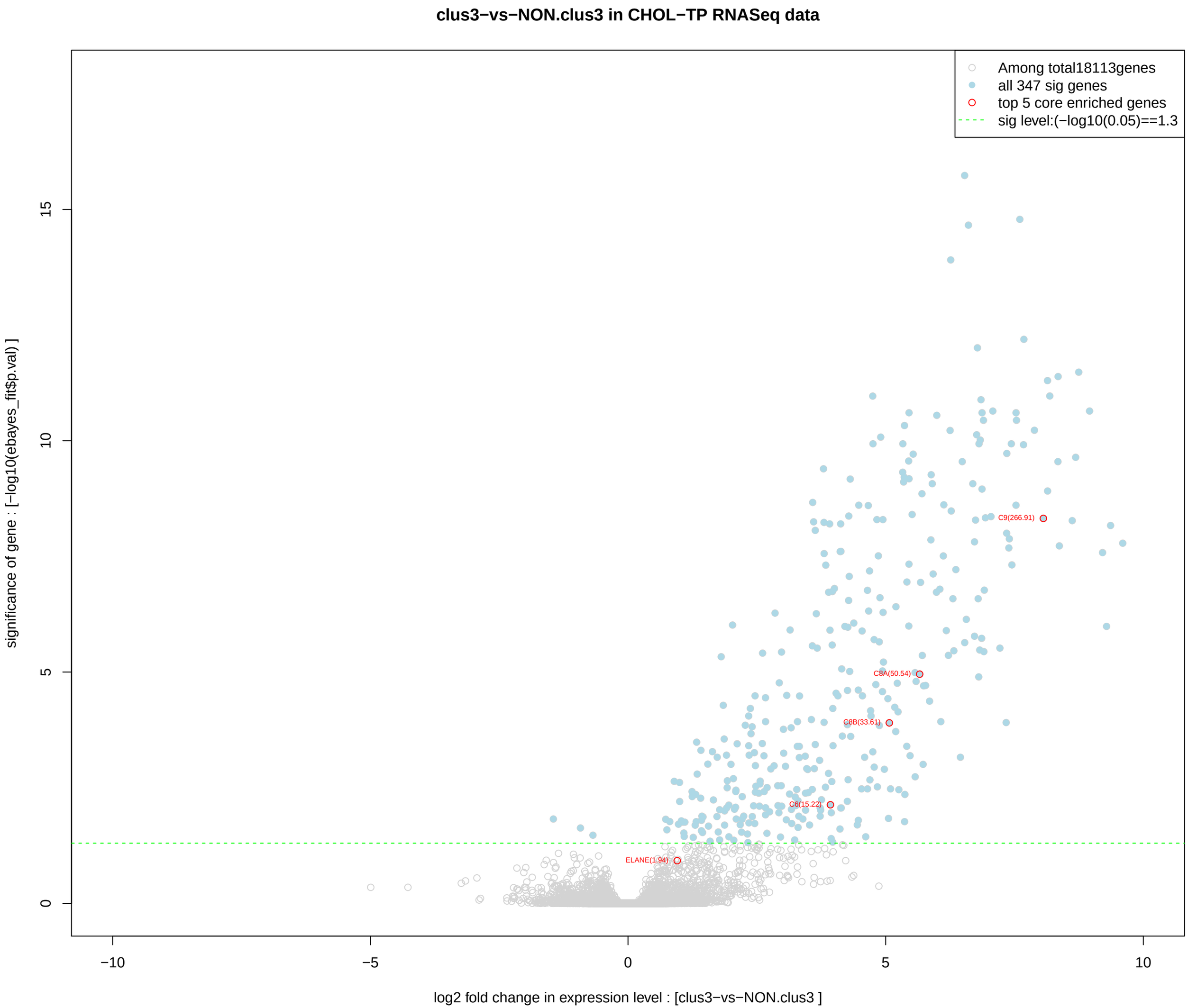
clus3−vs−NON.clus3 in CHOL−TP RNASeq data
Among total18113genes
all 347 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
C9(266.91)
C8A(50.54)
C8B(33.61)
C6(15.22)
ELANE(1.94)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
5
10
15
−10
−5
0
5
10