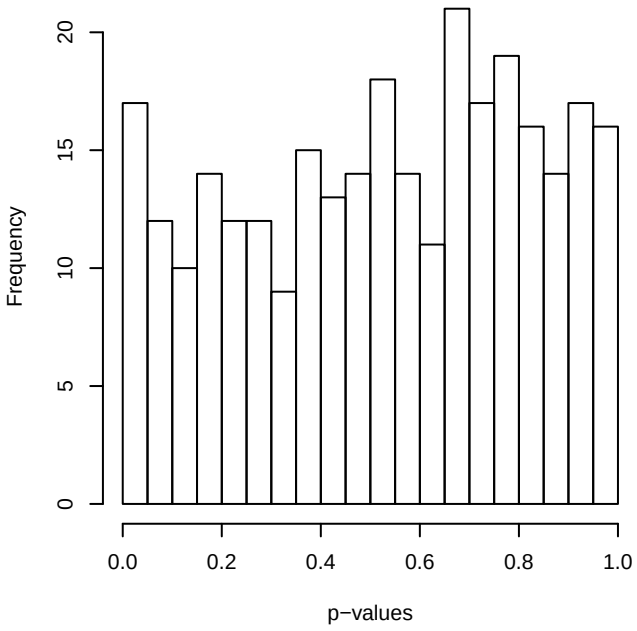
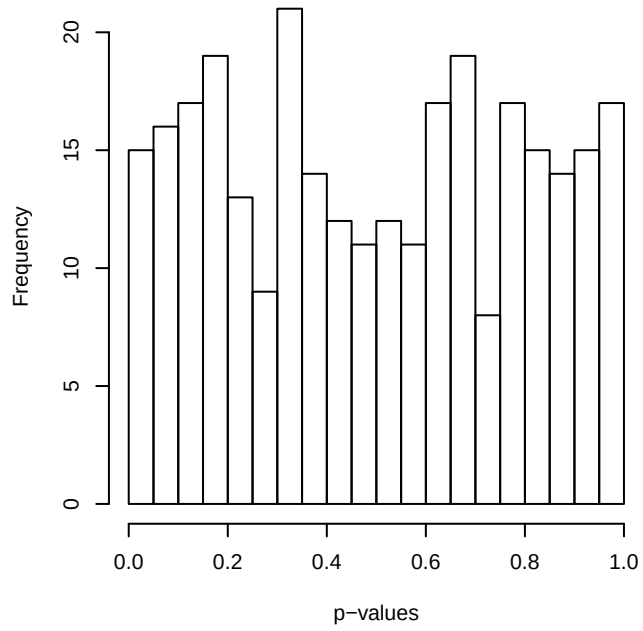
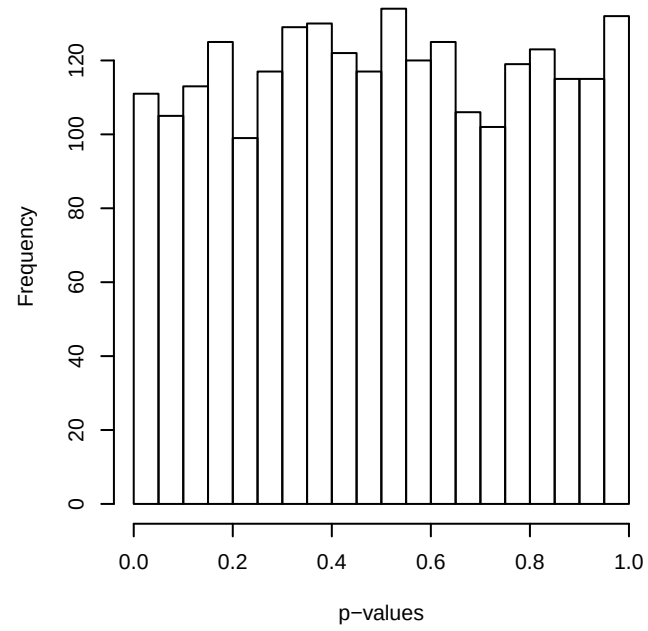
pval of mean−DE genes,jsd_zinb method



