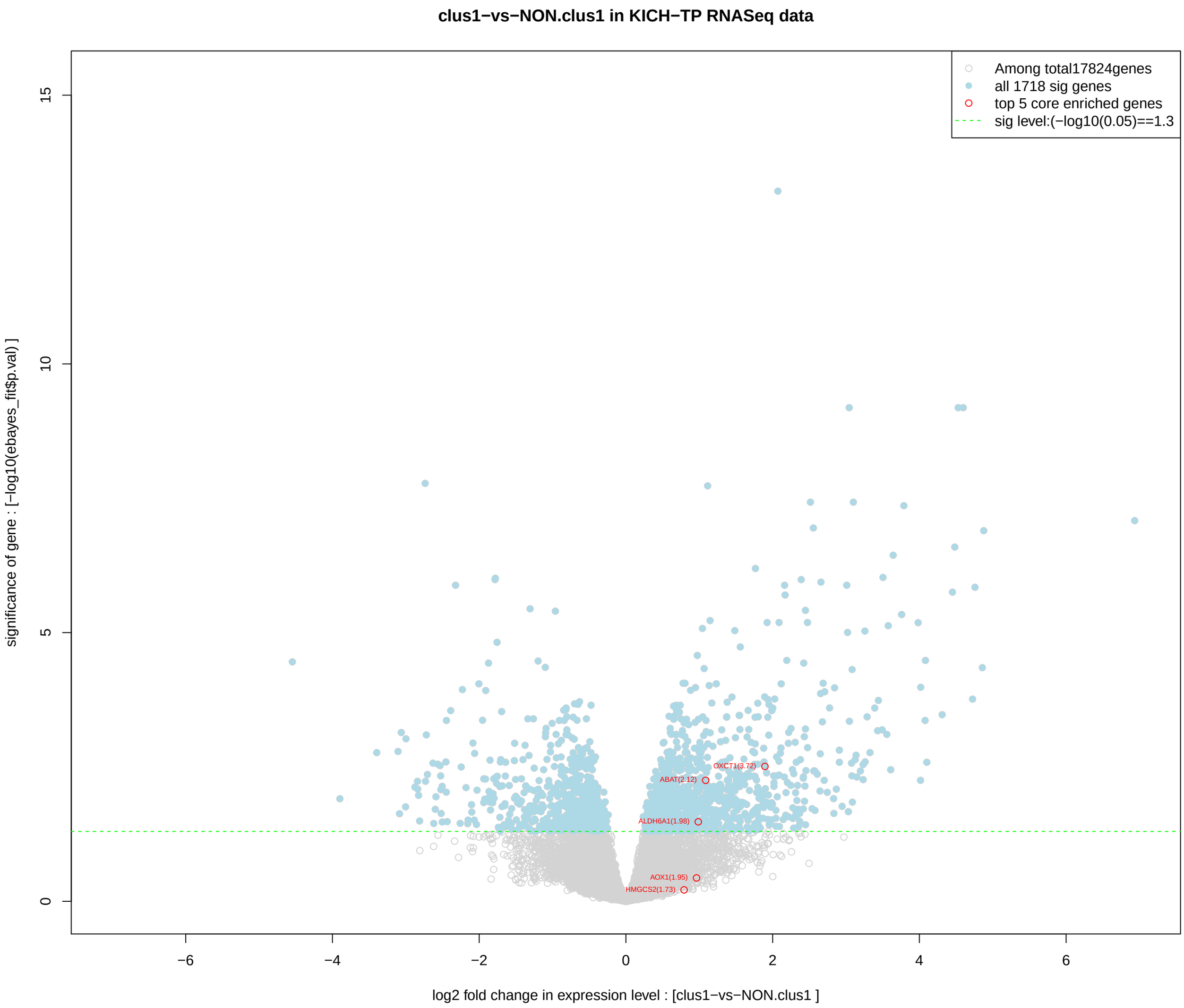
clus1−vs−NON.clus1 in KICH−TP RNASeq data
Among total17824genes
all 1718 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
OXCT1(3.72)
ABAT(2.12)
ALDH6A1(1.98)
AOX1(1.95)
HMGCS2(1.73)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
5
10
15
−6
−4
−2
0
2
4
6