# **Final Lab**

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# Part 1

I will now load my packages.

```
library(ggplot2)
wcgs <- read.csv("wcgs.csv")</pre>
```

We will now find a few summary statistics.

```
mean(wcgs$sbp)
```

[1] 128.6328

```
mean(wcgs$weight)
```

[1] 169.9537

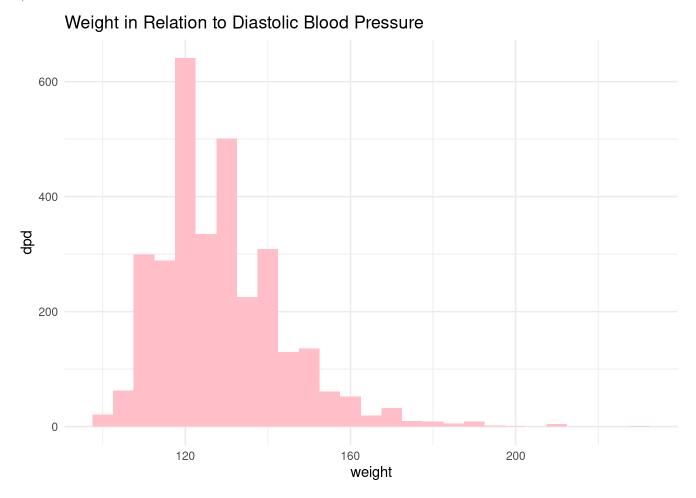
```
mean(wcgs$dbp)
```

[1] 82.01554

I found the mean of systolic blood pressure, weight, and diastolic blood pressure.

We will now create a visual graph showing the relation between variables weight and diastolic blood pressure.

```
ggplot(wcgs, aes(x=sbp)) +
  geom_histogram(binwidth = 5, fill="pink")+
  xlab("weight")+
  ylab("dpd")+
  ggtitle("Weight in Relation to Diastolic Blood Pressure") + theme_minimal()
```



This histogram shows the relation between weight and diastolic blood pressure.

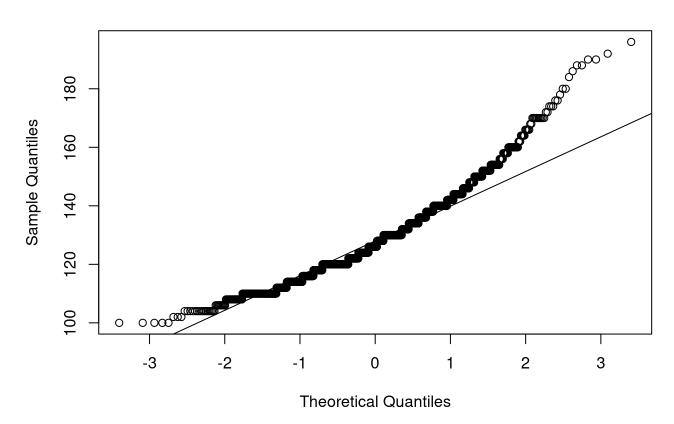
# Part 2

We will now conduct a hypothesis test for independent samples that considers the effects of smoking. Group 1 are the smokers while group 2 are the non-smokers.

```
group1<-with(wcgs,sbp[smoking == "Smoker"])
group2<-with(wcgs,sbp[smoking == "Non-smoker"])

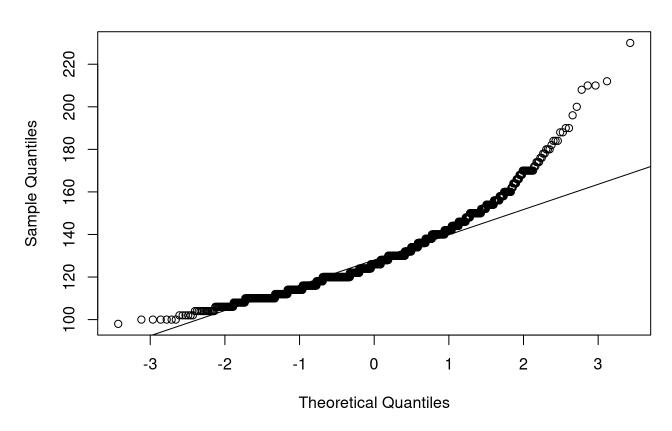
qqnorm(group1)
qqline(group1)</pre>
```

# **Normal Q-Q Plot**



qqnorm(group2)
qqline(group2)

### **Normal Q-Q Plot**



```
var.test(group1,group2)
```

F test to compare two variances

We will now conduct a t.test.

Two Sample t-test

```
t.test(sbp ~ smoking, data = wcgs, alternative = "two.sided", conf.level = 0.95,
```

```
data: sbp by smoking
t = -0.14729, df = 3152, p-value = 0.8829
```

```
alternative hypothesis: true difference in means between group Non-smoker and group Smoker is not equal to 0
95 percent confidence interval:
-1.1363606 0.9775651
sample estimates:
mean in group Non-smoker mean in group Smoker
128.5950 128.6744
```

Since the p-value is not less than 0.05, we fail to reject the null hypothesis. There is not enough evidence to suggest group 1 and group 2 had different systolic blood pressures.

