clus2−vs−NON.clus2 in STES−TP RNASeq data
Among total18748genes
all 10615 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CYP3A5(0.04)
CYP2C19(0.11)
PLA2G2A(0.04)
PLA2G10(0.12)
CYP2C9(0.13)
significance of gene : [−log10(ebayes_fit$p.val)]
0
10
20
30
−5
0
5
log2 fold change in expression level : [clus2−vs−NON.clus2]