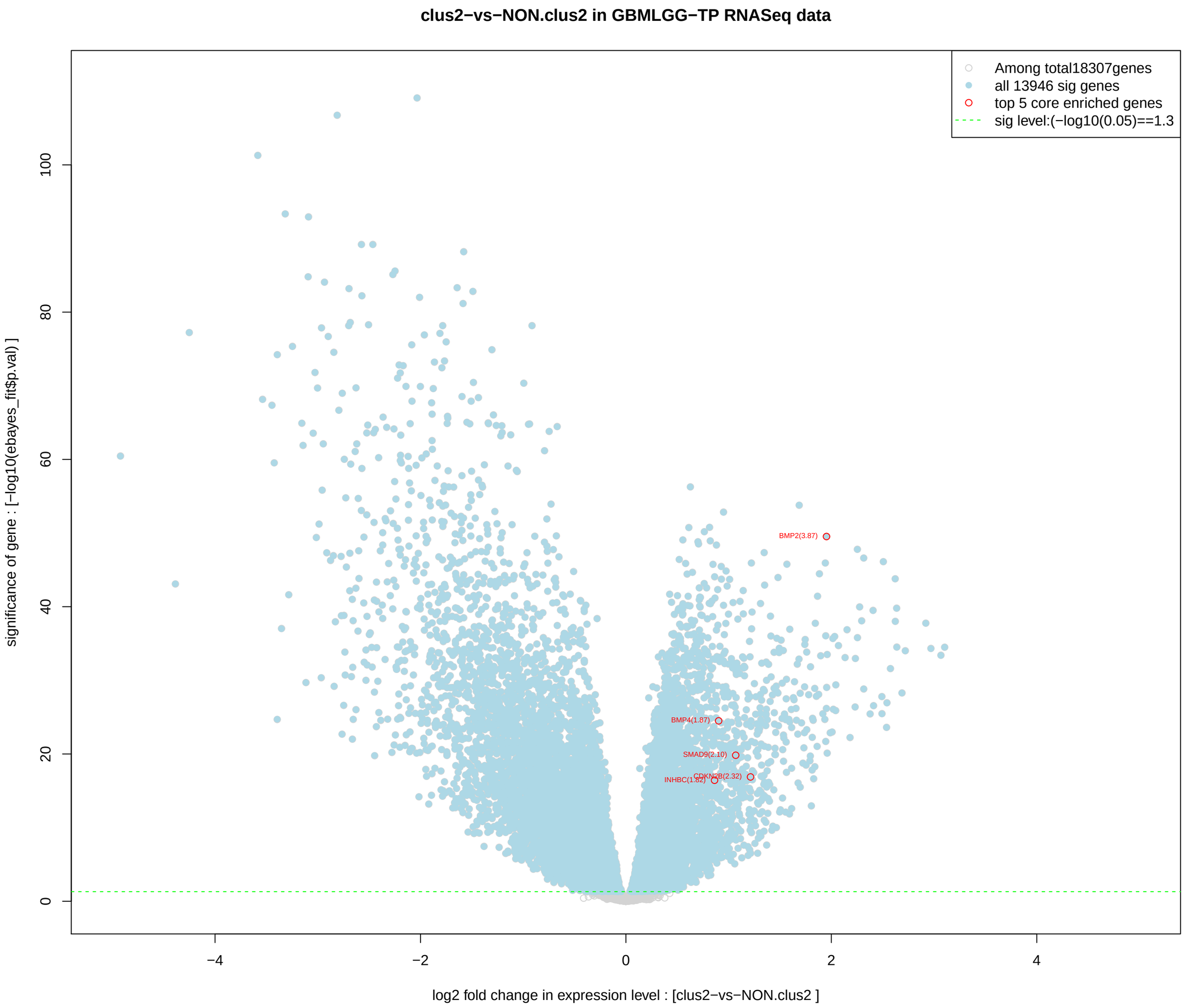
clus2−vs−NON.clus2 in GBMLGG−TP RNASeq data
Among total18307genes
all 13946 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
BMP2(3.87)
BMP4(1.87)
SMAD9(2.10)
CDKN2B(2.32)
INHBC(1.82)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
20
40
60
80
100
−4
−2
0
2
4