

# BuckVolExplr.R

*Audrey McCombs*

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```
library(ggplot2)
library(dplyr)
```

```
##
## 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
```

```
setwd("D:/Iowa State University/Debinski Lab/Nectar data/MAL")
```

```
buckvol15 <- read.csv("nectar analysis/data files/buckvol15.csv", header = T)
buckvol16 <- read.csv("nectar analysis/data files/buckvol16.csv", header = T)
buckvolboth <- rbind(buckvol15,buckvol16)
```

*#Data summaries*

```
summary(buckvol15)
```

##	date	plot	treatment	quad	volume
##	2015-06-22: 48	WSR10 : 50	C:229	WSR10NE: 14	Min. :0.01562
##	2015-06-23: 46	WH12 : 45	H:226	EC3NE : 13	1st Qu.:0.21818
##	2015-06-21: 39	EH4 : 44		WHSR9SW: 13	Median :0.36364
##	2015-06-24: 39	WHSR9 : 44		EC3SW : 12	Mean :0.44440
##	2015-06-20: 35	EC3 : 43		EHSR1SE: 12	3rd Qu.:0.54545
##	2015-06-30: 34	CH5 : 40		WH12NW : 12	Max. :3.03125
##	(Other) :214	(Other):189		(Other):379	

```
summary(buckvol16)
```

##	date	plot	treatment	quad	volume
##	2016-06-23:65	WSR10 : 51	C:190	EHSR1SW: 14	Min. :0.01515
##	2016-06-24:70	WHSR9 : 45	H:186	WSR10NW: 14	1st Qu.:0.04545
##	2016-06-25:58	WH12 : 43		WSR10SE: 14	Median :0.07576
##	2016-06-26:52	CC6 : 40		WSR10NE: 13	Mean :0.08464
##	2016-06-27:56	CH5 : 30		CC6NW : 12	3rd Qu.:0.10985
##	2016-06-29:53	CSR7 : 30		CSR7SW : 12	Max. :0.43939
##	2016-06-30:22	(Other):137		(Other):297	

```
summary(buckvolboth)
```

```
##           date           plot    treatment      quad      volume
## 2016-06-24: 70   WSR10 :101   C:419    WSR10NE: 27   Min.    :0.01515
## 2016-06-23: 65   WSR9   : 89   H:412    WSR10NW: 26   1st Qu.:0.07576
## 2016-06-25: 58   WH12   : 88           WSR10SE: 26   Median :0.16364
## 2016-06-27: 56   CC6    : 76           EHSR1SW: 23   Mean   :0.28162
## 2016-06-29: 53   CH5    : 70           ESR2NE  : 23   3rd Qu.:0.38182
## 2016-06-26: 52   EH4    : 69           WH12NW  : 23   Max.   :3.03125
## (Other)      :477   (Other):338           (Other):683
```

```
summarize(group_by(buckvol15, treatment), meanVol = mean(volume), sdVolume = sd(volume))
```

