clus1−vs−NON.clus1 in LAML−TB RNASeq data
Among total17259genes
all 8093 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PRKG2(5.89)
PLCB4(7.03)
EGFR(2.66)
GJA1(6.33)
GNAI1(4.53)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
5
10
15
20
−4
−2
0
2
4