clus3−vs−NON.clus3 in LUAD−TP RNASeq data
Among total18300genes
all 10473 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
MET(0.24)
EGFR(0.42)
PDGFD(0.62)
FGF10(0.77)
FGF5(0.63)
significance of gene : [−log10(ebayes_fit$p.val) ]
log2 fold change in expression level : [clus3−vs−NON.clus3 ]
0
10
20
30
40
50
−6
−4
−2
0
2
4
6