DATA 602 - Solutions to Assignment Two

```
require(ggplot2)
## Loading required package: ggplot2
require(mosaic)
## Loading required package: mosaic
## Warning: package 'mosaic' was built under R version 3.4.4
## Loading required package: dplyr
## Warning: package 'dplyr' was built under R version 3.4.4
##
## Attaching package: 'dplyr'
  The following objects are masked from 'package:stats':
##
##
##
       filter, lag
  The following objects are masked from 'package:base':
##
##
##
       intersect, setdiff, setequal, union
## Loading required package: lattice
## Loading required package: ggformula
## Warning: package 'ggformula' was built under R version 3.4.4
```

```
## Loading required package: ggstance
## Warning: package 'ggstance' was built under R version 3.4.4
##
## Attaching package: 'ggstance'
## The following objects are masked from 'package:ggplot2':
##
##
       geom_errorbarh, GeomErrorbarh
##
## New to ggformula? Try the tutorials:
    learnr::run tutorial("introduction", package = "ggformula")
##
    learnr::run_tutorial("refining", package = "ggformula")
##
## Loading required package: mosaicData
## Warning: package 'mosaicData' was built under R version 3.4.4
## Loading required package: Matrix
##
## The 'mosaic' package masks several functions from core packages in order to add
## additional features. The original behavior of these functions should not be affec
ted by this.
##
## Note: If you use the Matrix package, be sure to load it BEFORE loading mosaic.
## Attaching package: 'mosaic'
## The following object is masked from 'package:Matrix':
##
##
       mean
```

```
## The following objects are masked from 'package:dplyr':
##
## count, do, tally
```

```
## The following object is masked from 'package:ggplot2':
##
## stat
```

```
## The following objects are masked from 'package:stats':
##
## binom.test, cor, cor.test, cov, fivenum, IQR, median,
## prop.test, quantile, sd, t.test, var
```

```
## The following objects are masked from 'package:base':
##
## max, mean, min, prod, range, sample, sum
```

```
require(binom)
```

```
## Loading required package: binom
```

- **1.** (From Question 11, Assignment 1) **(2 marks)**. From Assignment 1, the delivery time is modeled by the Normal distribution with a mean of $\mu = 5.0$ hours and a standard deviation of $\sigma = 1.5$ hours. A random sample of n = 12 produced a sample mean of $\overline{X} = 5.6875$.
 - a. **Answer** Here one wishes to find $P(\overline{X} \geq 5.6875)$, where the distribution of \overline{X} will be exactly Normal with

$$\mu_{\overline{X}} = \mu_X = 5.0$$
 and $\sigma_{\overline{X}} = \frac{\sigma_X}{\sqrt{n}} = \frac{1.5}{\sqrt{12}} = 0.4330 \approx 0.433$

 $P(\overline{X} \le 5.6875)$ is then computed

```
1 - pnorm(5.6875, 5.0, (1.5/sqrt(12)))
```

```
## [1] 0.0561756
```

and

$$P(\overline{X} \ge 5.6875) = 0.0562$$

- 1 mark for the correct answer. Jiang, if the student does not divide by $\sigma_{\overline{X}}$, award zero marks.
- b. Answer:

$$\begin{split} P(0.5 \leq S \leq 1) = & P(0.5^2 \leq S^2 \leq 1^2) \\ = & P\left(\frac{(n-1)*0.5^2}{\sigma^2} \leq \frac{(n-1)S^2}{\sigma^2} \leq \frac{(n-1)*1^2}{\sigma^2}\right) \\ = & P\left(\frac{(12-1)*0.5^2}{1.5^2} \leq \chi_{df=12-1}^2 \leq \frac{(12-1)*1^2}{1.5^2}\right) \\ = & P(1.222 \leq \chi_{df=12-1}^2 \leq 4.889) \\ = & 0.0634 \end{split}$$

This probability is computed in R

[1] 0.06343895

- 1 mark for the correct answer.
- 2. (5 marks)
 - a. **Answer**. The mean and standard deviation of the distribution of \hat{p} is

$$\mu_{\hat{p}} = p = 0.80$$
 $\sigma_{\hat{p}} = \sqrt{\frac{p(1-p)}{500}} = \sqrt{\frac{0.80(1-0.80)}{500}} = 0.0179$

- (1 mark), 0.5 mark for each of the mean and standard deviation of the sample proportion.
- b. Answer Compute $P(\hat{p} \le 0.748)$ via R Studio

and
$$P(\hat{p} \le 0.748) = 0.001836 \approx 0.0018$$

• (1 mark) (Jiang, if the student computed $P(\hat{p} \le 0.744)$, then award marks, There was a typo in the original posting of this assignment.