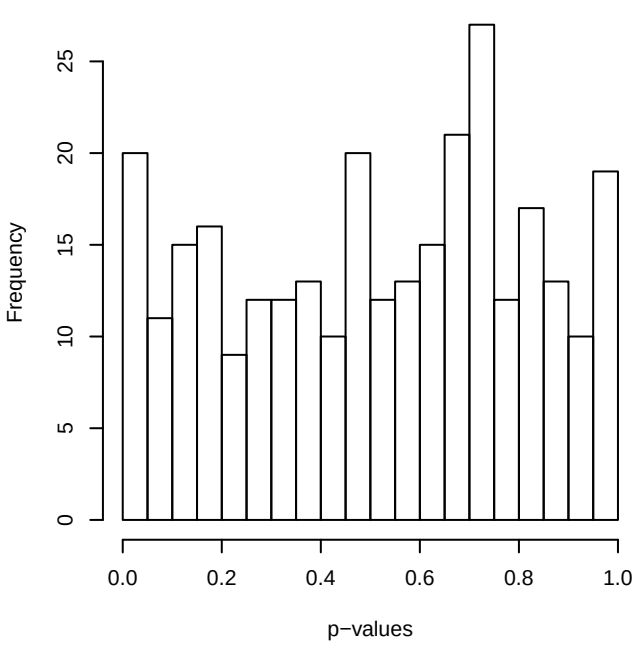
pval of var−DE genes,jsd_zinb method



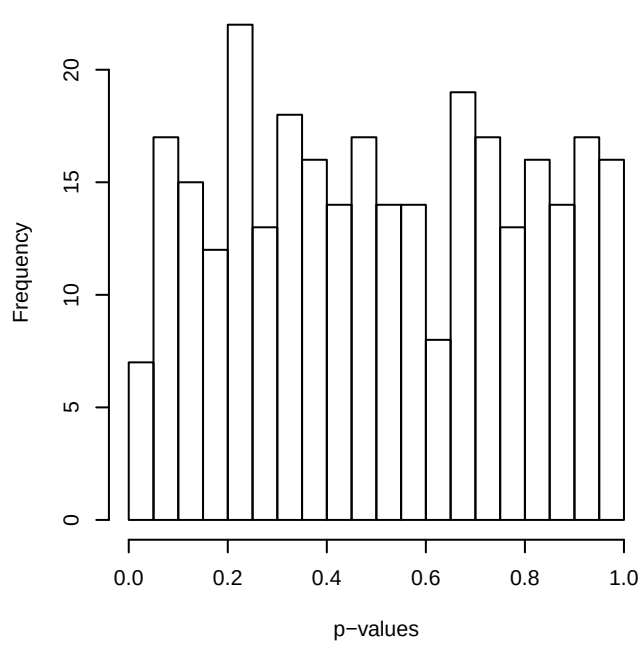
pval of non−DE genes,jsd_zinb method



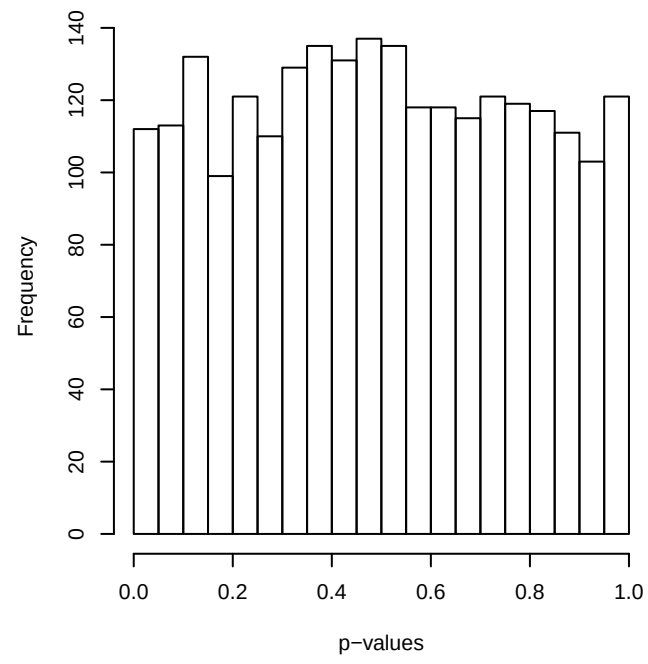
pval of mean−DE genes,jsd_empirical method



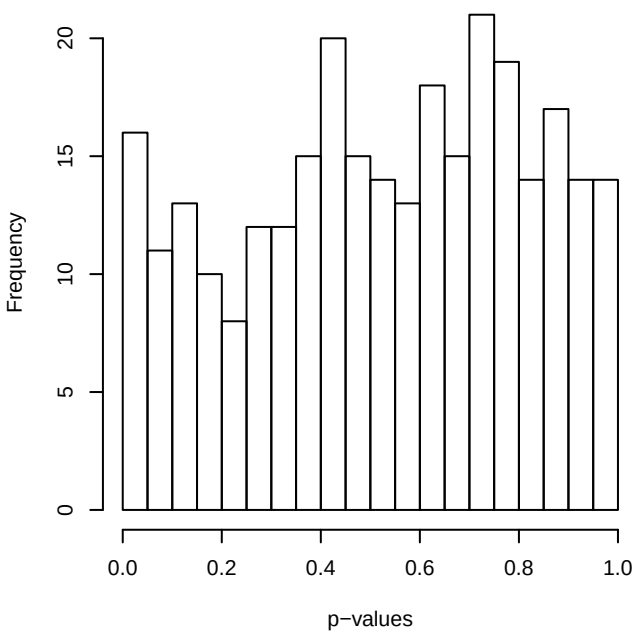
pval of var−DE genes,jsd_empirical method



