clus5−vs−NON.clus5 in KICH−TP RNASeq data
Among total17824genes
all 348 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PLAT(0.41)
IL6(0.29)
PLAU(0.42)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus5−vs−NON.clus5]
0
1
2
3
4
5
6
−4
−2
0
2
4