

# Week 5 Lab

AUTHOR  
Jessica Tran

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## Data

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For this lab, we will be using the data set `C02`. Take some time to get familiar with the data set using the help function.

```
?C02
```

## Problem 1

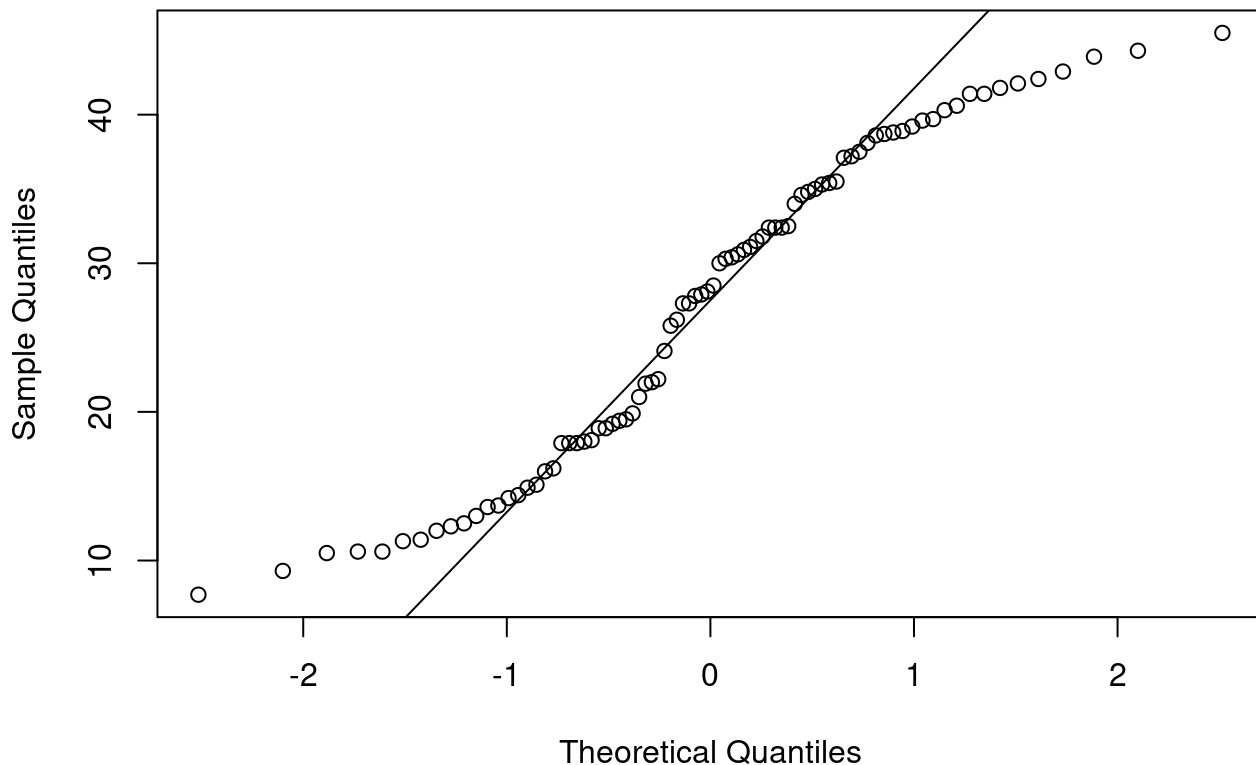
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Conduct a hypothesis test for a mean of one sample. Using the data set `C02` and the variable `uptake`, test the null hypothesis that the population mean `uptake` is mean 28 vs. the alternate hypothesis that the population mean `uptake` is less than 28 with a significance level of 0.05.

Check any assumptions:

```
qqnorm(C02$uptake)  
qqline(C02$uptake)
```

## Normal Q-Q Plot



If the assumptions are met, calculate the p-value:

```
t.test(C02$uptake, mu = 28, alternative = "less", conf.level = 0.95)
```

### One Sample t-test

```
data: C02$uptake
t = -0.6669, df = 83, p-value = 0.2533
alternative hypothesis: true mean is less than 28
95 percent confidence interval:
 -Inf 29.17585
sample estimates:
mean of x
 27.2131
```

## Problem 2

Conduct a hypothesis test for independent samples. Use the data set **C02** and the variables **Type** and **uptake**. Let group 1 be **Quebec** and group 2 be **Mississippi**. Test the null hypothesis that the population mean **uptake** of the **Quebec** samples is the **same** as that of the **Mississippi** samples vs. the alternate hypothesis that the population mean **uptake** of the **Quebec** samples is **different**

