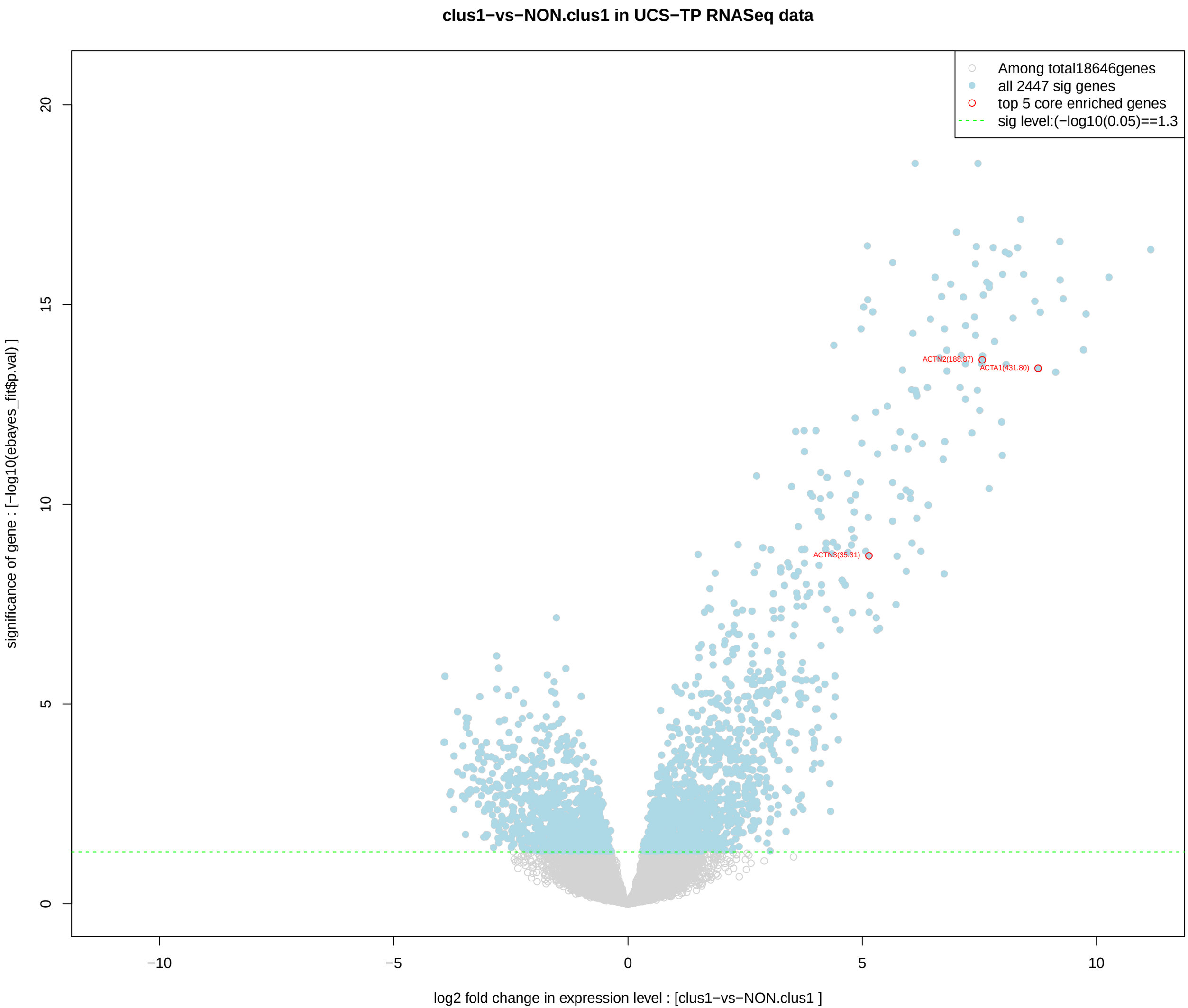
clus1−vs−NON.clus1 in UCS−TP RNASeq data
Among total18646genes
all 2447 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ACTN2(188.87)
ACTA1(431.80)
ACTN3(35.31)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
5
10
15
20
−10
−5
0
5
10