clus2−vs−NON.clus2 in MESO−TP RNASeq data
Among total18193genes
all 2197 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PLA2G7(0.27)
RELN(0.27)
LRP8(0.59)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
2
4
6
8
−4
−2
0
2
4