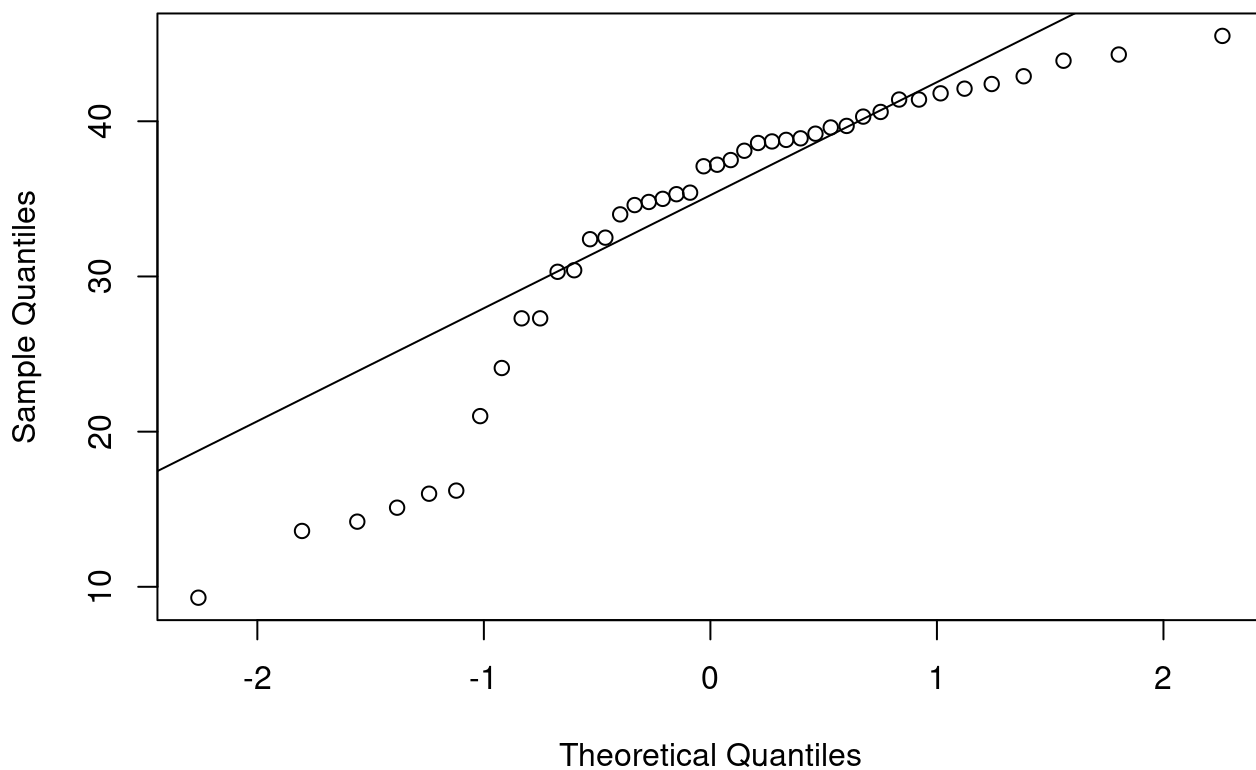
than that of the **Mississippi** samples with a significance level of 0.01. Comment out each line of code to explain what you are doing.

