clus3−vs−NON.clus3 in HNSC−TP RNASeq data
Among total18361genes
all 13069 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PLD2(1.93)
PLA2G2F(7.70)
PLA2G4E(7.47)
JMJD7−PLA2G4B(1.62)
PLA2G3(1.99)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
10
20
30
40
50
−4
−2
0
2
4