clus3−vs−NON.clus3 in UCEC−TP RNASeq data
Among total18525genes
all 9969 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CDC25B(0.46)
ESPL1(0.50)
E2F1(0.50)
E2F2(0.55)
ORC1L(0.58)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
20
40
60
80
100
−6
−4
−2
0
2
4
6