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
ITGB4(3.22)
ITGB6(3.92)
ITGB8(2.50)
ITGB3(2.40)
CACNA2D3(2.16)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
10
20
30
40
50
60
−4
−2
0
2
4