clus3−vs−NON.clus3 in TGCT−TP RNASeq data
Among total19051genes
all 14195 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
TGFB2(18.00)
ARNT2(9.68)
JUN(5.58)
HGF(8.18)
FIGF(3.61)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
10
20
30
40
50
−5
0
5