clus2−vs−NON.clus2 in PAAD−TP RNASeq data
Among total18457genes
all 1688 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
GSTA2(18.26)
ANPEP(4.28)
GSTA1(4.69)
GGT1(2.84)
GGT6(3.37)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
30
25
20
15
10
5
0
−5
0
5