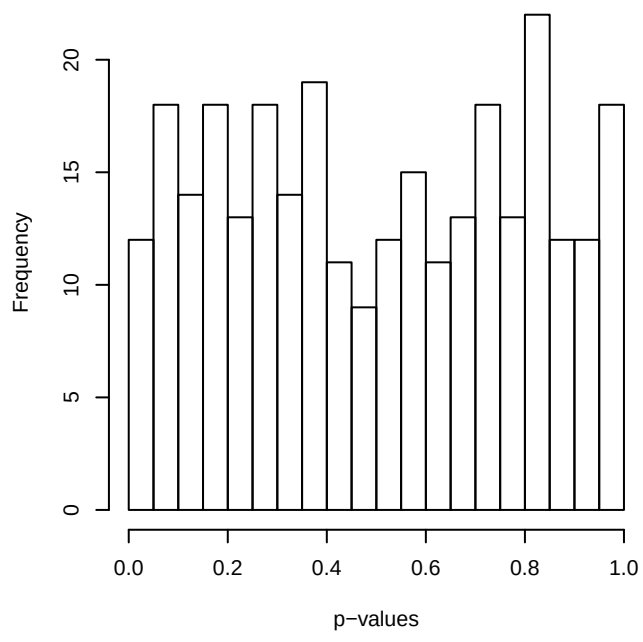
pval of non−DE genes,jsd_empirical method



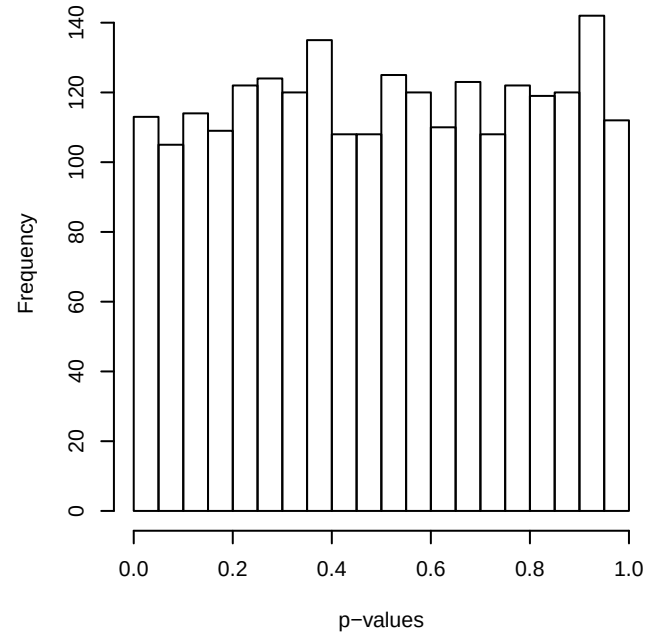
pval of mean−DE genes,klmean_zinb method



