clus4−vs−NON.clus4 in LGG−TP RNASeq data
Among total18315genes
all 13207 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
HS6ST3(12.86)
HS3ST2(5.61)
NDST3(5.92)
HS3ST4(4.97)
GPC5(4.91)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0
20
40
60
80
100
120
−4
−2
0
2
4