clus4−vs−NON.clus4 in THYM−TP RNASeq data
Among total18283genes
all 7807 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ERCC8(0.67)
PCNA(0.59)
RFC5(0.60)
POLE2(0.60)
GTF2H2(0.57)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0
5
10
15
20
25
30
−5
0
5