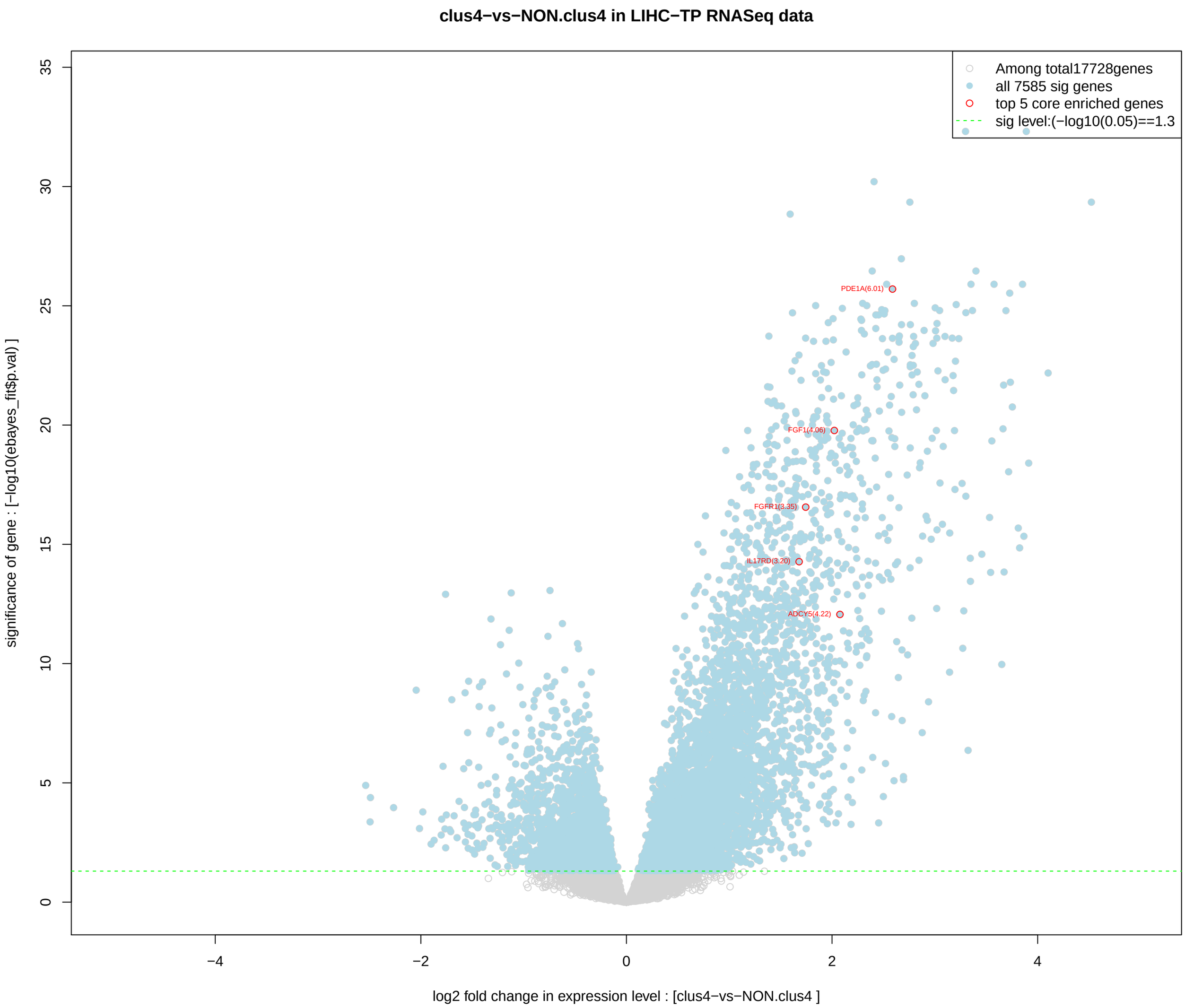
clus4−vs−NON.clus4 in LIHC−TP RNASeq data
Among total17728genes
all 7585 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PDE1A(6.01)
FGF1(4.06)
FGFR1(3.35)
IL17RD(3.20)
ADCY5(4.22)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0
5
10
15
20
25
30
35
−4
−2
0
2
4