Lab 11

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```
library(UsingR)
## Loading required package: MASS
## Loading required package: HistData
## Loading required package: Hmisc
## Loading required package: lattice
## Loading required package: survival
## Loading required package: Formula
## Loading required package: ggplot2
##
## Attaching package: 'Hmisc'
## The following objects are masked from 'package:base':
##
##
       format.pval, units
##
## Attaching package: 'UsingR'
## The following object is masked from 'package:survival':
##
##
       cancer
```

1) Does the data in normtemp support that average body temp is 98.6? Hypothesis test

```
## 98.14269 98.35577
## sample estimates:
## mean of x
## 98.24923
# The p-value is 2.411e^-07,
# we have enough evidence to reject the null hypothesis.
```

2) Waiting time of 100 bank customers

a) construct a 95% confidence interval

```
t.test(waiting_data, conf.level=0.95)$conf.int

## [1] 8.441023 11.312977

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

b) construct a 99% confidence interval

```
t.test(waiting_data, conf.level=0.99)$conf.int
## [1] 7.976271 11.777729
## attr(,"conf.level")
## [1] 0.99
```

c) Is there enough evidence to conclude that it takes on average more than 8 minutes to be served

3) Smokers and Non-Smokers. Is there evidence that the nonsmokers has a higher score than smokers?

```
u(ns) - u(s) = 0
# Null:
# Alternative: u(ns) - u(s) > 0
smokers = c(16,20,14,21,20,18,13,15,17,21)
nonsmokers = c(18,22,21,17,20,17,23,20,22,21)
t.test(nonsmokers, smokers, alt="greater")
##
## Welch Two Sample t-test
##
## data: nonsmokers and smokers
## t = 2.2573, df = 16.376, p-value = 0.01899
## alternative hypothesis: true difference in means is greater than 0
## 95 percent confidence interval:
## 0.5919017
                    Inf
## sample estimates:
## mean of x mean of y
       20.1
                  17.5
# With a p-value of 0.019,
# we have enough evidence to reject the null hypothesis.
```

4) Pulse rate Experiment

```
pulse_data = read.table("http://www.statsci.org/data/oz/ms212.txt", header=TRUE)
```

a) Test the hypothesis of whether there is a difference in pulse rate if the students were sitting

```
Null: u(norm) - u(sitting) = 0
```

```
# Alternative: u(norm) - u(sitting) != 0

# First version:
t.test(pulse_data$Pulse2~pulse_data$Ran, alt="greater")
```

##

```
## Welch Two Sample t-test
##
## data: pulse data$Pulse2 by pulse data$Ran
## t = 13.156, df = 57.309, p-value < 2.2e-16
## alternative hypothesis: true difference in means between group 1 and group 2 is greater than 0
## 95 percent confidence interval:
## 45.38385
## sample estimates:
## mean in group 1 mean in group 2
         126.84783
                          74.85714
# Second version:
R = subset(pulse_data, pulse_data$Ran=="1")
S = subset(pulse_data, pulse_data$Ran=="2")
t.test(R$Pulse2, S$Pulse2, alt="greater")
##
##
   Welch Two Sample t-test
##
## data: R$Pulse2 and S$Pulse2
## t = 13.156, df = 57.309, p-value < 2.2e-16
## alternative hypothesis: true difference in means is greater than 0
## 95 percent confidence interval:
## 45.38385
                  Tnf
## sample estimates:
## mean of x mean of y
## 126.84783 74.85714
# With a p-value of 2.2e^-16, we have enough evidence to reject the null hypothesis.
```

b) Test the hypothesis whether the average pulse rate for running students increased by 10 after they ran

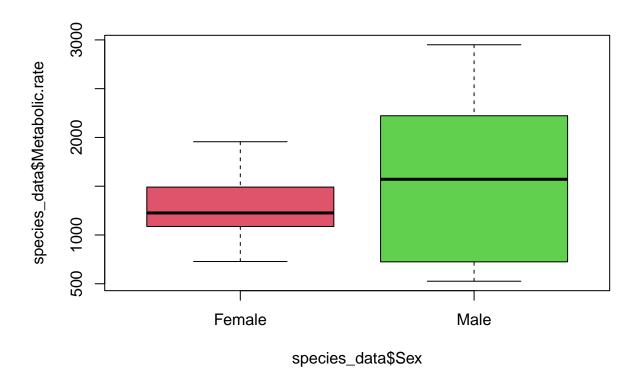
```
# Null:
                u(norm) - u(ran) = 10
# Alternative: u(norm) - u(ran) > 10
length(R$Age)
## [1] 46
t.test(R$Pulse2, R$Pulse1,
       alt="greater",
       mu=10,
       paired=T)
##
## Paired t-test
## data: R$Pulse2 and R$Pulse1
## t = 13.311, df = 45, p-value < 2.2e-16
## alternative hypothesis: true mean difference is greater than 10
## 95 percent confidence interval:
## 46.16911
                  Inf
## sample estimates:
## mean difference
##
           51.3913
```

```
# With a p-value of 2.2e^-16,
# we have evidence to reject the null hypothesis.
```

5) Furness and Bryant Metabolic rates of male and female breeding fulmars

a) Display the metabolic rate of female and male group using side-by-side boxplot

```
species_data = read.csv('C:\\repos\\STAT 50001\\Lab 11\\species.csv', sep='\t')
boxplot(species_data$Metabolic.rate ~ species_data$Sex, col=c(2,3))
```



b) Test the hypothesis whether there is a difference in metabolic rate based on gender

```
# Null: u(f) - u(m) = 0
# Alternative: u(f) - u(m) != 0
t.test(species_data$Metabolic.rate ~ species_data$Sex)

##
## Welch Two Sample t-test
##
## data: species_data$Metabolic.rate by species_data$Sex
## t = -0.77341, df = 10.466, p-value = 0.4564
```