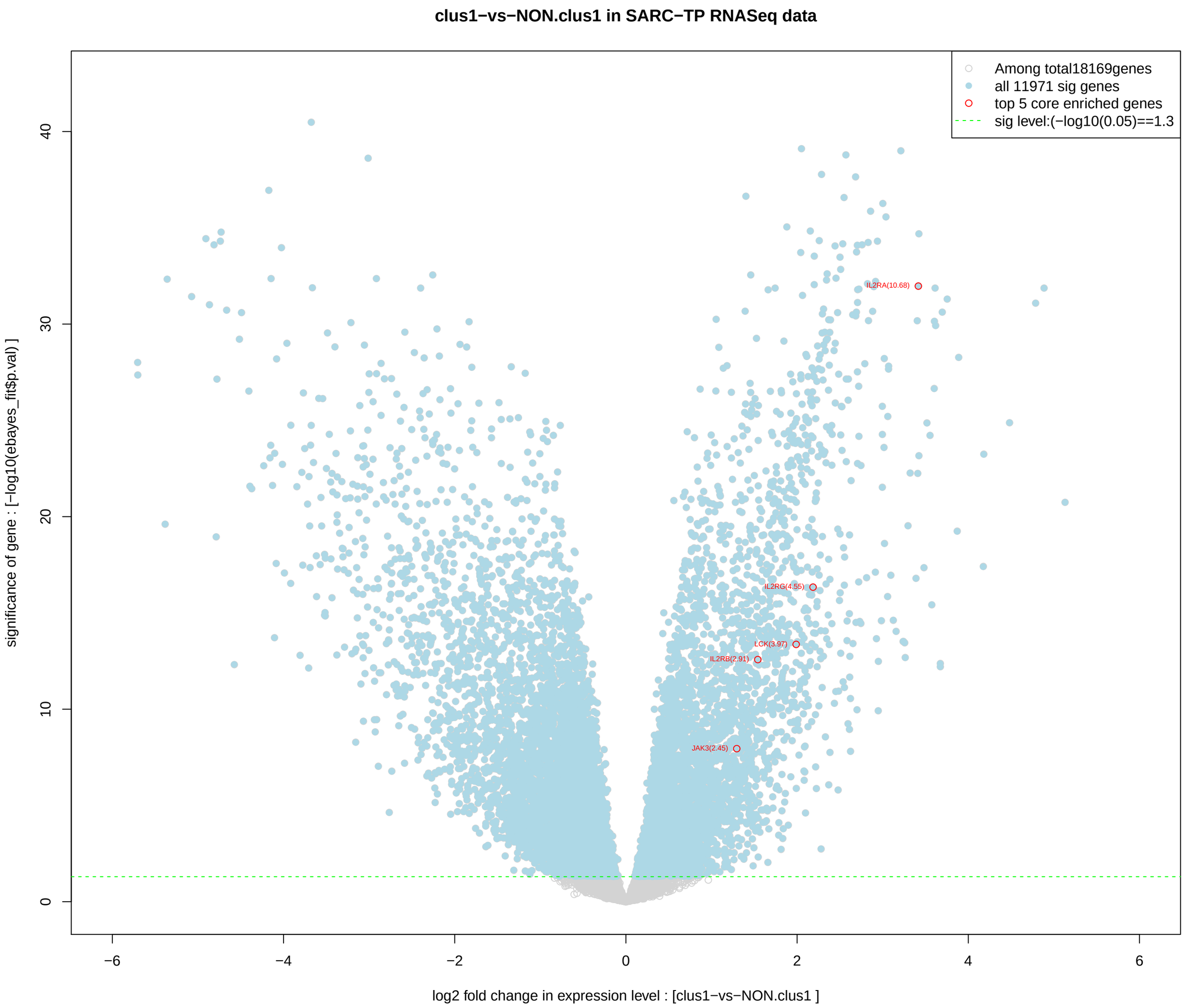
clus1−vs−NON.clus1 in SARC−TP RNASeq data
Among total18169genes
all 11971 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
IL2RA(10.68)
IL2RG(4.55)
LCK(3.97)
IL2RB(2.91)
JAK3(2.45)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
10
20
30
40
−6
−4
−2
0
2
4
6