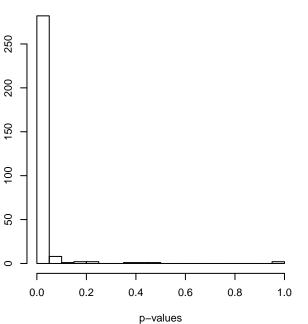
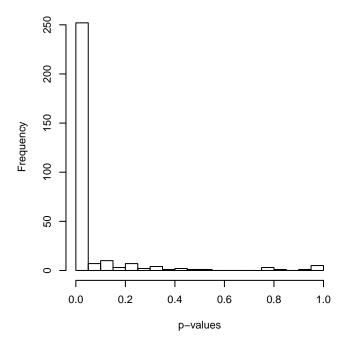
200

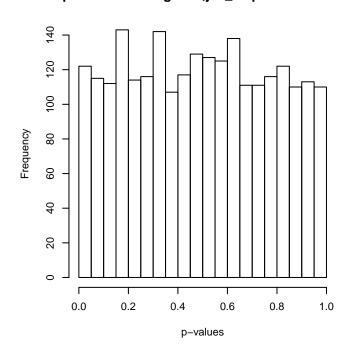
150

100

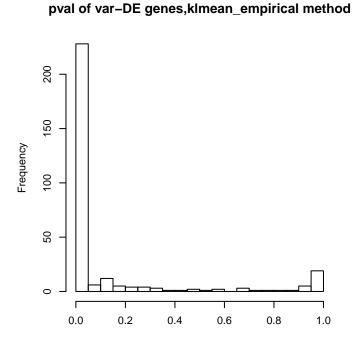
Frequency



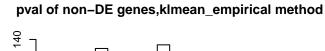


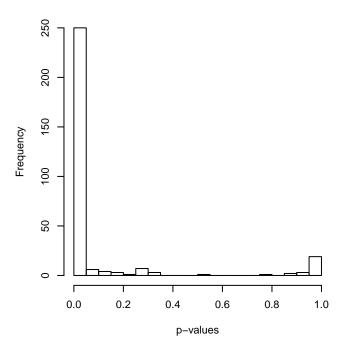


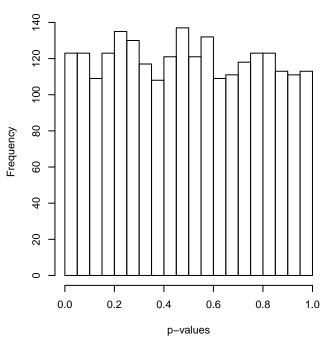
pval of mean-DE genes,klmean\_empirical method



p-values







250

200

150

100

50

20

15

10

2

0.0

0.2

0.4

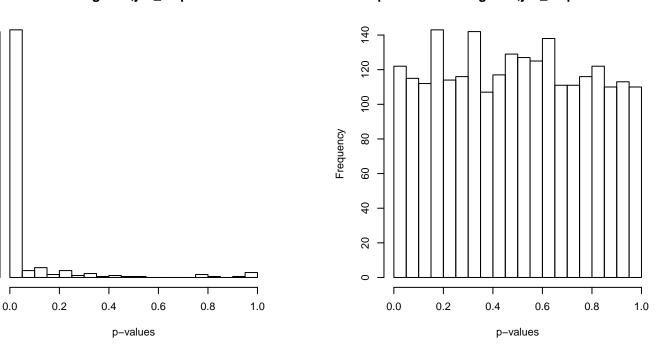
p-values

0.6

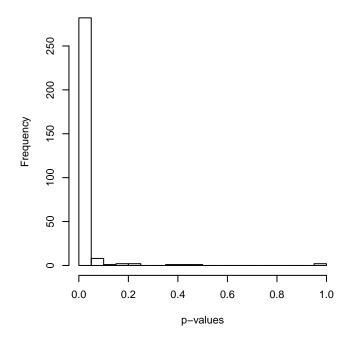
8.0

Frequency

Frequency



1.0



pval of mean-DE genes,deseq2 method

300

250

200

150

100

20

Frequency