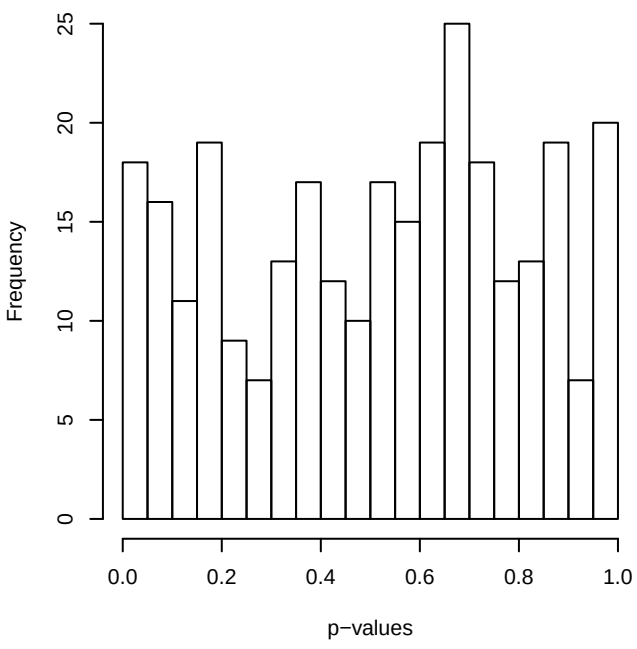
pval of var−DE genes,klmean_zinb method



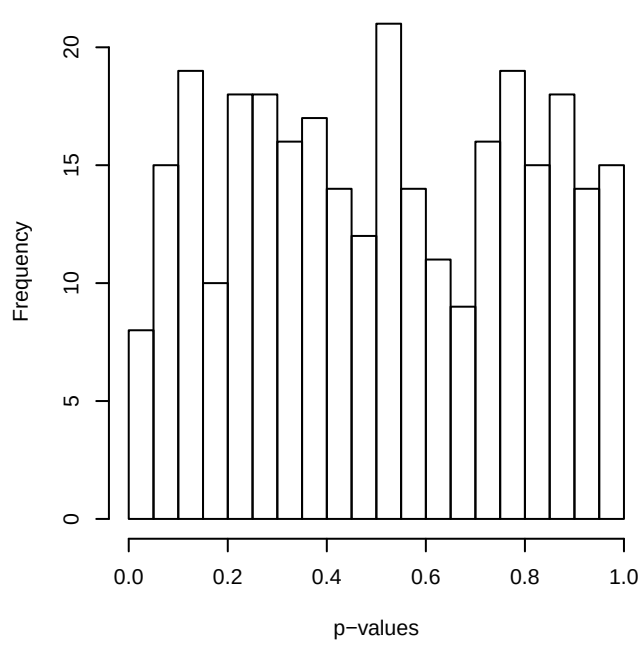
pval of non−DE genes,klmean_zinb method



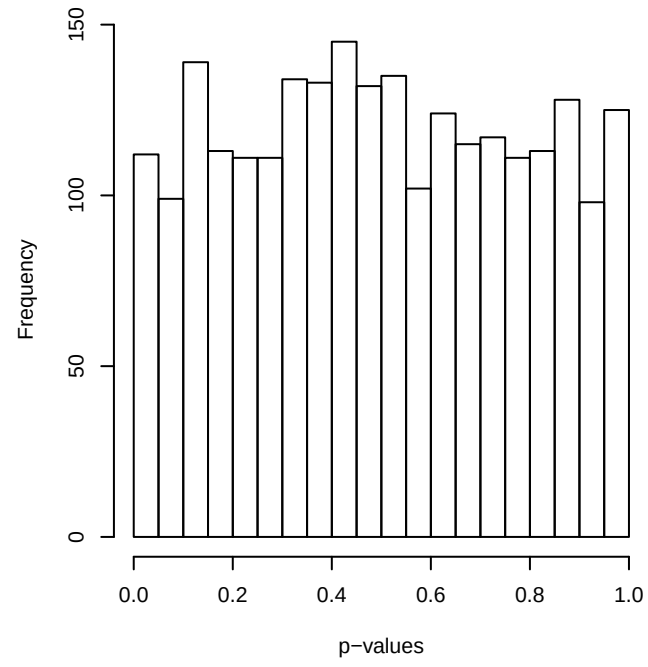
pval of mean−DE genes,klmean_empirical method

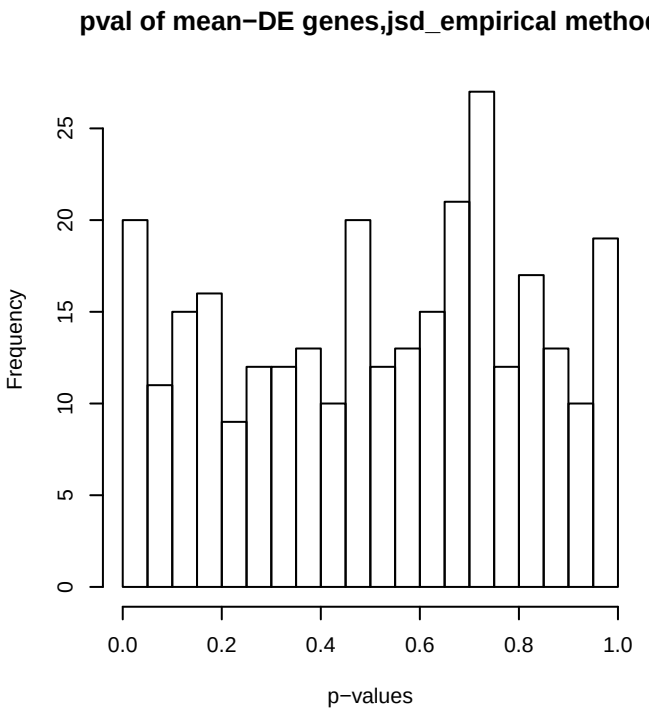


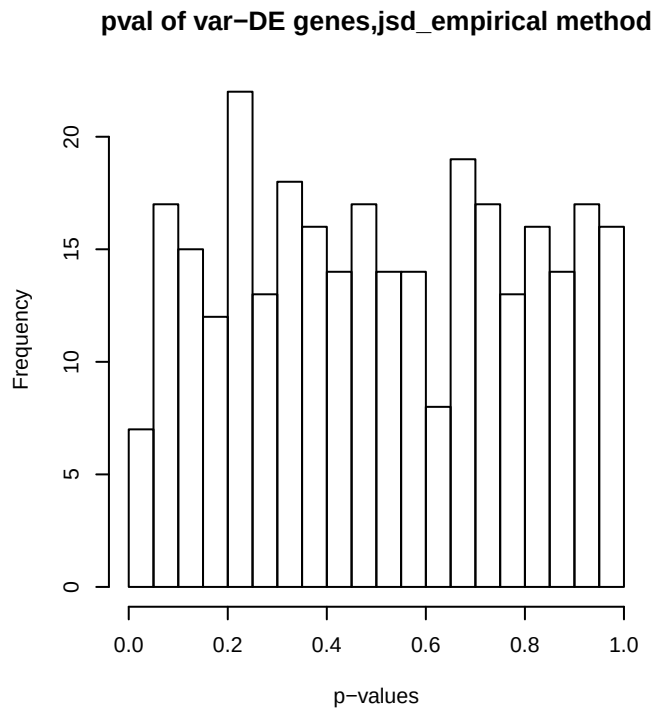
pval of var−DE genes,klmean_empirical method

