
clus1−vs−NON.clus1 in LIHC−TP RNASeq data
Among total17728genes
all 8882 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CARD18(2.46)
CASP5(2.09)
CCL13(2.76)
IL18(2.22)
CXCL1(4.53)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
5
10
15
20
25
30
−4
−2
0
2
4