clus2−vs−NON.clus2 in MESO−TP RNASeq data
Among total18187genes
all 2388 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CREB3L4(1.47)
DYNC1I1(1.93)
DYNC2H1(1.61)
CREB5(1.40)
AVPR2(1.35)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
2
4
6
8
10
−4
−2
0
2
4