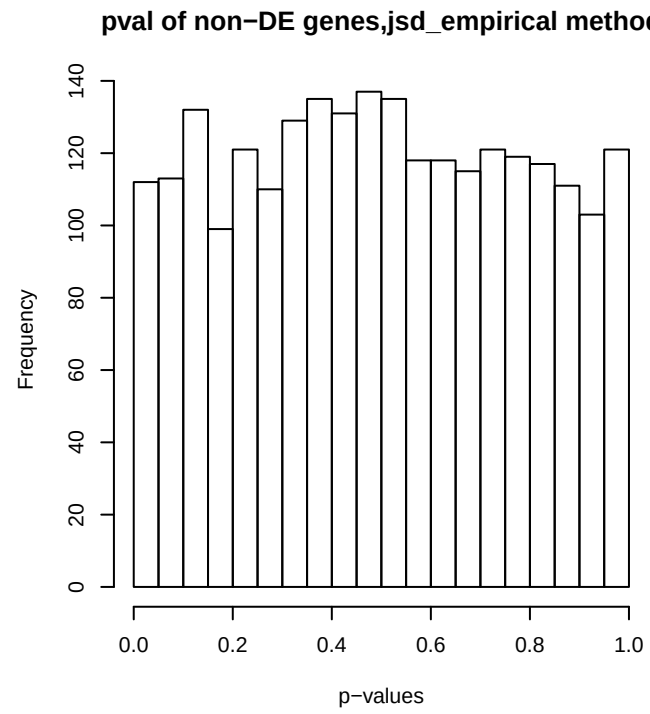


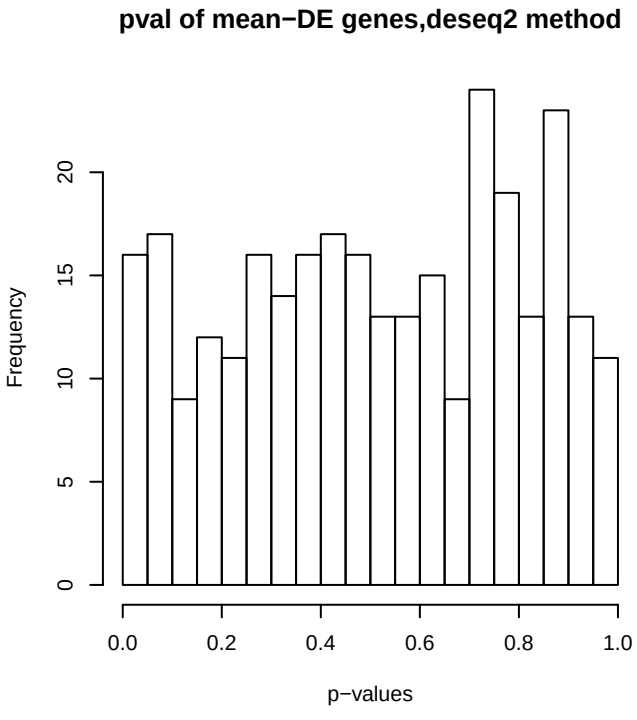
pval of non−DE genes,klmean_empirical method