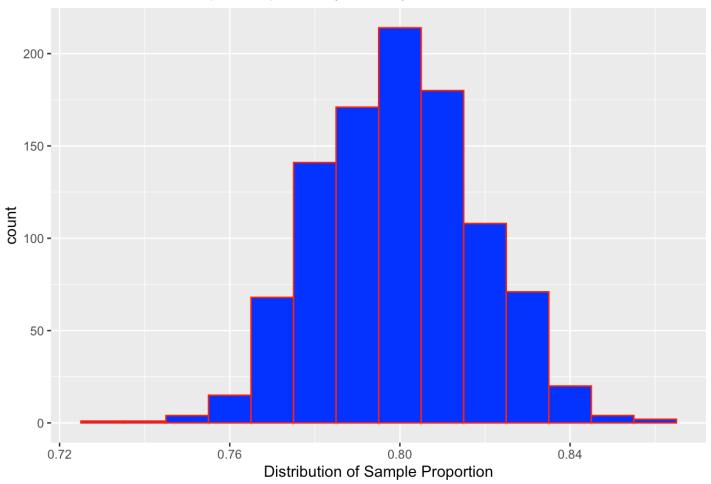
c. Answer:

```
ntimes = 1000
ntrials = 500
propsupport = numeric(ntimes)
propobserved = numeric(ntimes)
for(i in 1:ntimes)
{  propsupport[i] = (rbinom(1, ntrials, 0.80)/ntrials)
    if (propsupport[i] <= 0.744) propobserved[i] = 1 else propobserved[i] = 0
    }
ass2q2 = data.frame(propsupport, propobserved)
head(ass2q2, 4)</pre>
```

	propsupport <dbl></dbl>	propobserved <dbl></dbl>
1	0.788	0
2	0.814	0
3	0.776	0
4	0.832	0
4 rows		

ggplot(data=ass2q2, aes(x = propsupport)) + geom_histogram(fill='blue', col='red', bi nwidth=0.01) + xlab("Distribution of Sample Proportion") + ggtitle("Distribution of Sample Proportion (n = 500)")

Distribution of Sample Proportion (n = 500)



(2 marks) for the generation of the distribution of the sample proportion (which should roughly be the same as above)

proportion of sample proportions that are less than 0.748
sum(~ propobserved, data=ass2q2)/ntimes #OR

[1] 0.002

sum((propsupport <= 0.748))/ntimes

[1] 0.003

(1 mark) for computing the proportion of sample proportions that are less than the observed value of $\hat{p}=0.748$. (Jiang, results will differ from one student to the next, but they should be in neighbourhood of 0.001 - 0.003)

3. (3 marks)

Answer Solutions should be in the following structure.

One has to consider Billy's claim, that $\overline{X}>1$. To compute "how likely" Billy's claim is, we invoke the Central Limit Theorem, where the distribution of the mean number of matching numbers \overline{X} is approximately Normally distributed with a mean and standard deviation of

$$\mu_{\overline{X}} = \mu_X = 0.7347$$
 and $\sigma_{\overline{X}} = \frac{\sigma_X}{\sqrt{n}} = \frac{0.7599}{\sqrt{52}} = 0.105379 \approx 0.1054$

To compute the probability of Billy's claim:

```
options(scipen=999)
1 - pnorm(1, 0.7347, 0.1054)
```

```
## [1] 0.005916635
```

and $P(\overline{X} > 1) = 0.005917 \approx 0.0059$, which is very unlikely.

• (2 marks) for finding the probabilty of Billy's claim

Therefore, Billy's claim is not support from a probability perspective.

