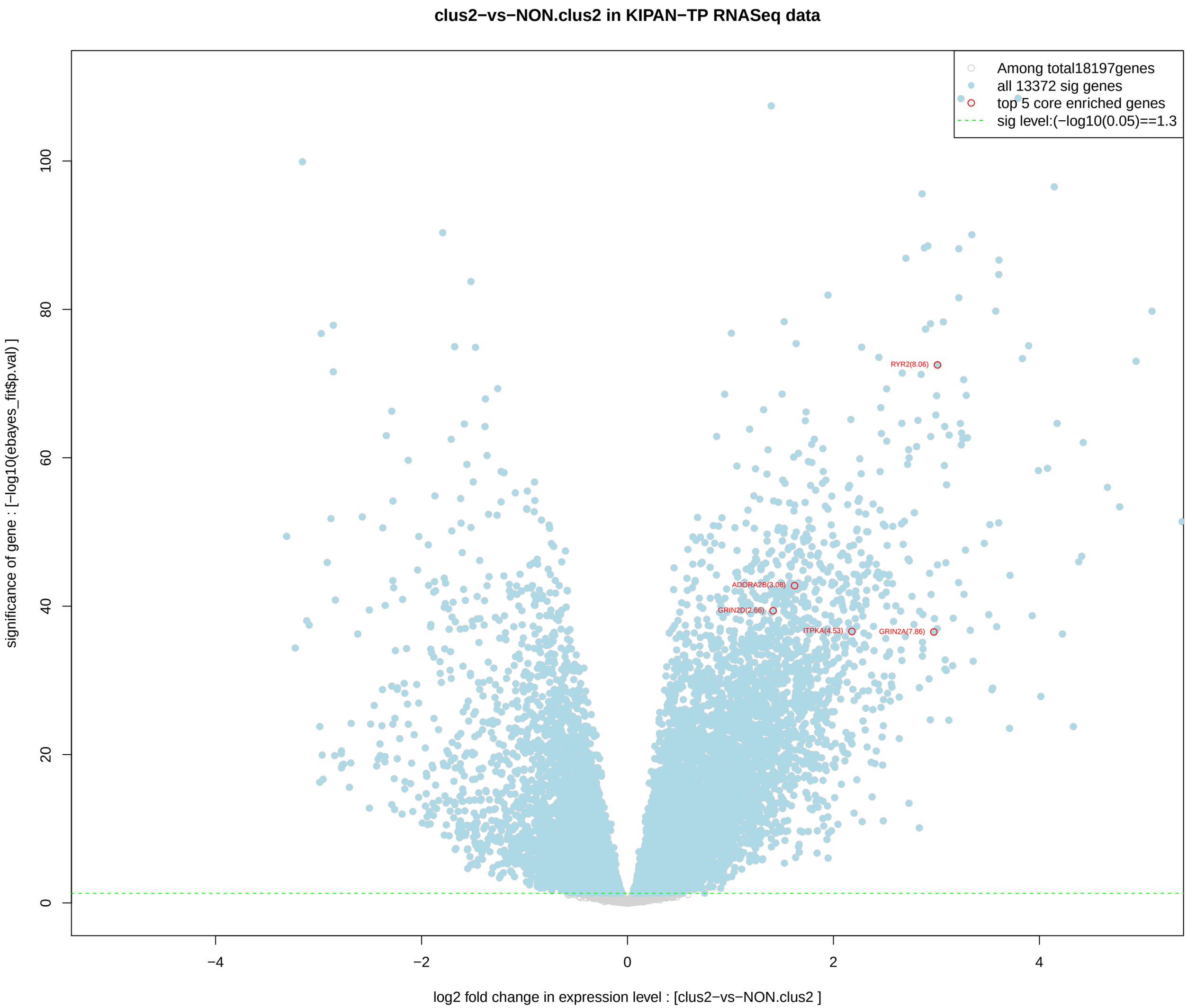
clus2−vs−NON.clus2 in KIPAN−TP RNASeq data
Among total18197genes
all 13372 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
RYR2(8.06)
ADORA2B(3.08)
GRIN2D(2.66)
ITPKA(4.53)
GRIN2A(7.86)
significance of gene : [−log10(ebayes_fit$p.val) ]
log2 fold change in expression level : [clus2−vs−NON.clus2 ]
0
20
40
60
80
100
−4
−2
0
2
4