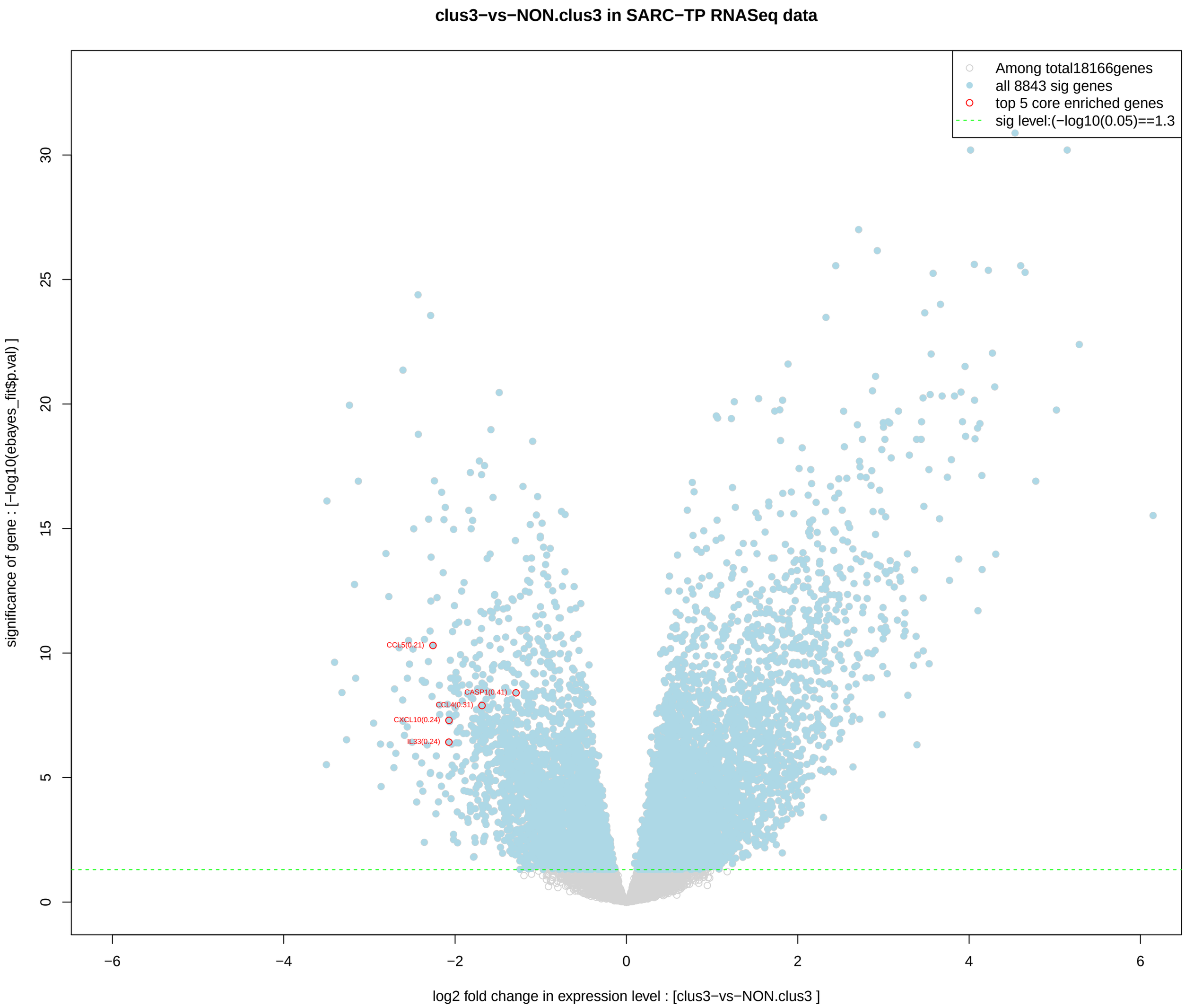
clus3−vs−NON.clus3 in SARC−TP RNASeq data
Among total18166genes
all 8843 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CCL5(0.21)
CASP1(0.41)
CCL4(0.31)
CXCL10(0.24)
IL33(0.24)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
5
10
15
20
25
30
−6
−4
−2
0
2
4
6