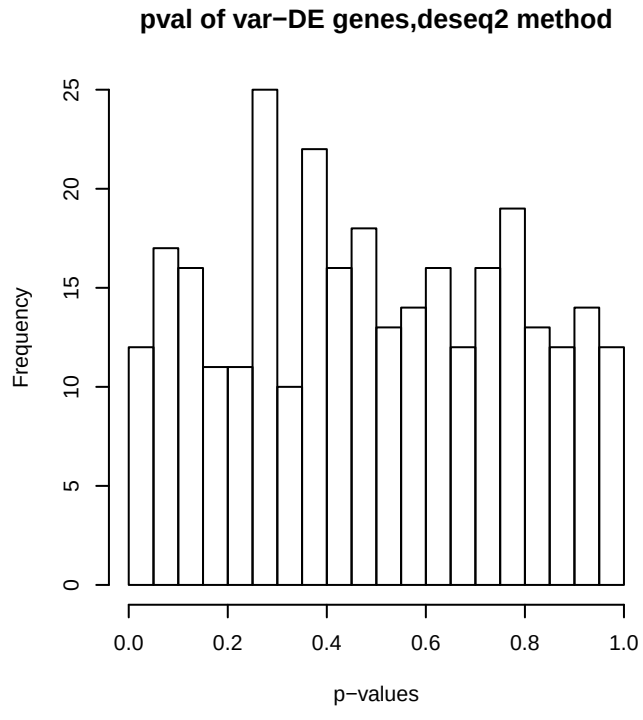


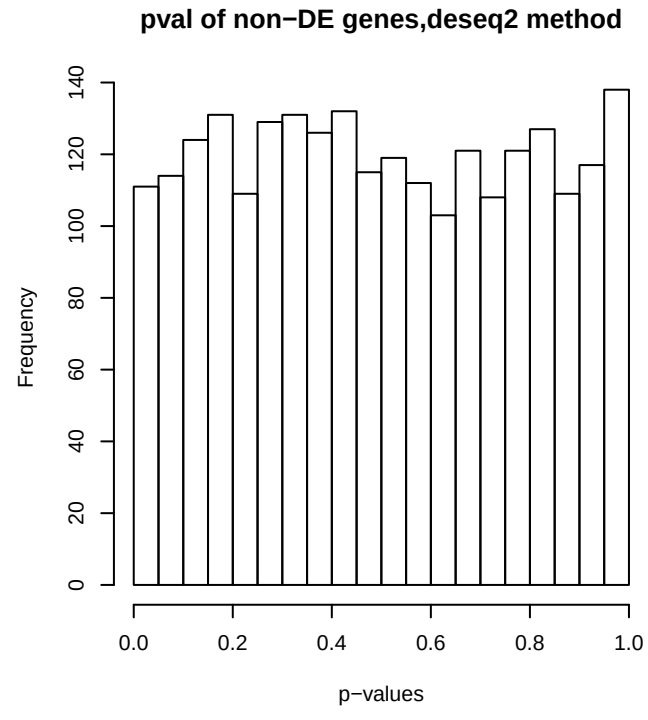

pval of mean−DE genes,jsd_empirical method
Frequency
p−values
0
5
10
15
20
25
0.0
0.2
0.4
0.6
0.8
1.0

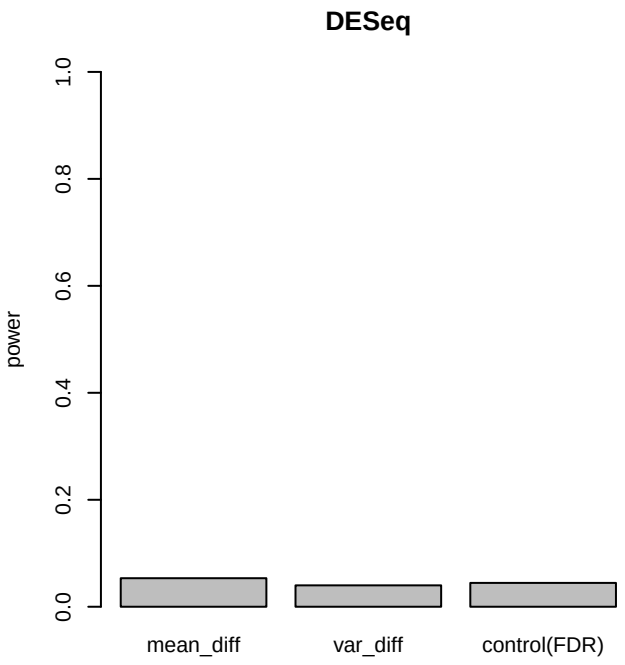

pval of var−DE genes,jsd_empirical method
Frequency
p−values
0
5
10
15
20
0.0
0.2
0.4
0.6
0.8
1.0

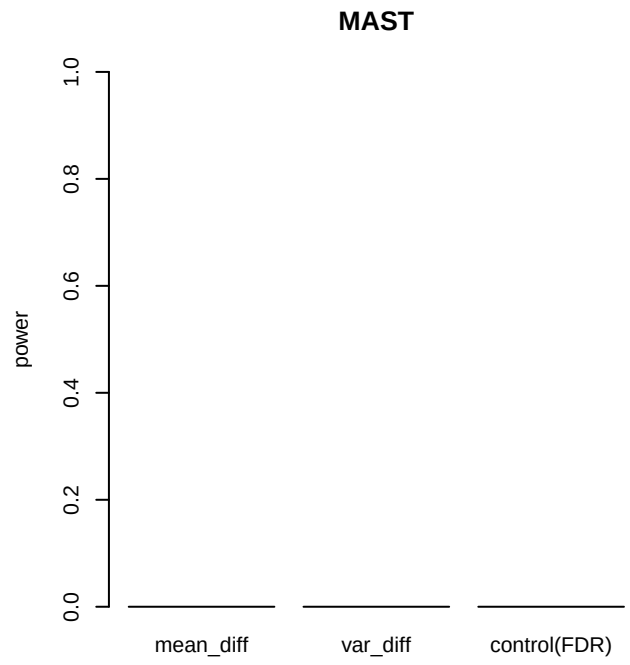

pval of non−DE genes,jsd_empirical method
Frequency
p−values
0
20
40
60
80
100
120
140
0.0
0.2
0.4
0.6
0.8
1.0

