
clus2−vs−NON.clus2 in PRAD−TP RNASeq data
Among total18256genes
all 12273 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
NUP210(1.55)
NUP54(1.22)
RANBP2(1.34)
SMN1(1.41)
NUP155(1.24)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
10
20
30
40
50
60
70
−4
−2
0
2
4