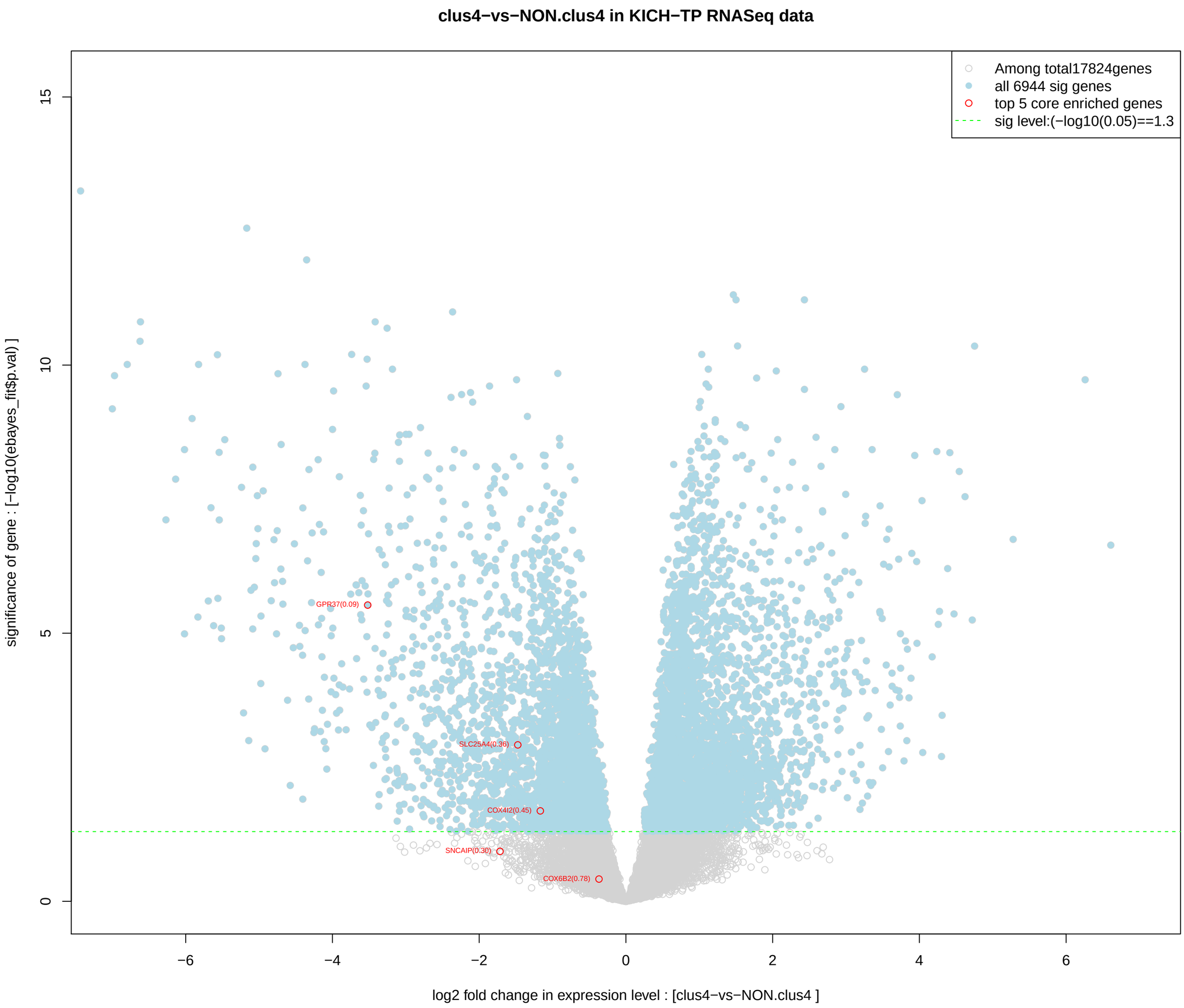
clus4−vs−NON.clus4 in KICH−TP RNASeq data
Among total17824genes
all 6944 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
GPR37(0.09)
SLC25A4(0.36)
COX4I2(0.45)
SNCAIP(0.30)
COX6B2(0.78)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0
5
10
15
−6
−4
−2
0
2
4
6