clus3−vs−NON.clus3 in PAAD−TP RNASeq data
Among total18457genes
all 8710 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CACNA1A(6.74)
GCK(7.97)
CACNA1B(8.58)
ABCC8(16.65)
KCNJ11(3.96)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
5
10
15
20
25
−4
−2
0
2
4