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

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

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

Among total17728genes
all 8561 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3

C2orf29(1.62)
PRKCD(1.53)
RPL21(1.58)
RPL36A(1.46)
RPL9(1.67)