STAT572 - Homework Assignment 4

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In-class 4: Monte Carlo Hypothesis Testing for $\hat{\sigma}$

```
% In-class; hypothesis test on the standard deviation parameter
load mcdata; n = length(mcdata);
% Population sigma is known.
sigma = 7.8; sigxbar = sigma/sqrt(n);
% Get the observed value of the test statistic.
Tobs = (n-1)*var(mcdata)/sigma^2;
M = 1000; % Number of Monte Carlo trials
Tm = zeros(1,M);
% Start the simulation.
for i = 1:M
\% Generate a rand sample under H_O where n is the sample size.
xs = sigma*randn(1,n) + mean(mcdata);
Tm(i) = (n-1)*var(xs)/sigma^2;
end
% This is a upper-tail test, so it is the 1-alpha quantile.
alpha = 0.05; cv = csquantiles(Tm,1-alpha);
% Plotting in-class
figure(2); hist(Tm)
set(get(gca,'child'),'FaceColor',[.9 .9 .9],'EdgeColor','black');
hold on
plot(cv,0,'*',Tobs,0,'bo')
legend('frequency','critical value','observed value')
hold off
% Calculating the p-value
g = [];
for i=1:length(Tm)
    if Tm(i) >= Tobs
        g = [g,Tm(i)];
    end
end
pval=length(g)/length(Tm);
```

The p-value calculated by the above algorithm is 0.2600. Given that we chose $\alpha = 0.05$, we fail to reject $H_o: \sigma = 7.8$ in favor of the alternative $H_a: \sigma > 7.8$.

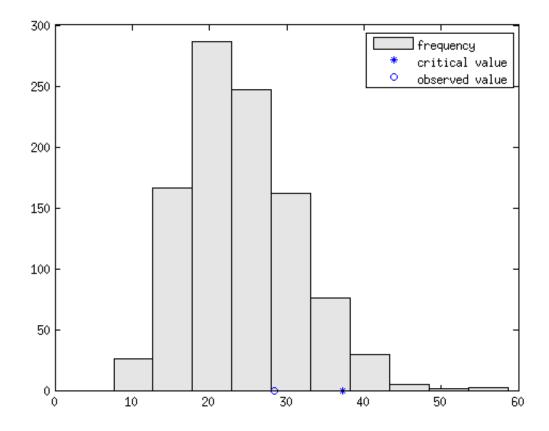


Figure 1:

In-class 5: Bootstrap Exercise on the Mean and 10% Trimmed Mean

```
% generate 200 rs from chi-square df=f
X = chi2rnd(5,1,200);

% PART A
n = length(X); % sample size
B = 400; % number of bootstrap replicates
% Get the value of the statistic of interest.
theta = mean(X);

% Use unidrnd to get the indices to the resamples.
inds = unidrnd(n,n,B);
% Extract these from the data.
xboot = X(inds);
thetab = mean(xboot); % get the mean for each column using seb = std(thetab);
```

```
biasEst = mean(thetab)-theta;
figure(1)
hist(thetab)
set(get(gca,'child'),'FaceColor',[.9 .9 .9],'EdgeColor','black');
title('Boostrapped Sample Mean Histogram')
ylabel('Frequency'); xlabel('Sample mean')
% PART B
thetat = trimmean(X,10); % use MATLAB trimmed mean function to estimate
thetatb = trimmean(xboot, 10); % generate boostrap trimmed mean
sebt = std(thetatb); % calculate the boostrap standard error
biastEst = mean(thetatb)-thetat; % estimate the bias
figure(2)
hist(thetatb)
set(get(gca,'child'),'FaceColor',[.9 .9 .9],'EdgeColor','black');
title('Boostrapped Trimmed Mean Histogram')
ylabel('Frequency'); xlabel('Trimmed mean')
par={'Sample Mean';'Trimmed Mean'}; Value = [mean(thetab); mean(thetatb)];
StdError = [seb; sebt]; Bias = [biasEst; biastEst];
T=table(Value,StdError,Bias,'RowNames',par); disp(T)
```

Implementing the above code we estimate the parameter, standard error and bias. This information is presented in figure 2. The histograms are presented in figures 3 and 4.



