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

Among total18197genes
all 14911 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3

CYP4A11(71.48)
CD36(11.32)
CYP4A22(3.09)
ANGPTL4(18.06)
FABP7(53.02)

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

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