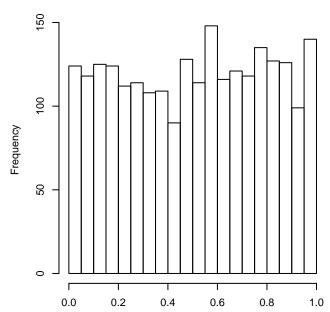
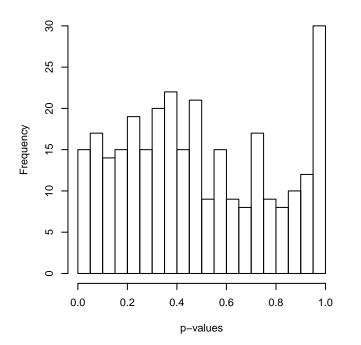


pval of non-DE genes,jsd_empirical method



Ledneucy 12 0.0 0.2 0.4 0.6 0.8 1.0 p-values

pval of var-DE genes,jsd_empirical method



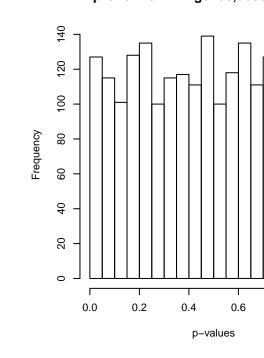
pval of mean-DE genes,jsd_empirical method

pval of non-DE genes,deseq2 method

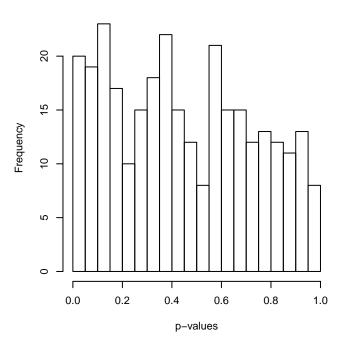
1.0

8.0

p-values



pval of var-DE genes,deseq2 method



pval of mean-DE genes,deseq2 method