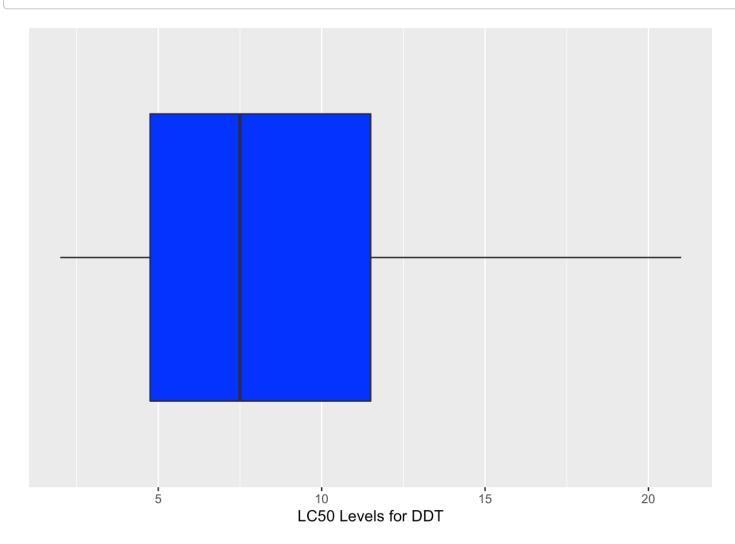
 (1 mark) for a comment on Billy's claim, the comment being supported from the probability computation

4. (10 marks)

```
lc50 = c(16, 5, 21, 19, 10, 5, 8, 2, 7, 2, 4, 9)
ass2q4df = data.frame(lc50)
head(ass2q4df, 4)
```

	lc50 <dbl></dbl>
1	16
2	5
3	21
4	19
4 rows	

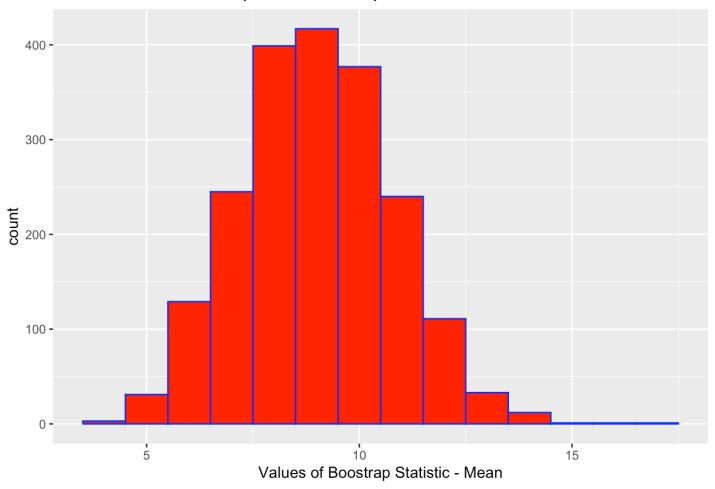
ggplot(ass2q4df) + geom_boxplot(mapping = aes(x = "var", y =lc50), fill= 'blue', na.r
m=TRUE) + xlab("") + ylab("LC50 Levels for DDT") + scale_x_discrete(breaks=NULL) + co
ord_flip()



a. Answer:

```
nsims = 2000
ntrials = 12
avelc50 = numeric(nsims)
for(i in 1:nsims)
{    avelc50[i] = mean(sample(lc50, ntrials, replace=TRUE))
}
ass2q4 = data.frame(avelc50)
ggplot(data = ass2q4, aes(x = avelc50)) + geom_histogram(fill='red', col='blue', binw idth=1) + xlab("Values of Boostrap Statistic - Mean") + ggtitle("Distribution of Boot strap Statistic: Sample Mean")
```

Distribution of Bootstrap Statistic: Sample Mean



• (2 marks) (Jiang, use your judgement here, as the results will vary from one student to the next. The distribution should be close to symmetrical with a central value around 9. If the student has roughly the same result award 2 marks; any moderate deviations penalize 1 mark)

b. Answer:

	quantile <dbl></dbl>	p <dbl></dbl>
2.5%	5.75	0.025
97.5%	12.50	0.975
2 rows		

The 95% bootstrap interval for μ , the mean amount of DDT required to kill 50% of the certain species of fish within 96 hours of exposure is somewhere beween 5.667 ppm and 12.585 ppm.

