

200

150

100

20

0.0

0.2

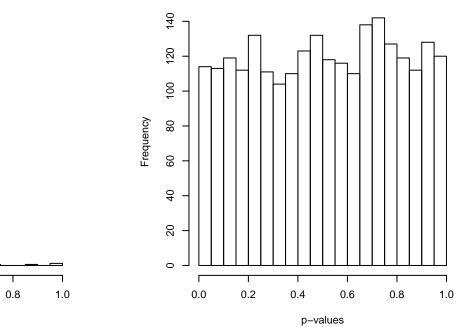
Frequency

pval of non-DE genes,deseq2 method

0.6

8.0

1.0

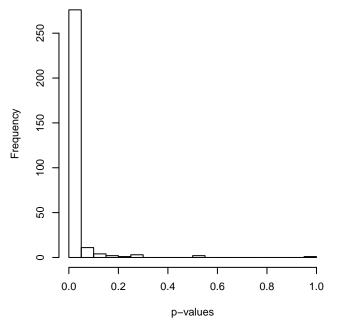


0.0

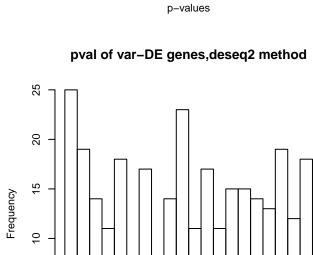
0.2

0.4

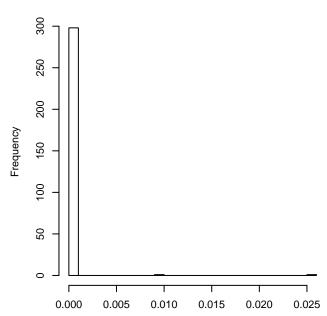
p-values



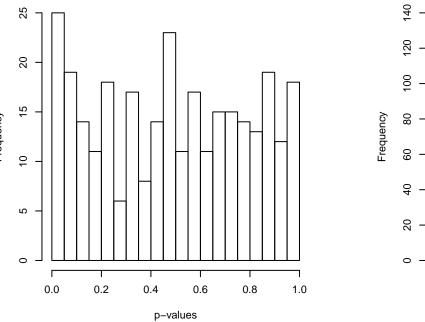
pval of mean-DE genes,deseq2 method



0.4



p-values



0.6