Figure 2: Mean and Trimmed Mean results.

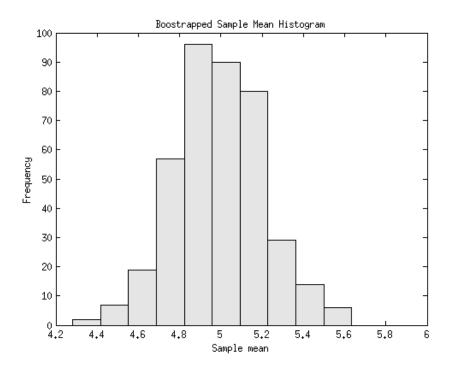


Figure 3: Frequency Histogram of the Bootstrapped Mean Parameter

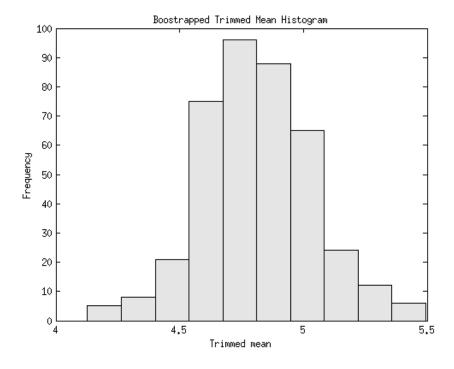


Figure 4: Frequency Histogram of the Bootstrapped 10% Trimmed Mean Parameter

```
% Based on Example 7.3 - Matinez
% Get several values for the mean under the alternative
% hypothesis. Note that we are getting some values
% below the null hypothesis.
mualt = 40:60;
% Note the critical value:
cv = 1.645;
\% Note the standard deviation for x-bar:
sig = 1.5;
\% It's easier to use the unstandardized version,
% so convert:
ct = cv*1.5 + 45;
\% Get a vector of ct values that is
% the same size as mualt.
ctv = ct*ones(size(mualt));
% Now get the probabilities to the left of this value.
% These are the probabilities of the Type II error.
beta = normcdf(ctv,mualt,sig);
% Plot probablity of a type II error vs. true mean
% under the alternative hypothesis
plot(mualt,beta);
xlabel('True Mean \mu')
ylabel('Probability of Type II Error')
axis([40 60 0 1.1])
```

In figure 5 below we see the relationship between the probability of a Type II Error and different values of the mean under the alternative hypothesis. We see precisely the opposite relationship between the power of the test, i.e. $1 - \beta$, and the means under the alternative which is shown in example 7.3 in the book (Martinez). This expected because of the definition Power= $1 - \beta$ and probability of a Type II Error= β .

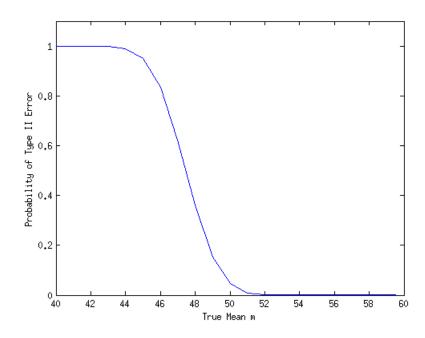


Figure 5: As the true mean of the population under the alternative hypothesis gets larger the probability of making a type II error goes to zero. This is precisely the opposite of the figure presented in example 7.3 in the book (Martinez).

```
% Based on example 7.3
% Generate the mean under the alternative
mualt = 40:60;
% Note the critical value:
cv = 1.645;
\% Note the standard deviation for x-bar:
sig=15./sqrt([50,100,200]);
% It's easier to use the unstandardized version,
ct = cv.*sig + 45;
% Get a vector of ct values that is
% the same size as mualt.
one = ones(size(mualt));
ctv = [ct(1)*one;ct(2)*one;ct(3)*one];
% Now get the probabilities to the left of this value.
% These are the probabilities of the Type II error.
beta50 = normcdf(ctv(1,:),mualt,sig(1));
beta100 = normcdf(ctv(2,:),mualt,sig(2));
beta200 = normcdf(ctv(3,:),mualt,sig(3));
```

```
% To get the power: 1-beta
pow50 = 1 - beta50;
pow100 = 1 - beta100;
pow200 = 1 - beta200;

plot(mualt,pow50,'r-o');
hold on
plot(mualt,pow100,'-*');
hold on
plot(mualt,pow200,'--');
xlabel('True Mean \mu')
ylabel('Power')
axis([40 60 0 1.1])
legend('Power for n=50','Power for n=100','Power for n=200',2)
hold off
```

