clus3−vs−NON.clus3 in PAAD−TP RNASeq data

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

- ○ Among total18457genes
- ● all 8710 sig genes
- ○ top 5 core enriched genes
- --- sig level:(−log10(0.05)==1.3

PIPOX(3.63)

OGDHL(6.95)

log2 fold change in expression level : [clus3−vs−NON.clus3]