clus3−vs−NON.clus3 in SKCM−TM RNASeq data

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

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

Legend:
- ○ Among total18048genes
- ● all 10565 sig genes
- ○ top 5 core enriched genes
- −−− sig level:(−log10(0.05)==1.3

Labeled points: PIK3CD(0.27), CD79A(0.19), CD19(0.22), PRKCB(0.34), CD79B(0.32)