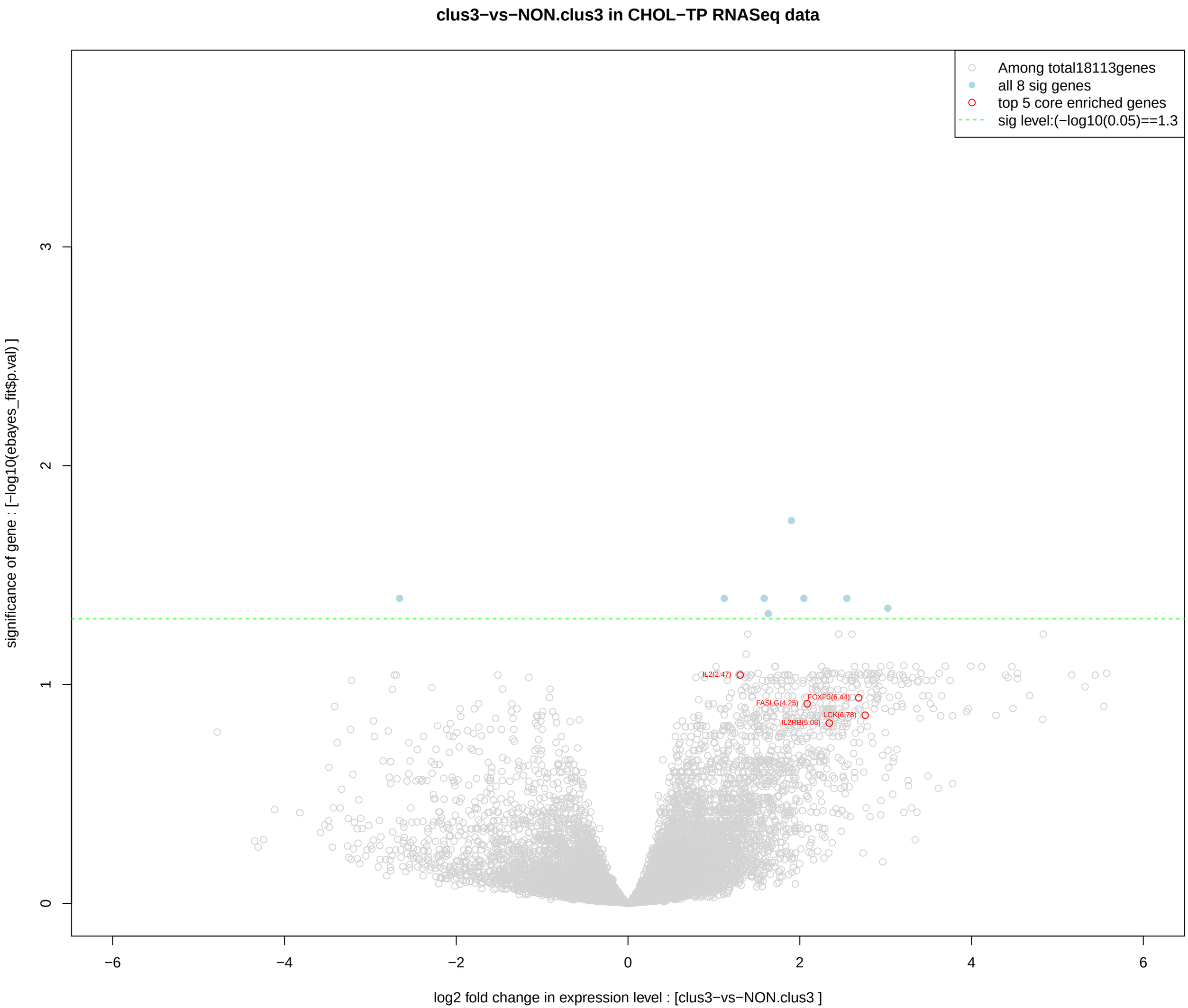
clus3−vs−NON.clus3 in CHOL−TP RNASeq data
Among total18113genes
all 8 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
IL2(2.47)
FASLG(4.25)
FOXP3(6.44)
LCK(6.78)
IL2RB(5.08)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
1
2
3
−6
−4
−2
0
2
4
6