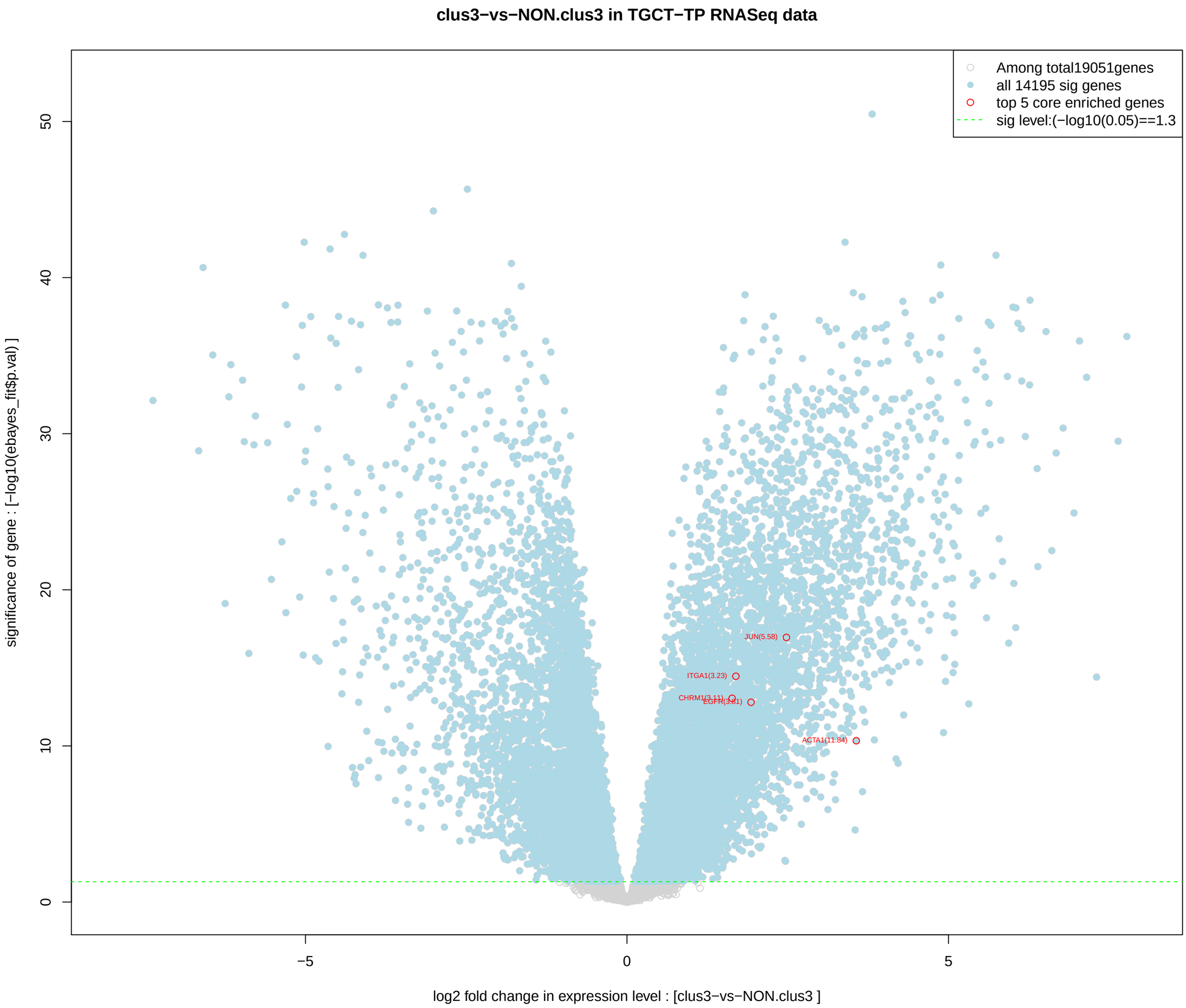
clus3−vs−NON.clus3 in TGCT−TP RNASeq data
Among total19051genes
all 14195 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
JUN(5.58)
ITGA1(3.23)
CHRM1(3.11)
EGFR(3.81)
ACTA1(11.84)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
10
20
30
40
50
−5
0
5