- (1 mark) for providing a 95% interval from their bootstrap distribution. As long as the student outlines/used either the **qdata()** or **quantile()** command to obtain the 2.5th and the 97.5th percentile from their bootstrap distribution in part (a), award full marks here.
- (1 mark) for the correct interpretation. Within this interpretation, ensure the student interprets their interval with the condition "from these data/based on these data (0.5 mark)" AND the student does indicate that the confidence interval is a narrowing down of the possible values of the population mean μ, that it, the mean is some value between the lower bound and the upper bound.
- c. Answer: Using the t.test() command

```
t.test(~ lc50, conf.level=0.95, data = ass2q4df)$conf
```

```
## [1] 4.91814 13.08186
## attr(,"conf.level")
## [1] 0.95
```

Using the *t*-interval, the 95% confidence interval is: $4.9181 \le \mu \le 13.082$.

• (2 marks). One mark for the correct value of the lower bound and 1 mark for the correct value of the upper bound.

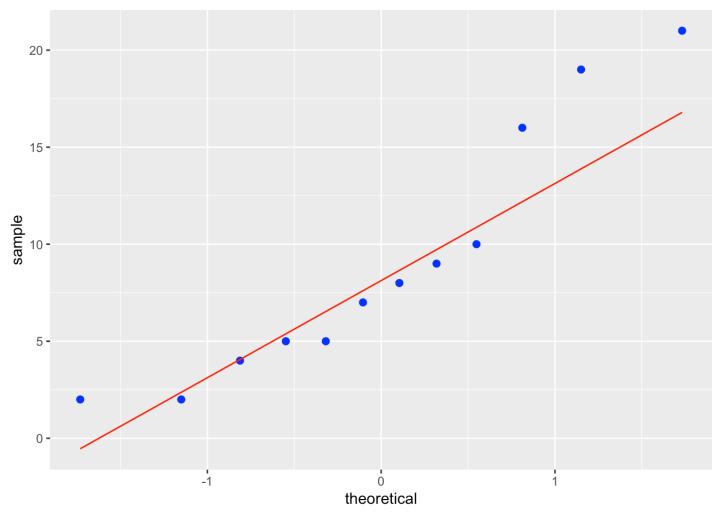
d. Answer:

• (2 marks) Jiang, you will get a variety of answers here. The bootstrap interval is "condition free", meaning it does not depend on the condition of Normality of the data and takes into account the "non-perfect" bootstrap distribuition of the sample mean which shows some skewness to the right. As long as the student has commented about this, award full marks.

e. Answer:

```
ggplot(data=ass2q4df, aes(sample = lc50)) + stat_qq(size=2, col='blue') + stat_qqline
(col='red')
```

2019-10-07, 7:38 PM



[1^]: http://angusreid.org/wp-content/uploads/2015/02/2015.02.13-Vaccinations.pdf (http://angusreid.org/wp-content/uploads/2015/02/2015.02.13-Vaccinations.pdf)

- (1 mark) for the generation of the Normal probability plot
- (1 mark) for a comment on the data following a Normal distribution (this will be subjective, it would appear that these data are not Normally distributed from the absence of linearity, but some may view the bulk-middle as being linear, and that is fine).

5. (7 marks)

a. **Answer:** A 95% confidence interval for p, using the "plus-2/plus-4" version is computed using the **binom.confint()** command:

binom.confint(571, 1866, method="agresti-coull")

method	x n	mean	lower	upper
<fctr></fctr>	<dbl> <dbl></dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>

```
1 agresti-coull 571 1866 0.3060021 0.2855056 0.3272958
1 row
```

```
#OR
binom.test(571, 1866, ci.method="plus4")
```

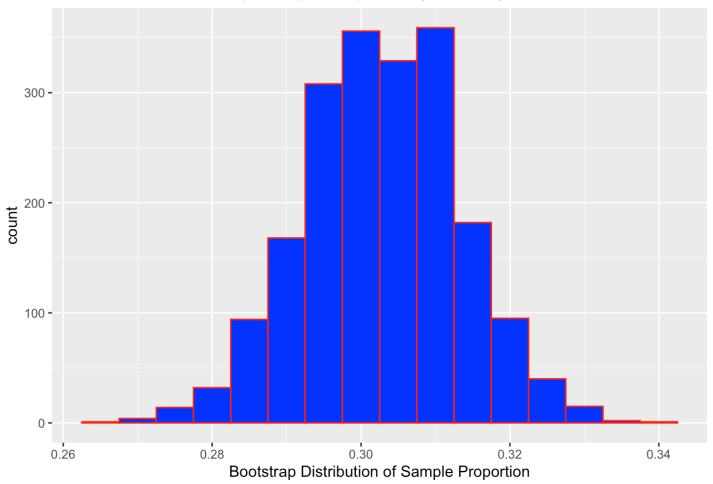
```
##
## Exact binomial test (Plus 4 CI)
##
## data: 571 out of 1866
## number of successes = 571, number of trials = 1866, p-value <
## 0.00000000000000022
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
## 0.2855226 0.3273117
## sample estimates:
## probability of success
## 0.3060021</pre>
```

