clus4−vs−NON.clus4 in THYM−TP RNASeq data
Among total18283genes
all 7807 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
ICOSLG(4.99)
TNFSF13B(4.65)
CD80(4.29)
AICDA(3.62)
TNFRSF13B(7.09)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0
5
10
15
20
25
30
−5
0
5