clus1−vs−NON.clus1 in PRAD−TP RNASeq data
Among total18256genes
all 11515 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CYP3A5(7.05)
ALDH3A1(4.63)
AOX1(3.67)
CYP3A7(2.69)
FMO2(2.89)
significance of gene : [−log10(ebayes_fit$p.val) ]
0
10
20
30
40
50
60
−4
−2
0
2
4
log2 fold change in expression level : [clus1−vs−NON.clus1 ]