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

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

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

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

Labeled points:
- IGF1(0.17)
- PIK3R5(0.34)
- PIK3CG(0.32)
- VEGFC(0.53)
- PIK3CD(0.49)