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
ESPL1(0.50)
AURKA(0.54)
CDC25C(0.65)
SGOL1(0.66)
RPS6KA6(0.60)
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