```
## Source: local data frame [2 x 3]
##
##   treatment    meanVol  sdVolume
##   (fctr)      (dbl)    (dbl)
## 1          C 0.5224432 0.4286155
## 2          H 0.3653227 0.3151584
```

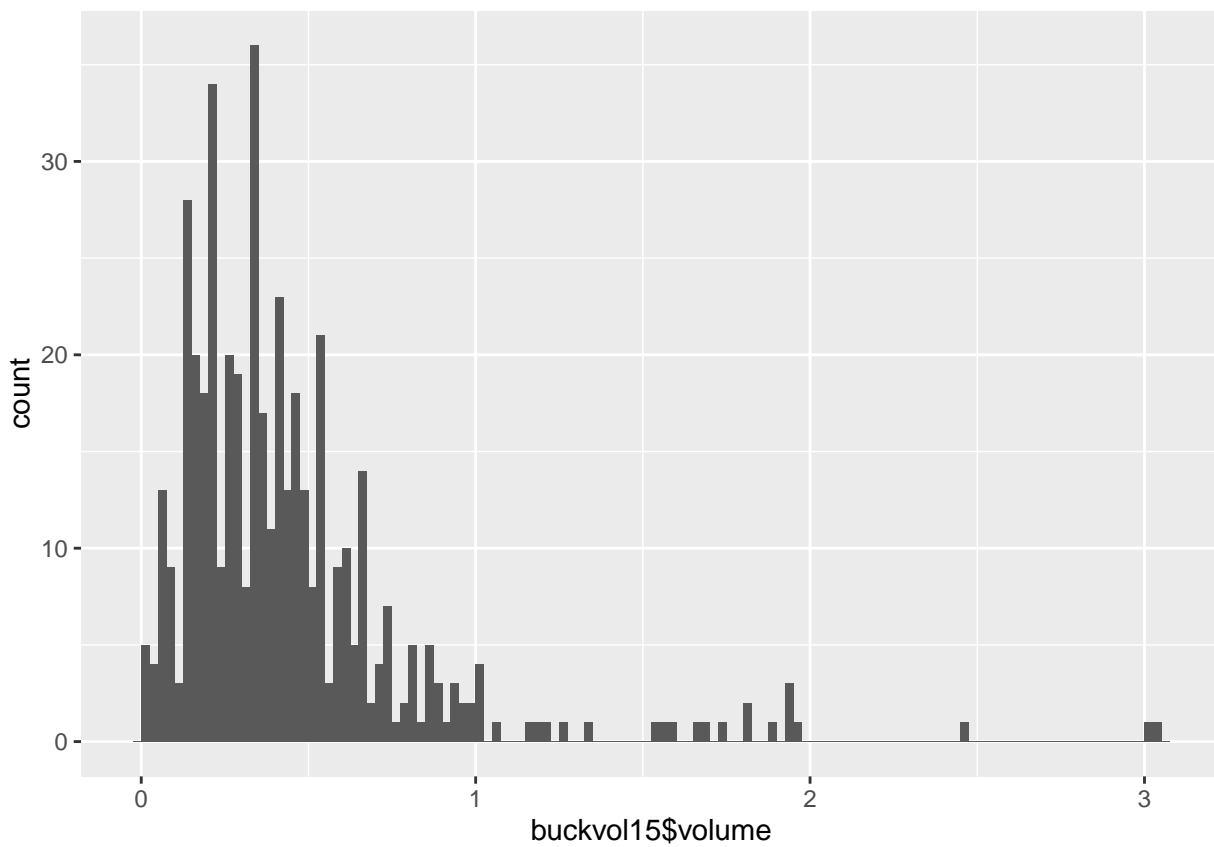
```
summarize(group_by(buckvol16, treatment), meanVol = mean(volume), sdVolume = sd(volume))
```

```
## Source: local data frame [2 x 3]
##
##   treatment    meanVol  sdVolume
##   (fctr)      (dbl)    (dbl)
## 1          C 0.08847687 0.06126738
## 2          H 0.08072662 0.04864575
```

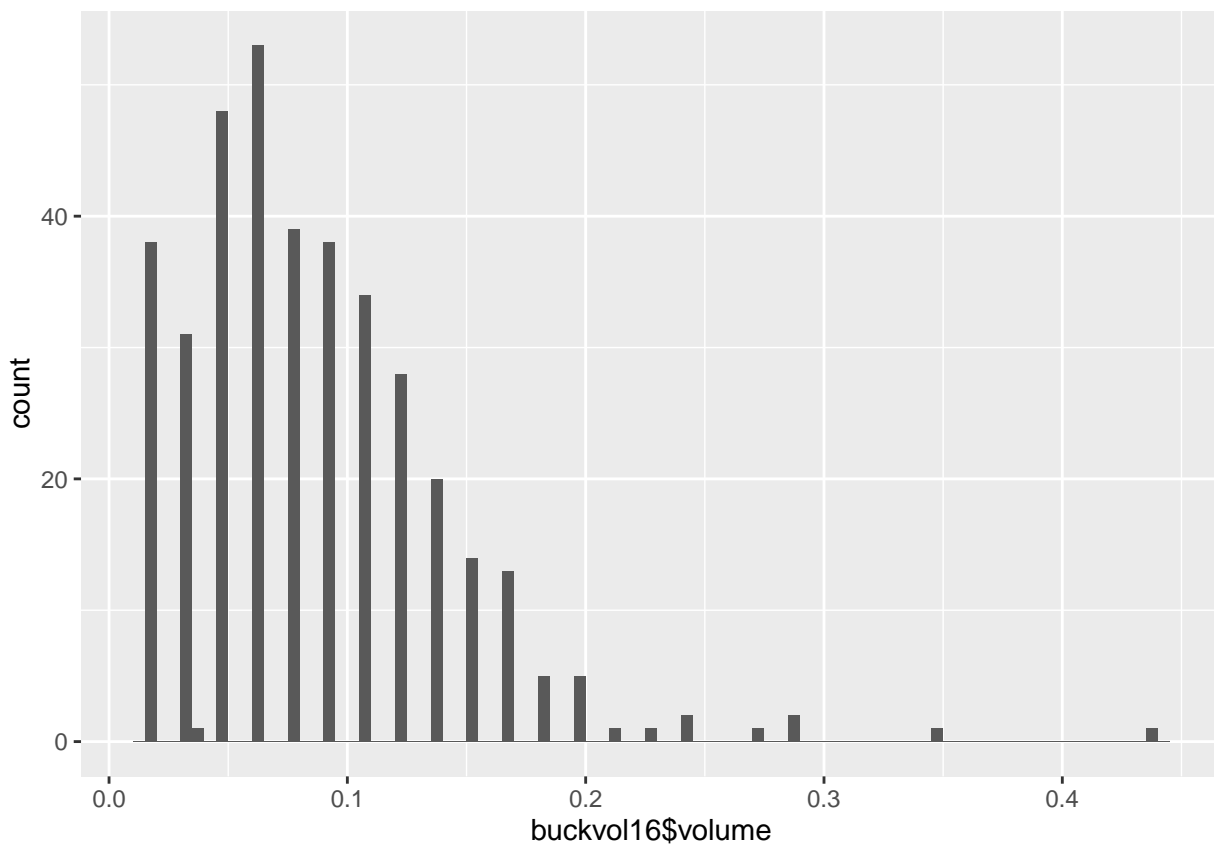
```
summarize(group_by(buckvolboth, treatment), meanVol = mean(volume), sdVolume = sd(volume))
```