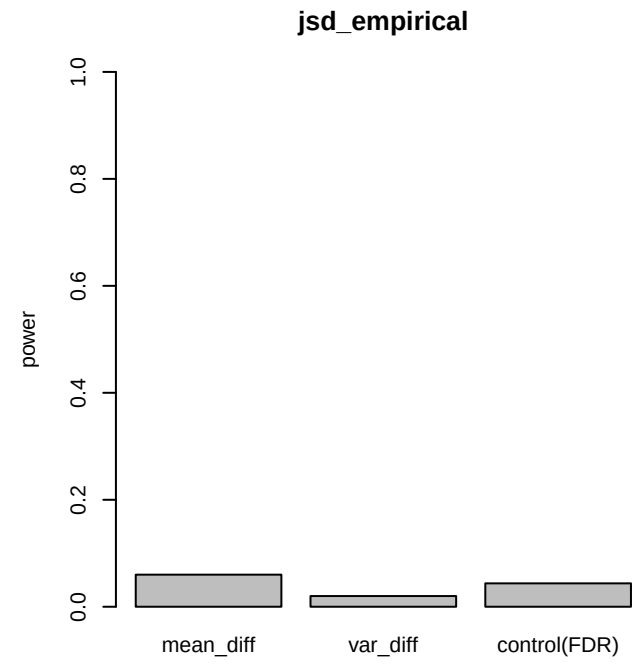

pval of mean−DE genes,deseq2 method
Frequency
p−values
0
5
10
15
20
0.0
0.2
0.4
0.6
0.8
1.0

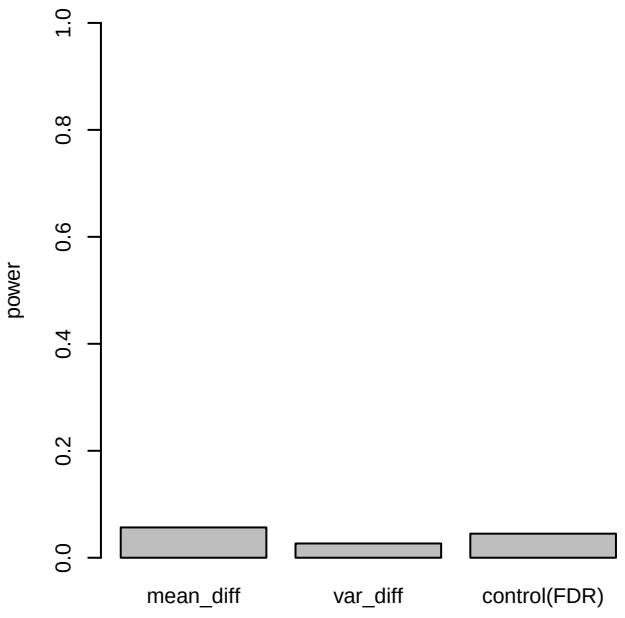

pval of var−DE genes,deseq2 method
Frequency
p−values
0
5
10
15
20
25
0.0
0.2
0.4
0.6
0.8
1.0

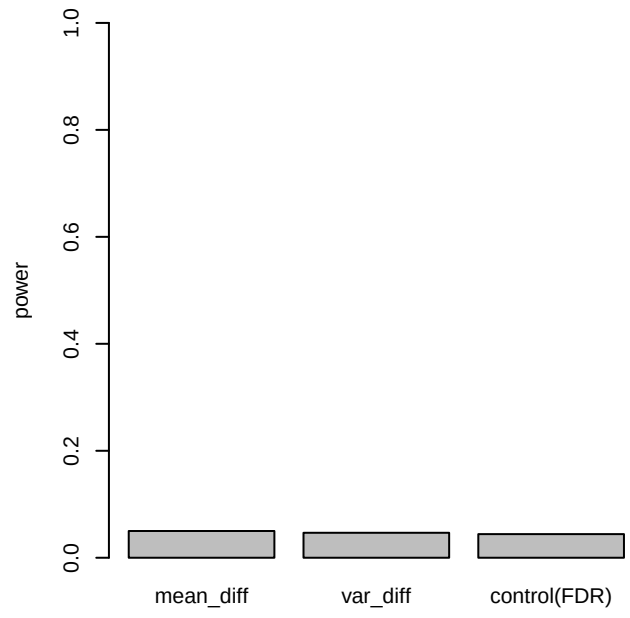

pval of non−DE genes,deseq2 method
Frequency
p−values
0
20
40
60
80
100
120
140
0.0
0.2
0.4
0.6
0.8
1.0

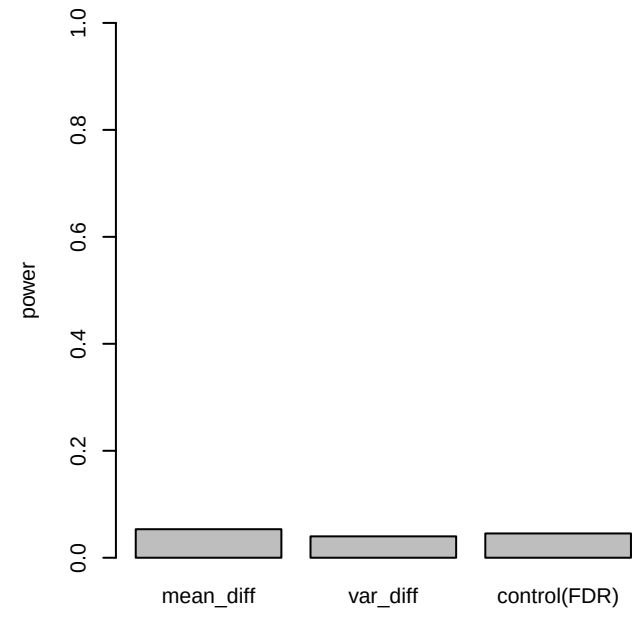

DESeq
power
0.0
0.2
0.4
0.6
0.8
1.0
mean_diff
var_diff
control(FDR)

