clus3−vs−NON.clus3 in HNSC−TP RNASeq data

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

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

Legend:
- ○ Among total18361genes
- ● all 13069 sig genes
- ○ top 5 core enriched genes
- − − sig level:(−log10(0.05)==1.3

Labeled genes: CACNB1(0.35), CACNA2D1(0.19), CACNA2D2(0.46), CACNG6(0.27), SGCG(0.25)