
clus1−vs−NON.clus1 in THCA−TP RNASeq data
Among total17971genes
all 12731 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
BCL2L1(1.79)
PMAIP1(3.30)
BID(1.85)
BBC3(1.49)
BIK(1.56)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
10
20
30
40
50
60
−4
−2
0
2
4