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

Legend:
- ○ Among total17717genes
- ● all 1056 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 : [clus3−vs−NON.clus3]

Labeled points: TCF7(0.13), SMO(0.37), WNT3(0.38), WNT4(0.25), WNT10B(0.35)