The 95% confidence interval for p is \$ 0.29 p 0.33\$.

- (2 marks). 1 mark for the correct lower bound, 1 mark for the correct upper bound
- b. **Answer** Below is the code to create a bootstrap distribution of the sample proportion \hat{p} :

```
nsims = 2000
nsize=1866
sampleprop = numeric(nsims)
ques2data = c(rep(0, 1886 - 571), rep(1, 571)) #create a data vectore of 571 1s and (
1866 - 571) 0s
for(i in 1:nsims)
{    sampleprop[i] = sum(sample(ques2data, nsize, replace=TRUE))/(nsize)
}
ques5df = data.frame(sampleprop)
# head(ques2df, 3)
ggplot(ques5df, aes(x = sampleprop)) + geom_histogram(col='red', fill='blue', binwidt
h=0.005) + xlab("Bootstrap Distribution of Sample Proportion") + ggtitle("Distributio
n of Bootstrap Sample Proportion (n = 1866)")
```

Distribution of Bootstrap Sample Proportion (n = 1866)



- (2 marks) Mark similar to Question 1(a). If the student provides a bootstrap distribution of the sample proportion which should appear to be similar to provided, award full (2) marks here Ajmery.
- c. **Answer:** To obtain a 95% bootstrap CI for p, obtain the 2.5 and 97.5 percentiles:

	quantile <dbl></dbl>	p <dbl></dbl>
2.5%	0.2824223	0.025
97.5%	0.3231511	0.975
2 rows		

The 95% confidence interval is $0.2819 \le p \le 0.3231$.

• **(1 mark)** Jiang, mark similar to Question 4(b). As long as the student obtains their interval from the bootstrap distribuion of \widheatp, award 1 mark.

- d. **Answer:** The bootstrap interval of [0.28819, 0.3231] compared to the plus2/plus4 of [0.28, 0.33]? Both are similar width, the bootstrap distribution showing some skewness to the left would be the preferred interval here.
- (2 marks) Jiang, please mark Question 5(d) similar to Question 4(e). As long as the student provides a "statistical justification" for their answer, award 2 marks.

6. (9 marks)

a. **Answer:** Below is the bootstrap distribution for the \widehat{p}_{HS} .

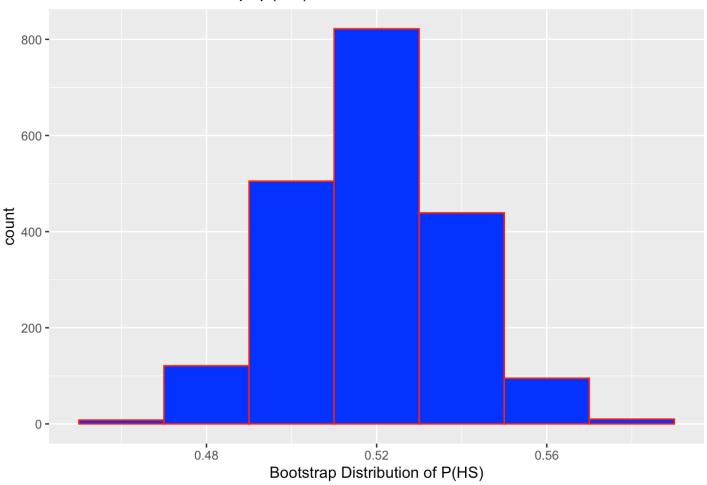
```
nsims = 2000
hsdata = c(rep(0,670-348), rep(1, 348)) #hs data with 348 1s and (670-348) 0s
hssampleprop = numeric(nsims)
for(i in 1:nsims)
{ hssampleprop[i] = sum(sample(hsdata, 670, replace=TRUE))/670}
gues6adf = data.frame( hssampleprop)
head(ques6adf, 3)
```

	hssampleprop <dbl></dbl>
1	0.5701493
2	0.5194030
3	0.5552239
3 rows	

Below is a histogram of the bootstrap statistic \widehat{p}_{HS}

```
ggplot(data=ques6adf, aes(x = hssampleprop)) + geom_histogram(col='red', fill='blue',
binwidth=0.02) + xlab("Bootstrap Distribution of P(HS)") + ggtitle("Distribution of B
ootstrap: p(HS)")
```

