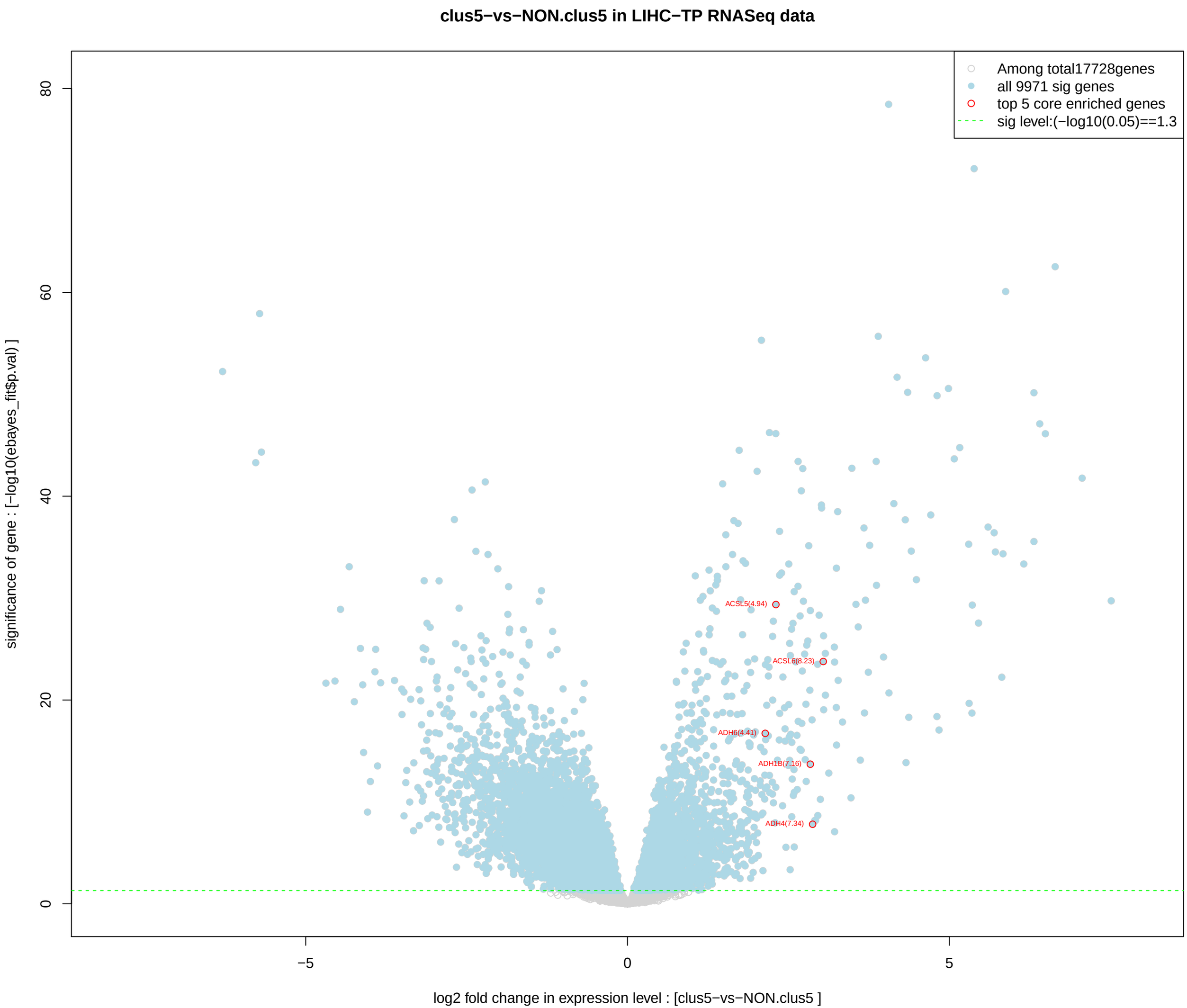
clus5−vs−NON.clus5 in LIHC−TP RNASeq data
Among total17728genes
all 9971 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ACSL5(4.94)
ACSL6(8.23)
ADH6(4.41)
ADH1B(7.16)
ADH4(7.34)
significance of gene : [−log10(ebayes_fit$p.val)]
0
20
40
60
80
−5
0
5
log2 fold change in expression level : [clus5−vs−NON.clus5]