clus2−vs−NON.clus2 in SARC−TP RNASeq data
Among total18166genes
all 12143 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
TFPI(0.14)
CFH(0.14)
PLAUR(0.19)
CFD(0.10)
C4BPB(0.21)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
20
40
60
80
−10
−5
0
5
10