clus2−vs−NON.clus2 in LUAD−TP RNASeq data
Among total18300genes
all 10864 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
EPHX2(0.50)
PXMP4(0.52)
ACSL5(0.56)
DAO(0.79)
BAAT(0.58)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
10
20
30
40
−4
−2
0
2
4