

clus2-vs-NON.clus2 in UCS-TP RNASeq data

- Among total 18646 genes
- all 2530 sig genes
- top 5 core enriched genes
- - - sig level: $(-\log_{10}(0.05)) = 1.3$

significance of gene : $[-\log_{10}(\text{ebayes_fit\$p.val})]$

