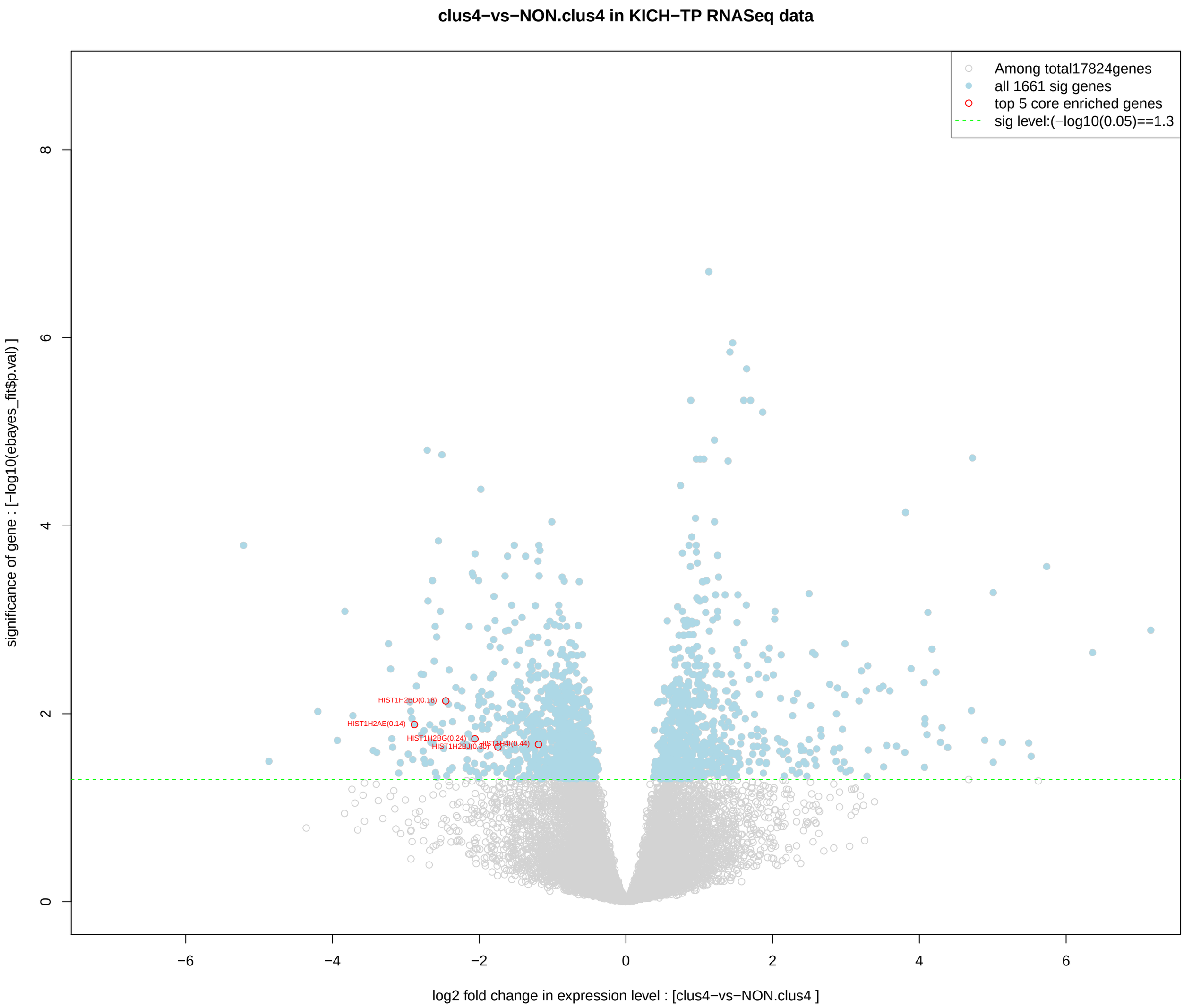
clus4−vs−NON.clus4 in KICH−TP RNASeq data
Among total17824genes
all 1661 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
HIST1H2BD(0.18)
HIST1H2AE(0.14)
HIST1H2BG(0.24)
HIST1H2BJ(0.30)
HIST1H4I(0.44)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0
2
4
6
8
−6
−4
−2
0
2
4
6