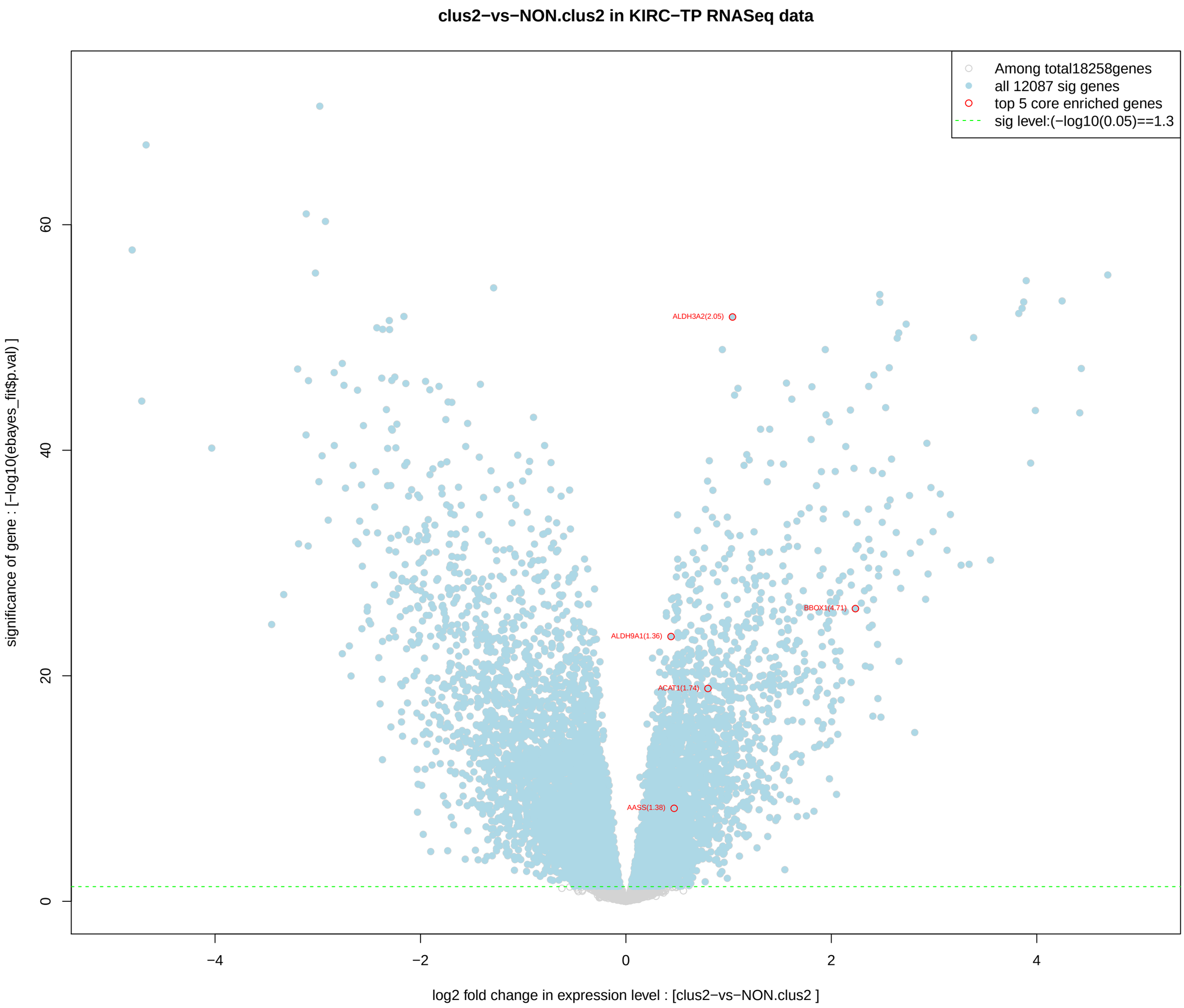
clus2−vs−NON.clus2 in KIRC−TP RNASeq data
Among total18258genes
all 12087 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ALDH3A2(2.05)
BBOX1(4.71)
ALDH9A1(1.36)
ACAT1(1.74)
AASS(1.38)
significance of gene : [−log10(ebayes_fit$p.val)]
0
20
40
60
−4
−2
0
2
4
log2 fold change in expression level : [clus2−vs−NON.clus2]