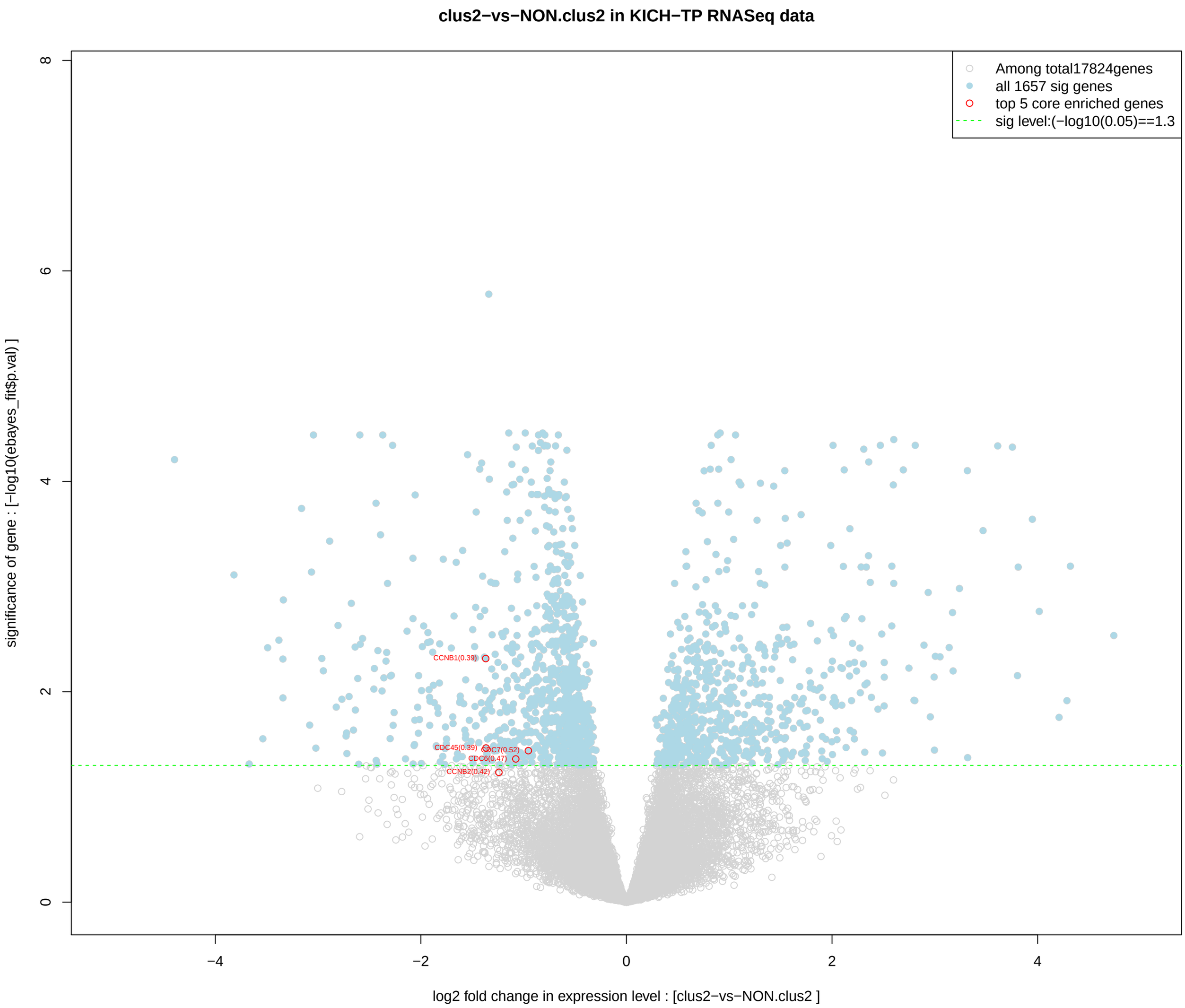
clus2−vs−NON.clus2 in KICH−TP RNASeq data
Among total17824genes
all 1657 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CCNB1(0.39)
CDC45(0.39)
CDC7(0.52)
CDC6(0.47)
CCNB2(0.42)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
2
4
6
8
−4
−2
0
2
4