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

significance of gene : [−log10(ebayes_fit$p.val)]

log2 fold change in expression level : [clus3−vs−NON.clus3]

GRIA2(10.80)
CAMK2B(9.13)
PPP1R1A(12.00)
GRM1(2.99)
GRIA1(2.32)