Distribution of Bootstrap: p(HS)



- (1 mark) Mark similar to how students were marked in generating the bootstrap distributions in both Question 1 and Question 2.
- b. **Answer:** Below is the bootstrap distribution for the \widehat{p}_{Uni} .

```
nsims = 2000
unidata = c(rep(0,376-274), rep(1, 274)) #university data with 274 1s and (376-274) 0
s
unisampleprop = numeric(nsims)
for(i in 1:nsims)
{
    unisampleprop[i] = sum(sample(unidata, 376, replace=TRUE))/376
}
ques6bdf = data.frame(unisampleprop)
head(ques6bdf, 3)
```

unisampleprop

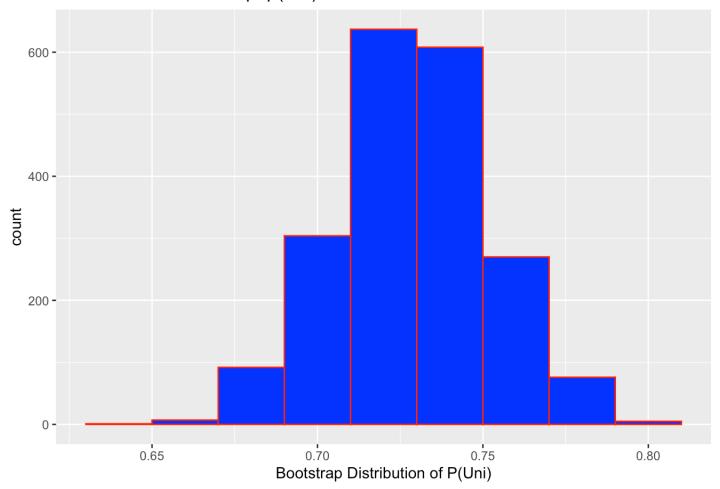
<dbl>

1	0.7287234
2	0.7393617
3	0.7367021
3 rows	

Below is a histogram of the bootstrap statistic $\widehat{p}_{\mathit{Uni}}$

ggplot(data=ques6bdf, aes(x = unisampleprop)) + geom_histogram(col='red', fill='blue'
, binwidth=0.02) + xlab("Bootstrap Distribution of P(Uni)") + ggtitle("Distribution of Bootstrap: p(Uni)")

Distribution of Bootstrap: p(Uni)



- (1 mark) Mark similar to how students were marked in generating the bootstrap distributions in part (a)
- c. Answer: Below is the bootstrap distribution for the $\hat{p}_{Uni}-\hat{p}_{HS}$.

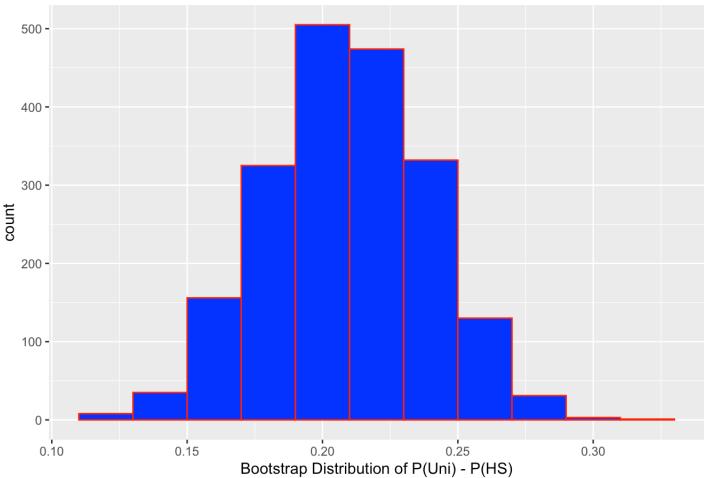
```
nsims = 2000
unidata = c(rep(0,376-274), rep(1, 274)) #university data with 274 1s and (376-274) 0
s
hsdata = c(rep(0,670-348), rep(1, 348)) #hs data with 348 1s and (670-348) 0s
unisampleprop = numeric(nsims)
hssampleprop = numeric(nsims)
diffsampleprop = numeric(nsims)
for(i in 1:nsims)
{
    unisampleprop[i] = sum(sample(unidata, 376, replace=TRUE))/376
    hssampleprop[i] = sum(sample(hsdata, 670, replace=TRUE))/670
    diffsampleprop[i] = unisampleprop[i] - hssampleprop[i]
}
ques6df = data.frame(unisampleprop, hssampleprop, diffsampleprop)
head(ques6df, 3)
```