## Part 3

We can check the assumptions by running <code>chisq.test()</code> . We can check the expected counts with the following code:

```
output <- chisq.test(table(wcgs$personality))
output$expected</pre>
```

```
A1 A2 B3 B4 788.5 788.5 788.5
```

We can find the p-value with the following code:

```
output$p.value
```

[1] 7.725469e-258

There were different proportions of personality types discovered.

## Part 4

We are determining the most correlated variable by:

1. First, we will filter the data to only select numerical values.

```
filtered_data <- Filter(is.numeric, wcgs)
head(filtered_data)</pre>
```

```
Χ
      id age height weight sbp dbp chol ncigs timechd
1 1 2001 49
                73
                                          25
                      150 110 76
                                   225
                                                1664
2 2 2002 42
                70
                      160 154 84
                                   177
                                          20
                                                3071
3 3 2003 42
                69
                      160 110 78
                                   181
                                          0
                                                3071
4 4 2004 41
                68
                      152 124 78
                                   132
                                          20
                                                3064
5 5 2005 59
                                   255
                70
                      150 144
                               86
                                          20
                                                1885
6 6 2006 44
                72
                      204 150 90
                                   182
                                           0
                                                3102
```

2. Second, we will calculate the correlation matrix.

```
cor(filtered_data, use = "pairwise.complete.obs")
```

```
Χ
                           id
                                      age
                                               height
                                                           weight
Χ
        1.000000000
                   0.956757817 -0.039143689 -0.056722904 -0.004811028
id
        0.956757817
                   1.000000000 -0.048160214 -0.052294226 -0.003780646
age
       -0.039143689 -0.048160214 1.000000000 -0.095375682 -0.034404537
       -0.056722904 -0.052294226 -0.095375682 1.000000000 0.532935466
weight
      -0.004811028 -0.003780646 -0.034404537 0.532935466 1.000000000
       -0.034136607 -0.044388131 0.165746397 0.018373573 0.253249623
sbp
dbp
       -0.051566439 -0.052577344 0.139197757 0.010275549 0.295920186
chol
        0.050372896 0.057845065 0.089188510 -0.088937779 0.008537442
ncigs
        timechd 0.048278246 0.041181424 -0.070919630 -0.009895169 -0.065350046
                         adb
                                    chol
                                              ncigs
                                                        timechd
              sbp
Χ
       -0.03413661 -0.05156644 0.050372896 0.013432770 0.048278246
id
       -0.04438813 -0.05257734 0.057845065 0.011667283 0.041181424
        0.16574640 0.13919776 0.089188510 -0.005033852 -0.070919630
age
height
        weight
        0.25324962 0.29592019 0.008537442 -0.081747507 -0.065350046
sbp
        1.00000000 0.77290641 0.123061297 0.029977529 -0.107884203
dbp
        0.77290641 1.00000000 0.129597108 -0.059342317 -0.110693969
        0.12306130 0.12959711 1.000000000 0.096031834 -0.095390054
chol
        0.02997753 -0.05934232 0.096031834 1.000000000 -0.093933141
ncigs
timechd -0.10788420 -0.11069397 -0.095390054 -0.093933141 1.000000000
```

The variable that is most correlated is dbp, and the secondmost correlated variable is weight. We will use these variables to build this model.

Now, we will build the models.

```
model <- lm(sbp ~ weight, data = wcgs)
summary(model)</pre>
```

```
Call:
```

```
lm(formula = sbp ~ weight, data = wcgs)
```

### Residuals:

```
Min 10 Median 30 Max -29.549 -10.097 -2.456 7.724 99.544
```

### Coefficients:

```
Estimate Std. Error t value Pr(>|t|)
(Intercept) 97.78884 2.11473 46.24 <2e-16 ***
weight 0.18148 0.01235 14.70 <2e-16 ***
---
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

Residual standard error: 14.63 on 3152 degrees of freedom

Multiple R-squared: 0.06414, Adjusted R-squared: 0.06384 F-statistic: 216 on 1 and 3152 DF, p-value: < 2.2e-16 Our model is sbp = 0.1814\*weight + 97.7888

```
model <- lm(sbp ~ dbp, data = wcgs)
summary(model)</pre>
```

### Call:

 $lm(formula = sbp \sim dbp, data = wcgs)$ 

### Residuals:

```
Min 10 Median 30 Max -30.237 -6.212 -1.394 5.386 62.581
```

#### Coefficients:

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 9.594 on 3152 degrees of freedom Multiple R-squared: 0.5974, Adjusted R-squared: 0.5973 F-statistic: 4677 on 1 and 3152 DF, p-value: < 2.2e-16

Our model is sbp = 1.2013\*dbp + 30.1102.

The diastolic blood pressure has a stronger correlation to systolic blood pressure than weight.

# Part 5

We found that there were not enough evidence to conlude that smokers and non-smokers had different systolic blood pressure. There were different proportions of personalities discovered so we could not complete a Chi-Squared test. The highest correlated variables with systolic blood pressure were weight followed by diastolic blood pressure. From this, we discovered that diastolic blood pressure has a stronger correlation to systolic blood pressure than weight.