Check any assumptions:

```
group1 <- with(CO2, uptake[Type == "Quebec"]) # select just group 1
group2 <- with(CO2, uptake[Type == "Mississippi"]) # select just group 2

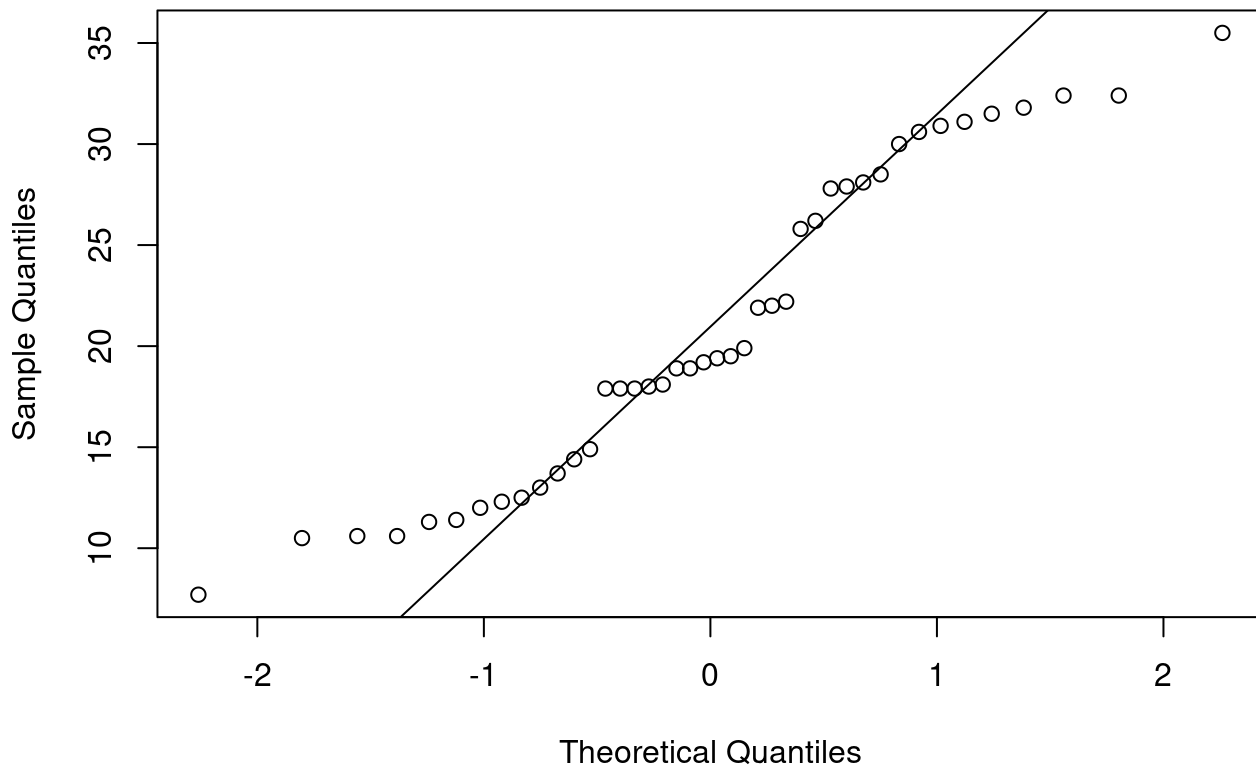
qqnorm(group1)                                # QQ plot for group 1
qqline(group1)
```

### Normal Q-Q Plot



```
qqnorm(group2)                                # QQ plot for group 2
qqline(group2)
```

## Normal Q-Q Plot



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

F test to compare two variances

data: group1 and group2

F = 1.532, num df = 41, denom df = 41, p-value = 0.1763

alternative hypothesis: true ratio of variances is not equal to 1

95 percent confidence interval:

0.8234755 2.8500701

sample estimates:

ratio of variances

1.53198

If the assumptions are met, calculate the p-value:

```
t.test(uptake ~ Type, data = C02, alternative = "two.sided", conf.level = 0.99, p
```

Two Sample t-test

data: uptake by Type

t = 6.5969, df = 82, p-value = 3.835e-09

alternative hypothesis: true difference in means between group Quebec and group Mississippi is not equal to 0  
99 percent confidence interval:  
7.598856 17.720192  
sample estimates:  
mean in group Quebec mean in group Mississippi  
33.54286 20.88333

## Problem 3

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Conduct a hypothesis test for paired samples. Use the data set `C02` and the variable `uptake`. Let group 1 be `chilled` and group 2 be `nonchilled`. Test the null hypothesis that the population mean `uptake` of the `chilled` samples is the same as that of the `nonchilled` samples vs. the alternate hypothesis that the population mean `uptake` of the `chilled` samples is different than that of the `nonchilled` samples with a significance level of 0.05. Comment out each line of code to explain what you are doing.

