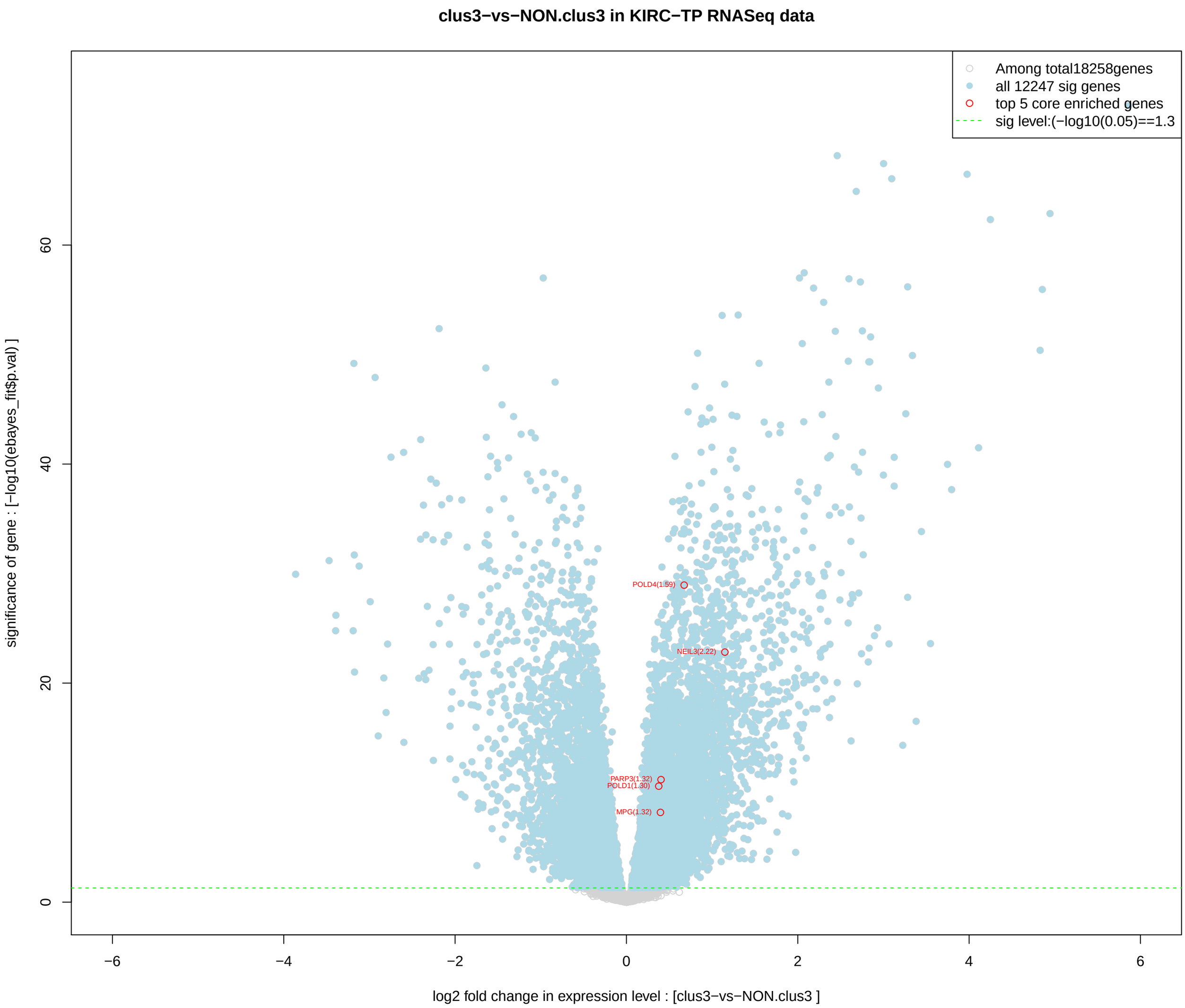
clus3−vs−NON.clus3 in KIRC−TP RNASeq data
Among total18258genes
all 12247 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
POLD4(1.59)
NEIL3(2.22)
PARP3(1.32)
POLD1(1.30)
MPG(1.32)
significance of gene : [−log10(ebayes_fit$p.val)]
0
20
40
60
−6
−4
−2
0
2
4
6
log2 fold change in expression level : [clus3−vs−NON.clus3]