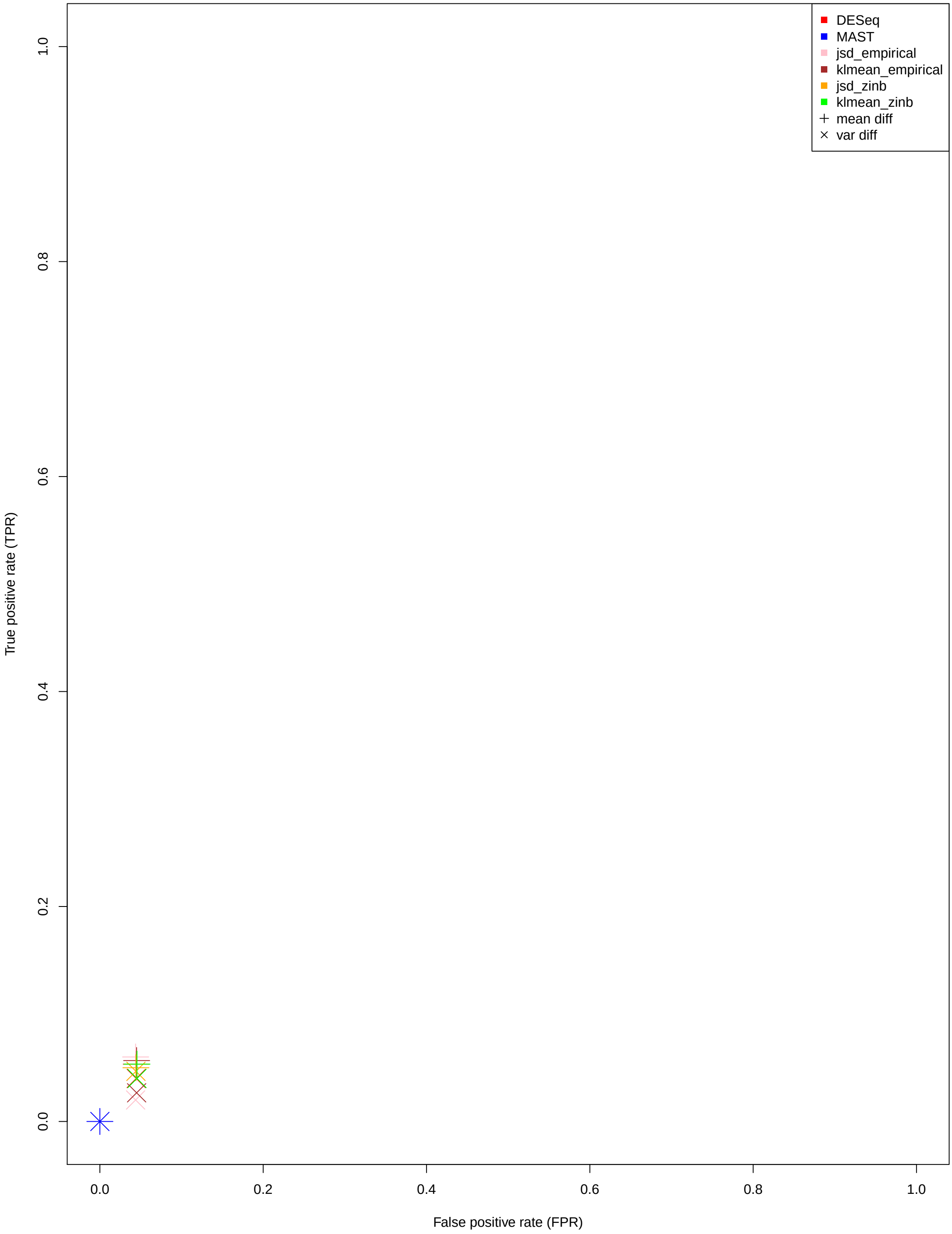

MAST
power
0.0
0.2
0.4
0.6
0.8
1.0
mean_diff
var_diff
control(FDR)


jsd_empirical
power
0.0
0.2
0.4
0.6
0.8
1.0
mean_diff
var_diff
control(FDR)


klmean_empirical
power
0.0
0.2
0.4
0.6
0.8
1.0
mean_diff
var_diff
control(FDR)


jsd_zinb
power
0.0
0.2
0.4
0.6
0.8
1.0
mean_diff
var_diff
control(FDR)


klmean_zinb
power
0.0
0.2
0.4
0.6
0.8
1.0
mean_diff
var_diff
control(FDR)


DESeq
MAST
jsd_empirical
klmean_empirical
jsd_zinb
klmean_zinb
mean diff
var diff
True positive rate (TPR)
False positive rate (FPR)
0.0
0.2
0.4
0.6
0.8
1.0