clus2−vs−NON.clus2 in UCEC−TP RNASeq data

significance of gene : [−log10(ebayes_fit$p.val)]

- ○ Among total18525genes
- ● all 7958 sig genes
- ○ top 5 core enriched genes
- --- sig level:(−log10(0.05)==1.3

HERC5(0.44)
UBE2L6(0.62)
ISG15(0.65) NBD(0.77)
SAA1(0.49)

log2 fold change in expression level : [clus2−vs−NON.clus2]