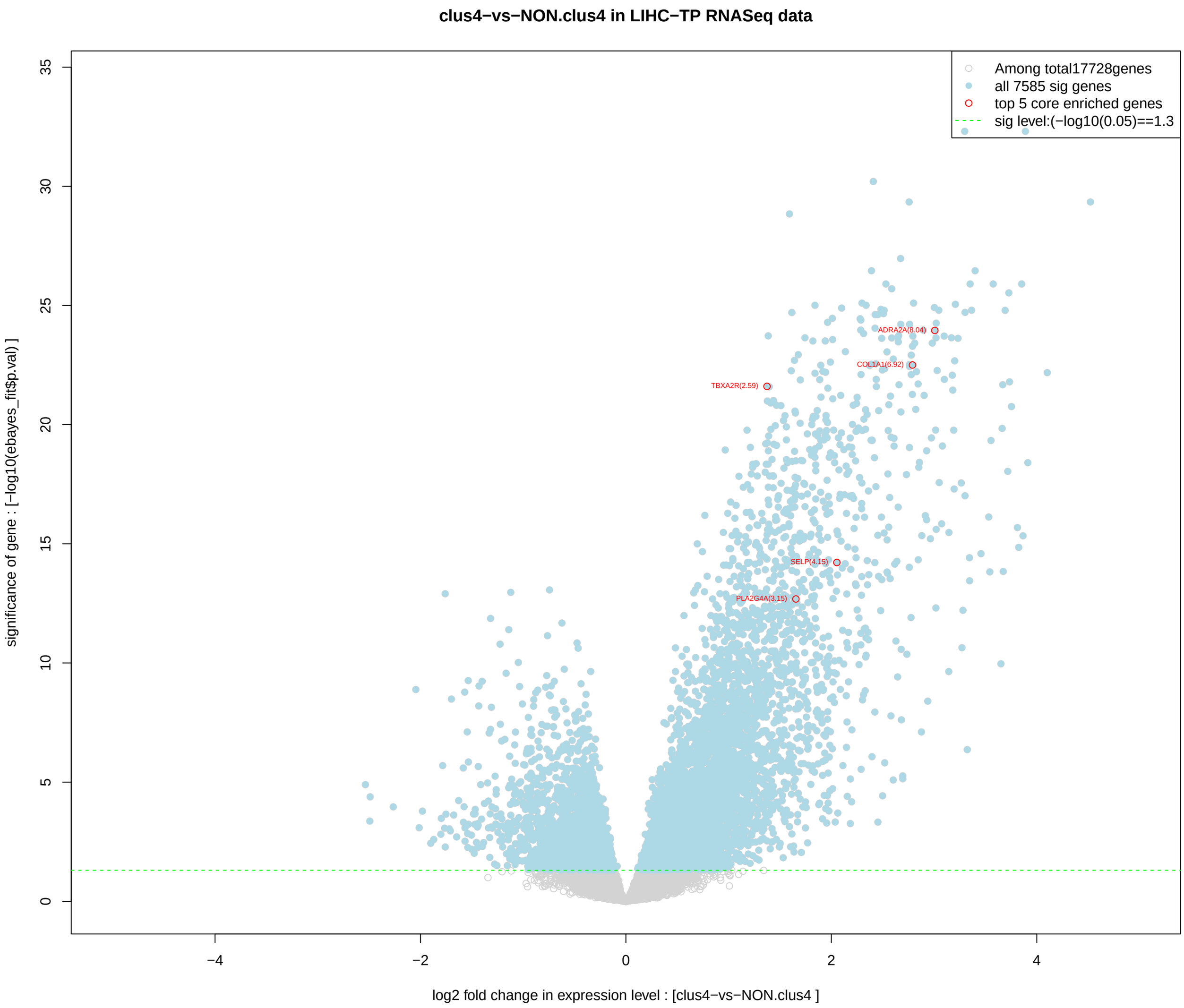
clus4−vs−NON.clus4 in LIHC−TP RNASeq data
Among total17728genes
all 7585 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ADRA2A(8.04)
COL1A1(6.92)
TBXA2R(2.59)
SELP(4.15)
PLA2G4A(3.15)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
35
30
25
20
15
10
5
0
−4
−2
0
2
4