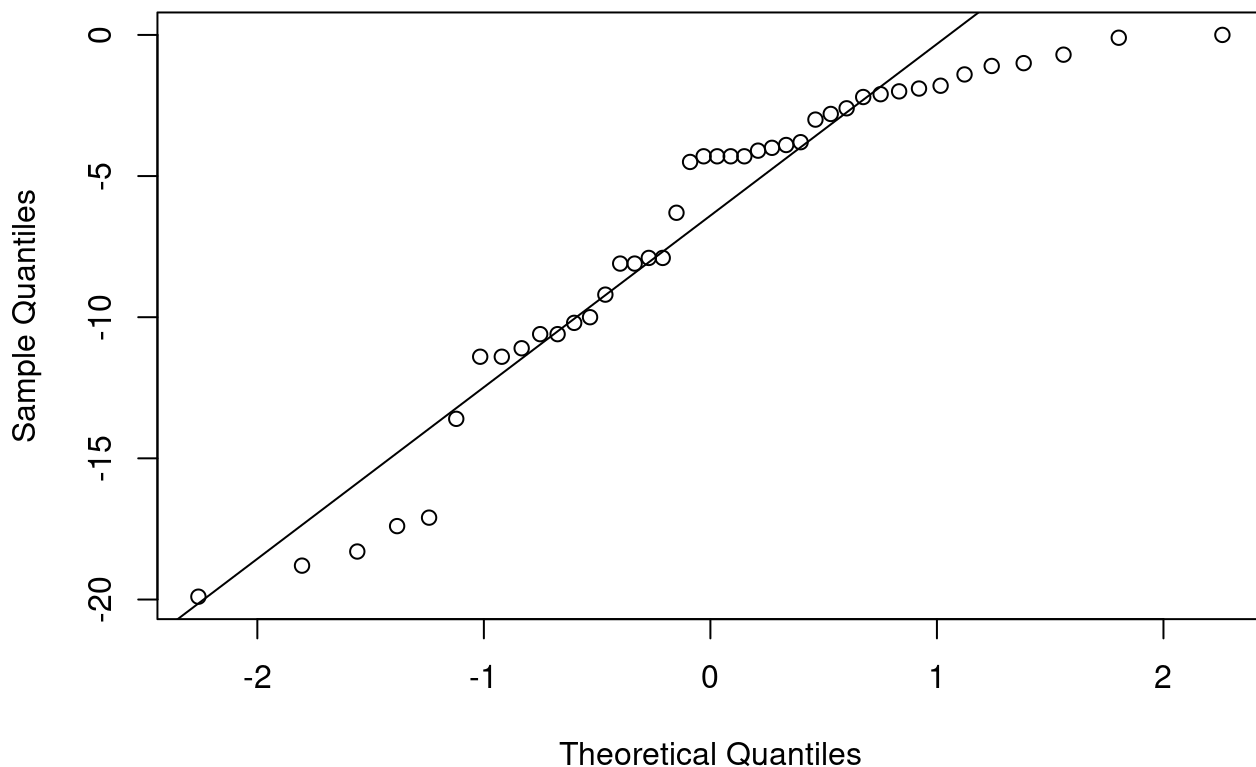
Check any assumptions:

```
group1 <- with(C02, uptake[Treatment == "chilled"]) # select just group 1
group2 <- with(C02, uptake[Treatment == "nonchilled"]) # select just group 2

differenced_data <- group1 - group2

qqnorm(differenced_data)
qqline(differenced_data)
```

## Normal Q-Q Plot



If the assumptions are met, calculate the p-value:

```
t.test(uptake ~ Treatment, data = C02, alternative = "two.sided", conf.level = 0.
```

Paired t-test

data: uptake by Treatment

t = 7.939, df = 41, p-value = 8.051e-10

alternative hypothesis: true mean difference is not equal to 0

95 percent confidence interval:

5.114589 8.604458

sample estimates:

mean difference

6.859524

## Problem 4

Conduct a hypothesis test for one proportion. We are interested in knowing what proportion of plants have an uptake greater than  $25 \mu\text{mol}/m^2 \text{ sec}$ . Out of the 84 plants, 49 of them had an uptake greater than  $25 \mu\text{mol}/m^2 \text{ sec}$ . At a significance level of 1%, can you conclude that a majority of plants have an update greater than  $25 \mu\text{mol}/m^2 \text{ sec}$ . Comment out each line of code to explain what you are doing.

Check any assumptions:

```
84*(0.5)      #np
```

```
[1] 42
```

```
84*(1-0.5)     #n(1-p)
```

```
[1] 42
```

If the assumptions are met, calculate the p-value:

```
prop.test(49, 84, p = 0.5, alternative = "greater", correct = FALSE)
```

1-sample proportions test without continuity correction

```
data: 49 out of 84, null probability 0.5
X-squared = 2.3333, df = 1, p-value = 0.06332
alternative hypothesis: true p is greater than 0.5
95 percent confidence interval:
 0.4936065 1.0000000
sample estimates:
      p
0.5833333
```

## Submitting

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Submit the following to Canvas:

- Your rendered PDF titled Lastname\_5R. Make sure your name is at the top of the document.
- Your .qmd file