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

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

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

Among total17976genes
all 6692 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3

ACAT1(0.67)
ALDH2(0.82)
AADAT(0.61)
AASDH(0.79)
ACAT2(0.73)