clus1−vs−NON.clus1 in CESC−TP RNASeq data
Among total18170genes
all 6590 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
LEF1(2.61)
AKT3(1.95)
AXIN2(2.40)
CTNNA2(1.88)
TCF7L1(1.74)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
5
10
15
20
25
−4
−2
0
2
4