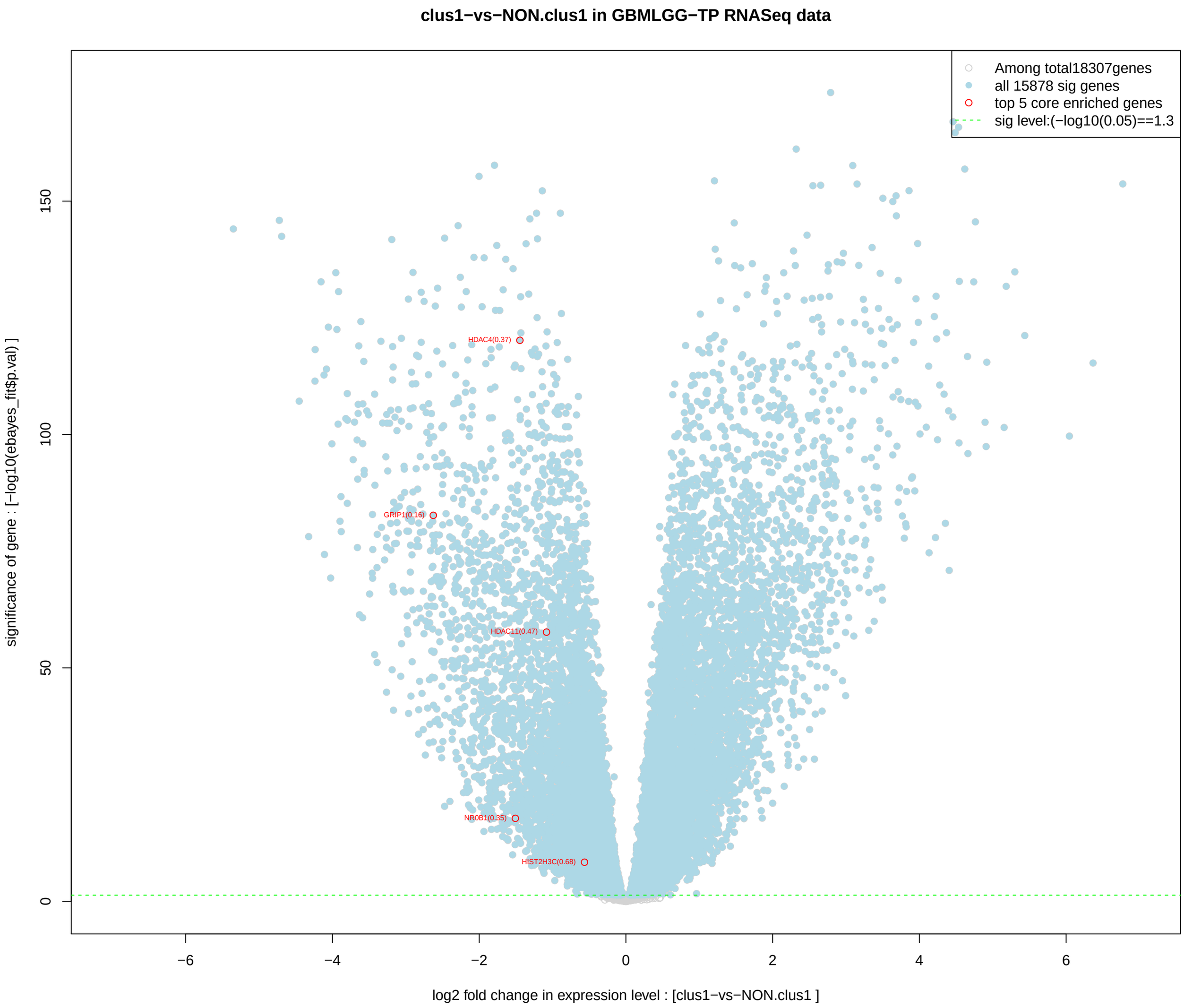
clus1−vs−NON.clus1 in GBMLGG−TP RNASeq data
Among total18307genes
all 15878 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
HDAC4(0.37)
GRIP1(0.16)
HDAC11(0.47)
NR0B1(0.35)
HIST2H3C(0.68)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
50
100
150
−6
−4
−2
0
2
4
6