In figure 6 we see three different power curves for three different sample sizes, n = 50, n = 100 and n = 200. Notice that the larger the sample size the greater the power of the test. For example, for every possible value of μ under the alternative, the curve corresponding to n = 200 is above the other curves. Similarly, the curve that corresponds to n = 100 is above the curve corresponding to n = 50. This is expected given that the standard error of the sample mean is $\frac{\sigma}{\sqrt{n}}$. This tells us that the larger the sample size n, the smaller the standard error of the parameter estimate.

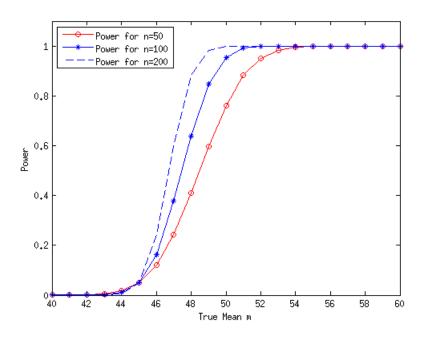


Figure 6: Power of the test for n = 50, 100, 200

Exercise 7.5

```
% Two-tailed test for Example 7.6
load mcdata
n = length(mcdata);
% Population sigma is known.
sigma = 7.8;
sigxbar = sigma/sqrt(n);
% Get the observed value of the test statistic.
Tobs = (mean(mcdata)-454)/sigxbar;
% This command generates the normal probability plot.
normplot(mcdata)
M = 1000; % Number of Monte Carlo trials
% Storage for test statistics from the MC trials.
Tm = zeros(1,M);
% Start the simulation.
for i = 1:M
% Generate a random sample under H_0
% where n is the sample size.
xs = sigma*randn(1,n) + 454;
Tm(i) = (mean(xs) - 454)/sigxbar;
end
% Get the critical value for alpha.
% This is a two-tail test, so we need the
% alpha and 1-alpha quantile.
alpha = 0.05/2;
cv1 = csquantiles(Tm,alpha);
cv2 = csquantiles(Tm,1-alpha);
fprintf('Critical Values\t\t%9.3f\t%9.3f\n',cv1,cv2)
```

Implementing the above code allows us two get two critical values at the points where $P(X < x_l) = \alpha/2$ and $P(X > x_u) = \alpha/2$. The calculated values are $x_l = -1.891$ (lower) and $x_u = 1.867$. The value $\alpha = 0.05$. The resulting test statistic Tobs= -2.5641 thus we reject the null hypothesis in favor of the alternative that the mean is not equal to 454. See figure 7 below for MATLAB output.

```
Command Window

Critical Values -1.891 1.867

>> Tobs

Tobs = -2.5641

fx >>
```

Figure 7: Empirical Critical Values

MATLAB Code

Exactly the same as Example 7.8 from the book. I only changed the variable M. I show just the initial part of the code:

```
% Example 7.8
clc
load mcdata; n = length(mcdata);
% Population sigma is known.
sigma = 7.8; sigxbar = sigma/sqrt(n);

M = 100000; alpha = 0.05;
% Get the critical value, using z as test statistic.
cv = norminv(alpha,0,1);
% Start the simulation.
.
```

Changing M increases the size of the sample generated by the simulation. This caused the estimation of the type I error, $\hat{\alpha}$ to approach its true value. Figure 8 shows the results for M=1000 and M=100000. As you can see the estimate when M=100000 is much closer to its true value, $\alpha=0.05$. The respective absolute differences from $\alpha=0.05$ are 0.4×10^{-2} , and -0.3×10^{-3} .

```
Command Window

>> M, alphahat

M =

1000

alphahat =

0.0460

fe >> |

>> M, alphahat

M =

100000

alphahat =

0.0503

fe >>
```

