clus5−vs−NON.clus5 in LGG−TP RNASeq data

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

log2 fold change in expression level : [clus5−vs−NON.clus5]

Legend:
- ○ Among total18315genes
- ● all 5763 sig genes
- ○ top 5 core enriched genes
- - - - sig level:(−log10(0.05)==1.3

Labeled points:
- PPAP2C(5.47)
- ENPP2(5.78)
- PLD1(3.19)
- LPCAT2(1.88)
- PLA2G7(1.81)