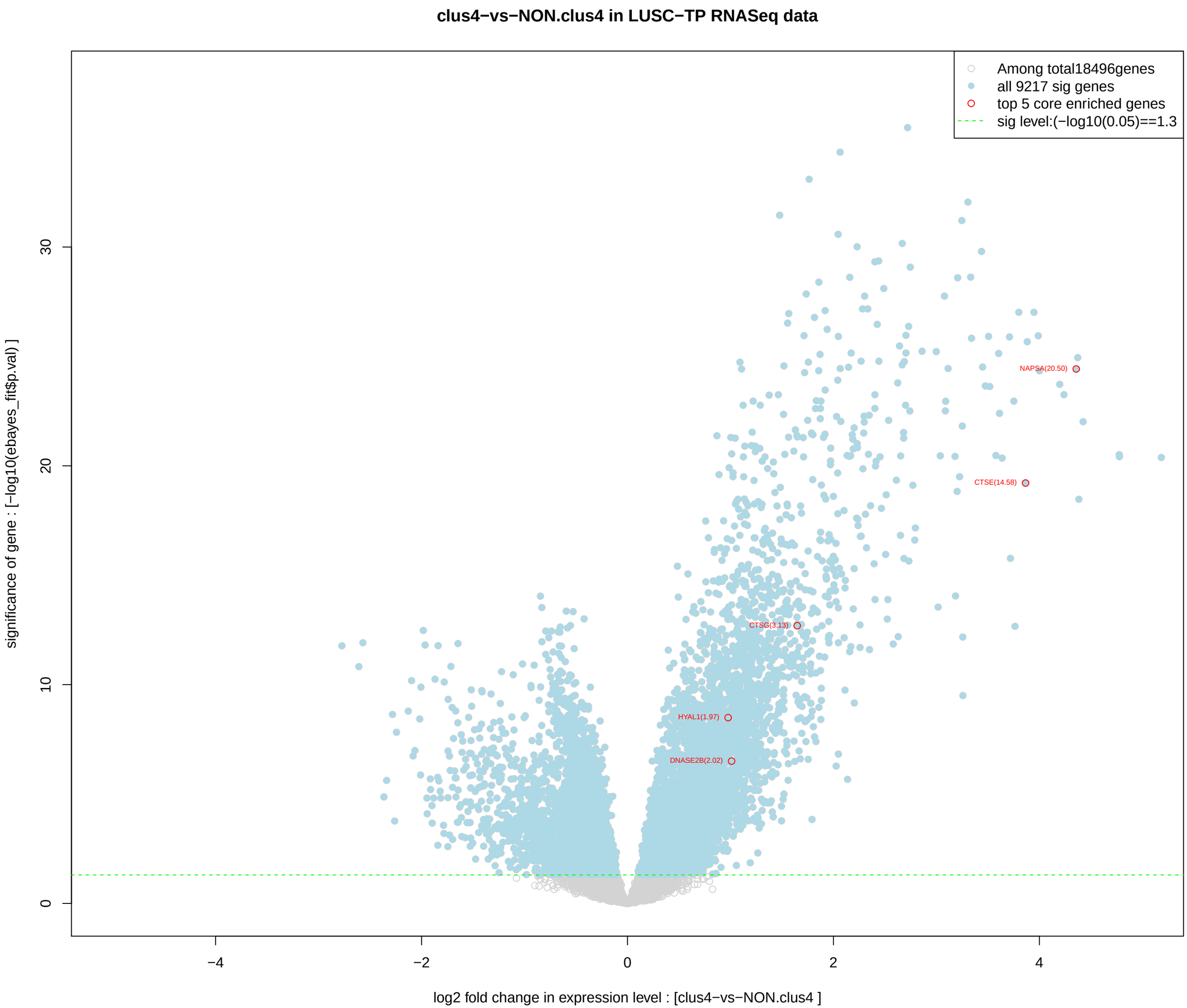
clus4−vs−NON.clus4 in LUSC−TP RNASeq data
Among total18496genes
all 9217 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
NAPSA(20.50)
CTSE(14.58)
CTSG(3.13)
HYAL1(1.97)
DNASE2B(2.02)
significance of gene : [−log10(ebayes_fit$p.val)]
0
10
20
30
−4
−2
0
2
4
log2 fold change in expression level : [clus4−vs−NON.clus4]