	unisampleprop <dbl></dbl>	hssampleprop <dbl></dbl>	diffsampleprop <dbl></dbl>
1	0.7393617	0.5358209	0.2035408
2	0.7606383	0.5029851	0.2576532
3	0.7420213	0.4955224	0.2464989
3 rows			

Below is a histogram of the bootstrap statistic $\hat{p}_{\mathit{Uni}} - \hat{p}_{\mathit{HS}}$

```
ggplot(data=ques6df, aes(x = diffsampleprop)) + geom_histogram(col='red', fill='blue'
, binwidth=0.02) + xlab("Bootstrap Distribution of P(Uni) - P(HS)") + ggtitle("Distribution of Bootstrap: p(Uni) - p(HS)")
```

Distribution of Bootstrap: p(Uni) - p(HS)



• (3 marks) Mark similar to how students were marked in generating the bootstrap distributions in both Question 1 and Question 2.

d. Answer:

From this, the 95% bootstrap interval is

	quantile <dbl></dbl>	p <dbl></dbl>
2.5%	0.1513814	0.025
97.5%	0.2653872	0.975
2 rows		

$$0.1508 \le p_{Uni} - p_{Hs} \le 0.2683$$

• (2 mark) for the provision of the bootstrap interval, 1 mark for the lower bound, and 1 mark for the upper bound.

No, one cannot infer that $p_{Uni} = p_{HS}$, as the confidence interval has a lower bound of approximately 0.149 and an upper bound of about 0.27. Because this confidence inteval has a lower bound that exceeds 0, one can infer that p_{Uni} **EXCEEDS** p_{HS} by anywhere from 14.9% to 27%.

• (2 marks) Jiang, as long as the student makes a statement that the CI does not capure zero, hence one cannot conclude that $p_{Uni} = p_{HS}$, award 2 marks.

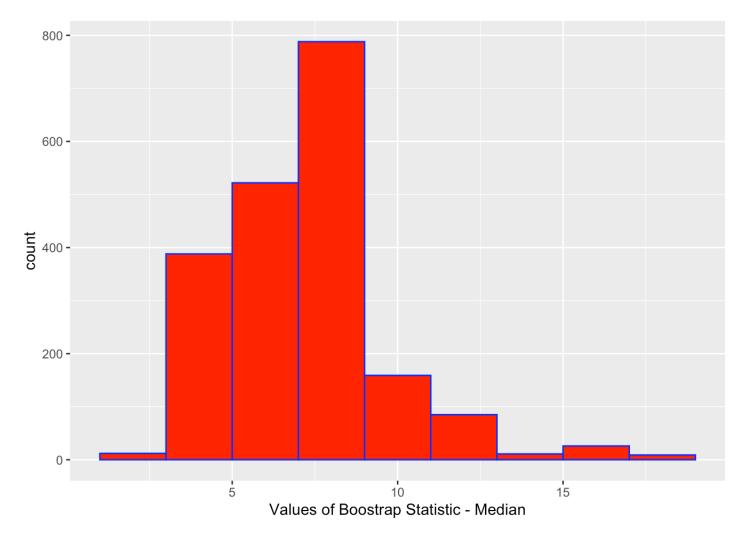
7. (3 marks)

Re-using the code from Question 4, changing the mean to median

```
lc50 = c(16, 5, 21, 19, 10, 5, 8, 2, 7, 2, 4, 9)
nsims = 2000
ntrials = 12
medianlc50 = numeric(nsims)
for(i in 1:nsims)
{    medianlc50[i] = median(sample(lc50, ntrials, replace=TRUE))
}
ass3q4 = data.frame(medianlc50)
```

The bootstrap distribution of the sample median \widetilde{X} appears below.

```
ggplot(data = ass3q4, aes(x = medianlc50)) + geom_histogram(fill='red', col='blue', b
inwidth=2) + xlab("Values of Boostrap Statistic - Median")
```



• (2 marks) for creating the boostrap distribution of the sample median.

