clus1−vs−NON.clus1 in TGCT−TP RNASeq data
Among total19051genes
all 14401 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
FZD7(0.07)
FZD2(0.14)
CHP2(0.07)
CER1(0.02)
WIF1(0.06)
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
70
−5
0
5