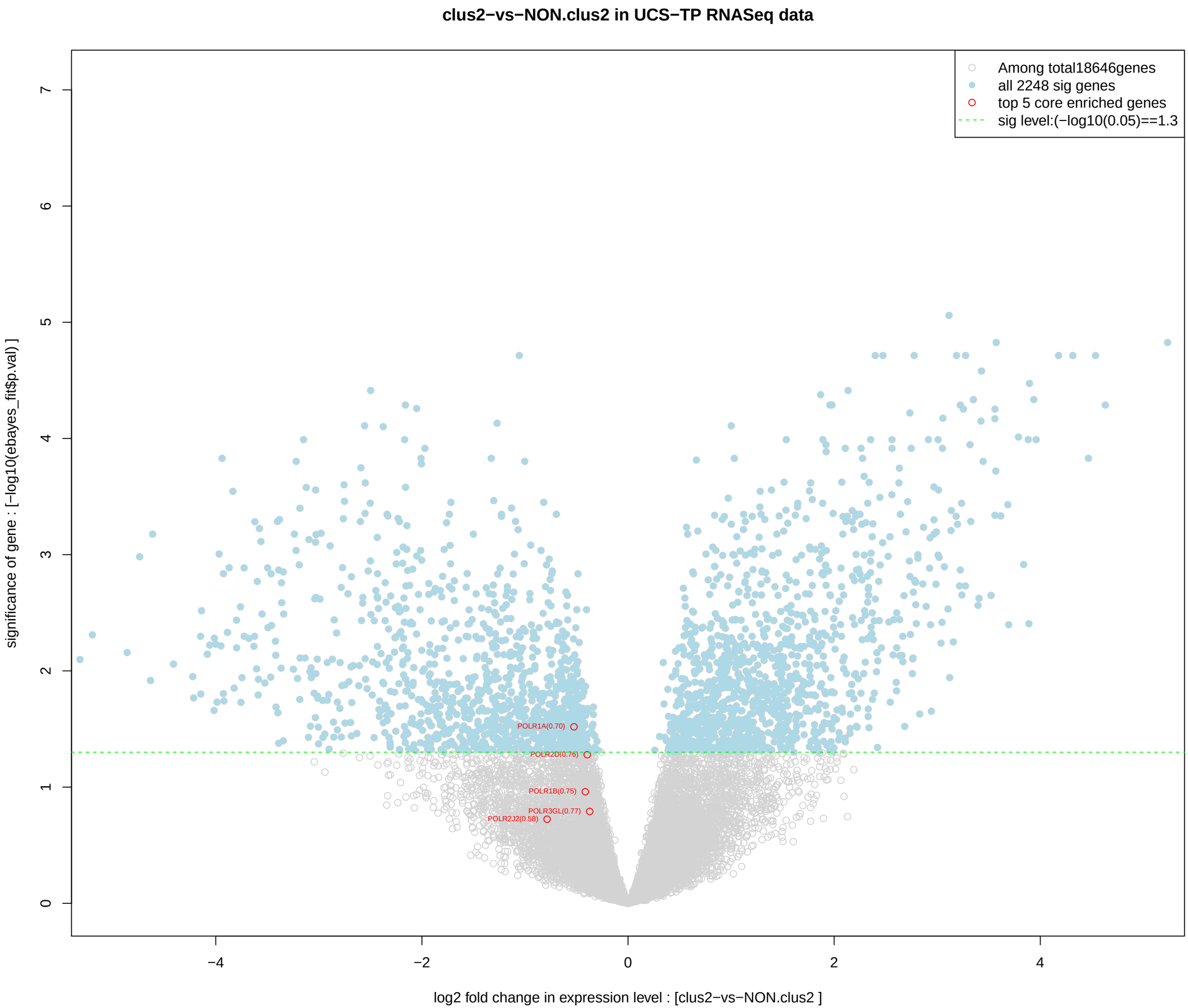
clus2−vs−NON.clus2 in UCS−TP RNASeq data
Among total18646genes
all 2248 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
POLR1A(0.70)
POLR2D(0.76)
POLR1B(0.75)
POLR3GL(0.77)
POLR2J2(0.58)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
1
2
3
4
5
6
7
−4
−2
0
2
4