
clus2−vs−NON.clus2 in LIHC−TP RNASeq data
Among total17728genes
all 9501 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PCK2(3.17)
ACO1(1.82)
SDHD(1.56)
PCK1(6.06)
OGDHL(2.83)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
10
20
30
40
−6
−4
−2
0
2
4
6