From this, a 99% confidence interval for $\widehat{\mu}$ is $3.5 \le \widetilde{\mu} \le 16$.

	quantile <dbl></dbl>	p <dbl></dbl>
0.5%	3	0.005
99.5%	16	0.995
2 rows		

quantile(ass3q4\$median1c50, c(0.005, 0.995))

```
## 0.5% 99.5%
## 3 16
```

• (1 marks) 0.5 mark for the correct lower bound and 0.5 mark for the correct upper bound based on the student's usage of the qdata() or the quantile() command

8. (7 marks)

a. **Answer:** 95% confidence interval for p_{NDP} is

```
binom.test(126, 1003, ci.method="plus4")$conf
```

```
## [1] 0.1065370 0.1476835
## attr(,"conf.level")
## [1] 0.95
## attr(,"method")
## [1] "plus4"
```

and

$$0.1065 \le p_{NDP} \le 0.1476$$

- 2 marks, 1 for the correct lower bound and 1 for the correct upper bound
- b. **Answer:** The bootstrap distribution of $\frac{X_{NDP}+2}{n+4}$ is provided below

```
Nsims = 2000

nsize = 1003

ndpdata= c(rep(0, 1003 - 126), rep(1, 126)) #puts the data in the form of (1003 - 126)

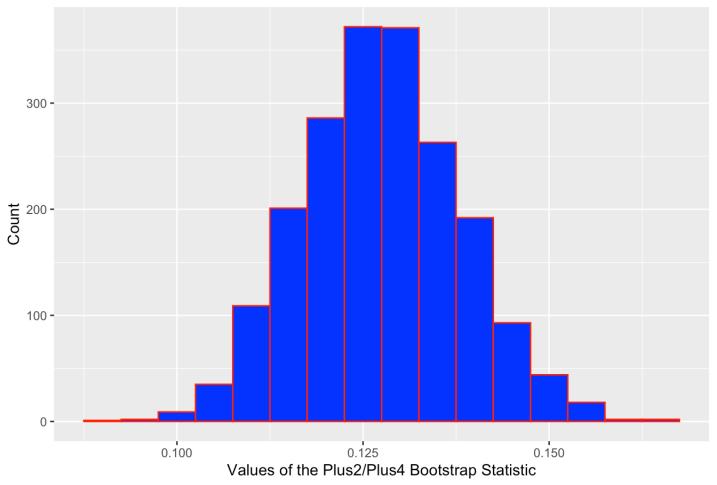
) 0s and 126 1s
```

Here are the contents of the bootstrap:

```
bootptilde = numeric(Nsims)
for(i in 1:Nsims)
{
   bootptilde[i] = (sum(sample(ndpdata, nsize, replace=TRUE)) + 2)/(nsize + 4)
}
bootq8 = data.frame(bootptilde)
```

ggplot(data=bootq8, aes(x = bootptilde)) + geom_histogram(col='red', fill='blue', bin
width = 0.005) + xlab("Values of the Plus2/Plus4 Bootstrap Statistic") + ylab("Count"
) + ggtitle("Distribution of Bootstrap Plus2/Plus4")

Distribution of Bootstrap Plus2/Plus4



- 2 marks for generating the bootstrap distribution of $\frac{X_{NDP}+2}{n+4}$
- c. **Answer:** The 95% bootstrap interval for p from the result in (b) is

qdata(~bootptilde, c(0.025, 0.975), data=bootq8)

	quantile <dbl></dbl>	p <dbl></dbl>
2.5%	0.1082423	0.025
97.5%	0.1489573	0.975
2 rows		

the 95% bootstrap interval for p is then

$$0.1072 \le p \le 0.1470$$

- 1 mark for the bootstrap inteval. Again, results will vary from one student to the next.
- d. Answers here will vary Jiang. As long as the student provides a sound, statistical commentary based on the result they obtained from (a) and (c), awared **2 marks**.