clus3−vs−NON.clus3 in CESC−TP RNASeq data
Among total18170genes
all 9458 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ACY3(9.74)
ABP1(15.23)
ALDH3B1(4.18)
HNMT(2.35)
DDC(3.34)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
10
20
30
40
50
−5
0
5