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

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

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

Legend:
- ○ Among total18307genes
- ● all 14306 sig genes
- ○ top 5 core enriched genes
- −−− sig level:(−log10(0.05)==1.3

Labeled points:
- HLA−DMB(0.33)
- CD80(0.43)
- IL10(0.38)
- HLA−DOA(0.36)
- HLA−DQA1(0.26)