
clus3−vs−NON.clus3 in UCS−TP RNASeq data
Among total18646genes
all 313 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
UGT2A3(8.39)
MGAM(4.22)
UGT2B10(2.48)
ENPP3(6.99)
UGT2B7(7.54)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
2
4
6
8
−6
−4
−2
0
2
4
6