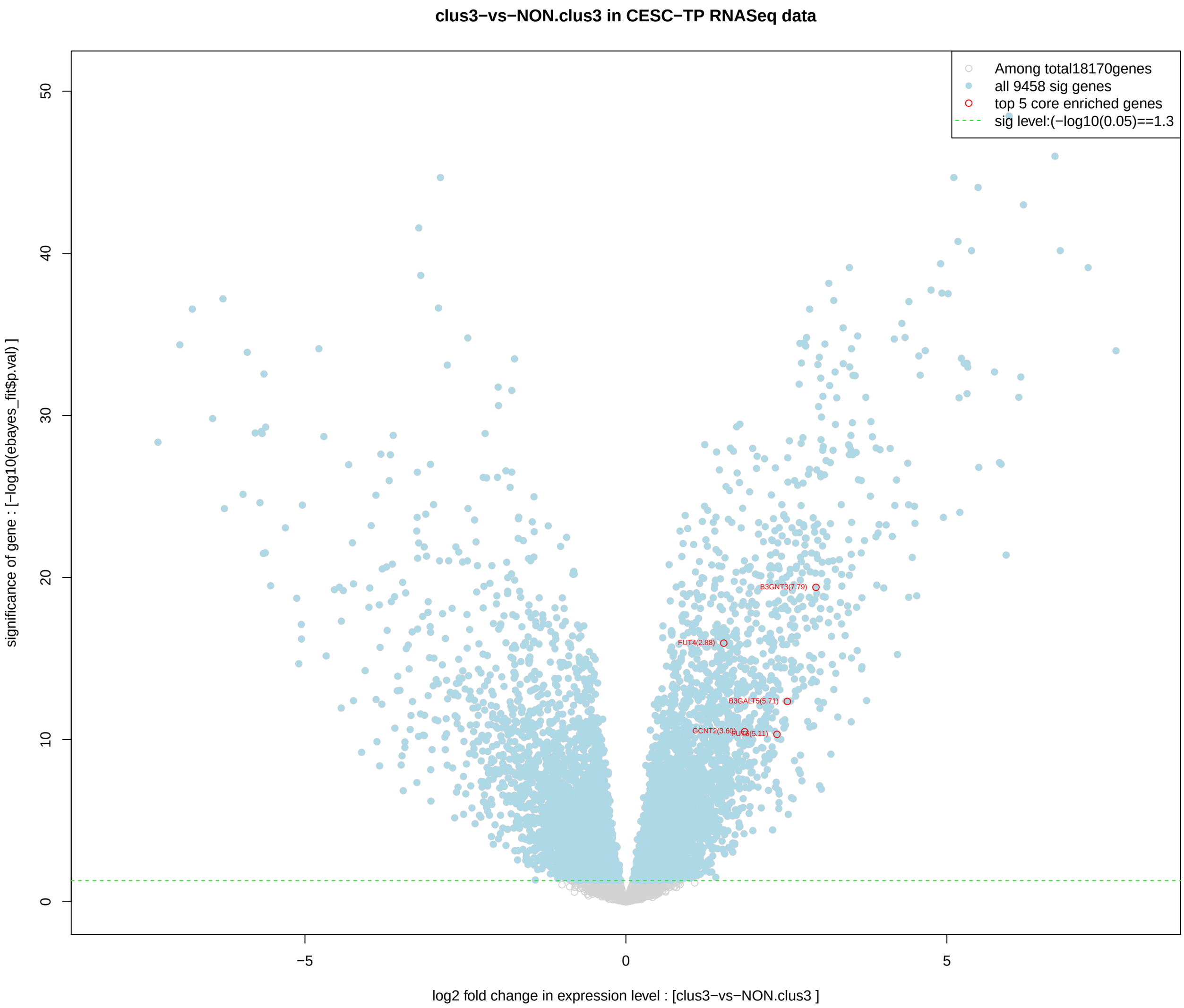
clus3−vs−NON.clus3 in CESC−TP RNASeq data
Among total18170genes
all 9458 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
B3GNT3(7.79)
FUT4(2.88)
B3GALT5(5.71)
GCNT2(3.60)
FUT9(5.11)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
10
20
30
40
50
−5
0
5