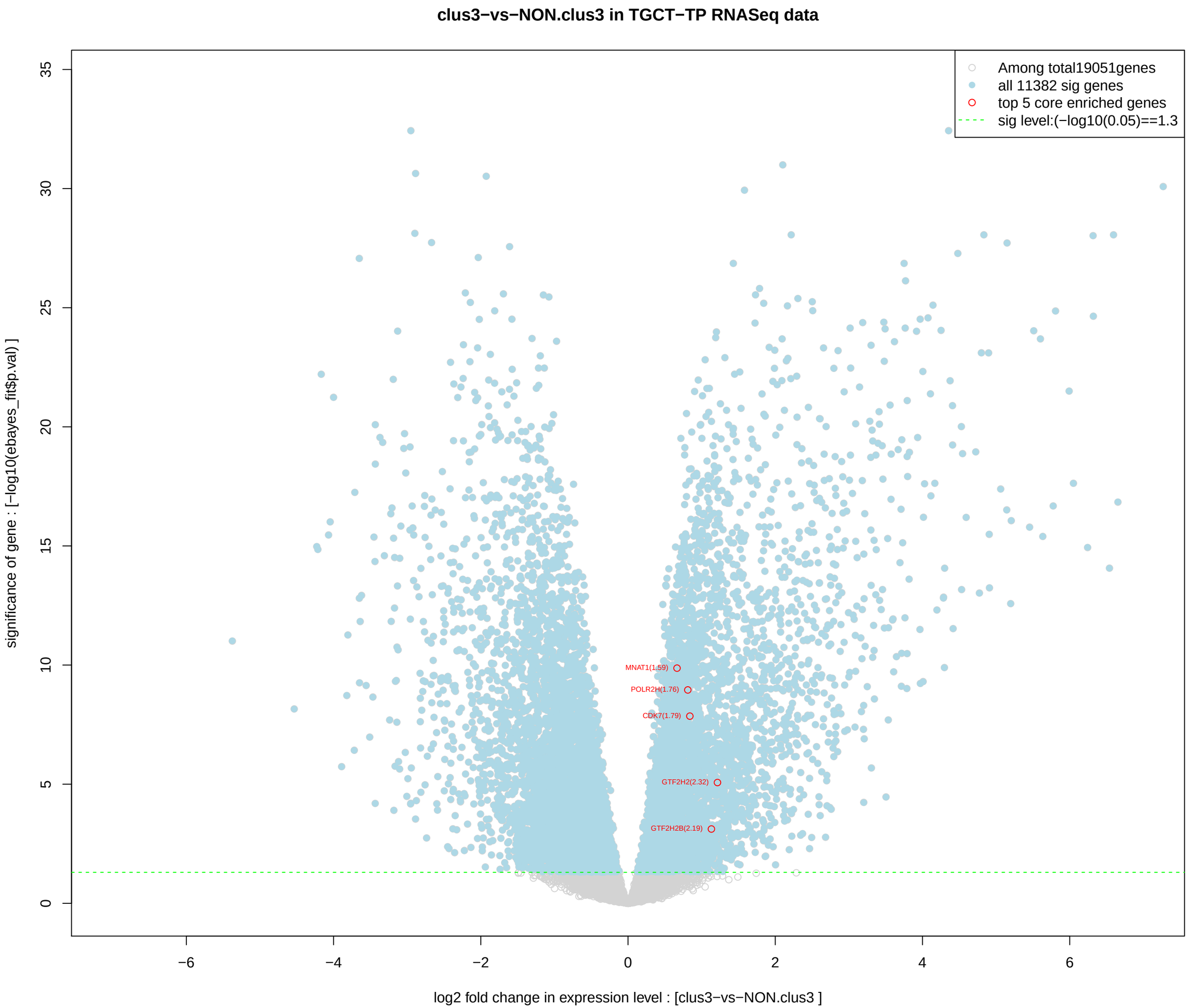
clus3−vs−NON.clus3 in TGCT−TP RNASeq data
Among total19051genes
all 11382 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
MNAT1(1.59)
POLR2H(1.76)
CDK7(1.79)
GTF2H2(2.32)
GTF2H2B(2.19)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
5
10
15
20
25
30
35
−6
−4
−2
0
2
4
6