clus2−vs−NON.clus2 in CHOL−TP RNASeq data

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

log2 fold change in expression level : [clus2−vs−NON.clus2]

Legend:
- ○ Among total18113genes
- ● all 489 sig genes
- ○ top 5 core enriched genes
- --- sig level:(−log10(0.05)==1.3

Labeled genes: IKZF3(0.18), PIK3CG(0.32), IL2RA(0.24), FASLG(0.43), IL2RB(0.41)