Figure 8:

```
% Code for exercise 7.7
% get the dataset
load forearm;
% PART A
n = length(forearm); % sample size
B = 1000; % number of bootstrap replicates
% Get the value of the statistic of interest.
theta1 = skewness(forearm);
theta2 = kurtosis(forearm);
theta3 = var(forearm,1);
\% Use unidrnd to get the indices to the resamples.
inds = unidrnd(n,n,B);
% Extract these from the data.
foreboot = forearm(inds);
theta1b = skewness(foreboot); % get the skewnwness for each column using
theta2b = kurtosis(foreboot); % get the kurtosis for each column using
theta3b = var(foreboot,1); % get the 2nd moment for each column using
seb1 = std(theta1b);
seb2 = std(theta2b);
biasEst1 = mean(theta1b)-theta1;
biasEst2 = mean(theta2b)-theta2;
% PART B
```

```
% find the bootstrap percentile interval for the central 2nd moment
alpha = 0.1;
k = B*alpha/2;
theta3b = sort(theta3b);
blo = theta3b(k);
bhi = theta3b(B-k);

fprintf('Skewness Standard Error and Bias %9.3f\t%4.3f\n\n',seb1,biasEst1)
fprintf('Kurtosis Standard Error and Bias %9.3f\t%4.3f\n\n',seb2,biasEst2)
fprintf('Pct Interval for 2nd Central Moment (%2.3f, %3.3f)\n',blo,bhi)
```

The resulting percentile interval is 1.029, 1.460) which means, based on the bootstrap procedure, given that $\alpha = 0.1$, we should expect, on repeated sampling, that 90% of the calculated sample 2nd central moments statistics will fall within this interval.

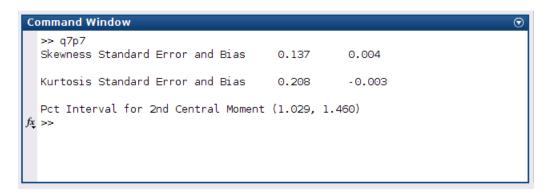


Figure 9: Results, exercise 7.7.

Exercise 7.11

```
% Exercise 7.11
clear;
% load population data
load lawpop

% find true variances for the lsat and gpa populations
popVar = var(lawpop,1);

% load sample data
load law

% BOOTSTRAP Procedure
B = 10000; % number of bootstrap replicates

% get the bootstrap values for the variance based on the sample data
bvals = bootstrp(B, @(x) var(x,1),law);
```

% get bootstrap estimates for the standar error and the bias seb = std(bvals); biasb = mean(bvals)-popVar;

The above code implements the bootstrap procedure and calculates bootstrap estimates of the variance, variance standard error, and bias. The MATLAB results are shown in figure 10 below. To get a sense of how well the bootstrap estimates perform, we look at the ratio of the standard error and the ratio of the bias as a percent of the corresponding population variance. We can see that for the LSAT measure, the standard error is about 24% of the true variance, while for the GPA measure the standard error is about 35%, hence the estimates of GPA variance are more dispersed around the true value. We note a similar result for the bias. The LSAT estimate bias is about 4% of the true variance, while the estimate bias of GPA variance is about 46% of the true parameter. We can see that the bootstrap variance estimate of LSAT is more stable that that of the GPA data.

```
>> q7pll
Population Variance: LSAT= 1463.272, GPA=0.035

Bootstrap Variance Estimate: LSAT= 1522.299, GPA=0.052

Bootstrap Variance Std Error: LSAT= 354.871, GPA=0.012

Bootstrap Variance Bias: LSAT= 59.027, GPA=0.016

Bootstrap SE as pct of Pop Var: LSAT= 0.243, GPA=0.347

Bootstrap Bias as pct of Pop Var: LSAT= 0.040, GPA=0.455
```

Figure 10: Results, exercise 7.11.