```
## Source: local data frame [2 x 3]
##
##   treatment    meanVol  sdVolume
##   (fctr)      (dbl)    (dbl)
## 1          C 0.3256566 0.3856019
## 2          H 0.2368400 0.2748577
```

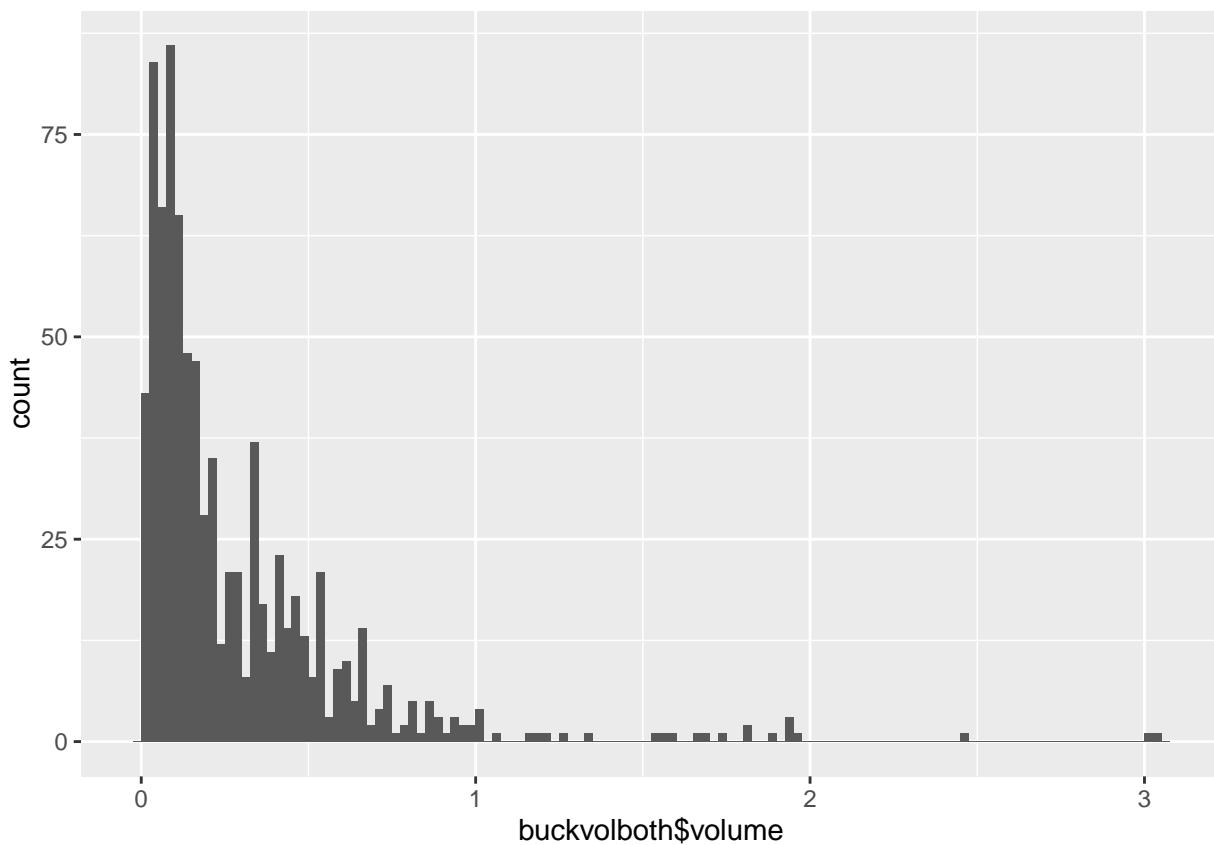
```
qplot(buckvol15$volume, binwidth = .025)
```



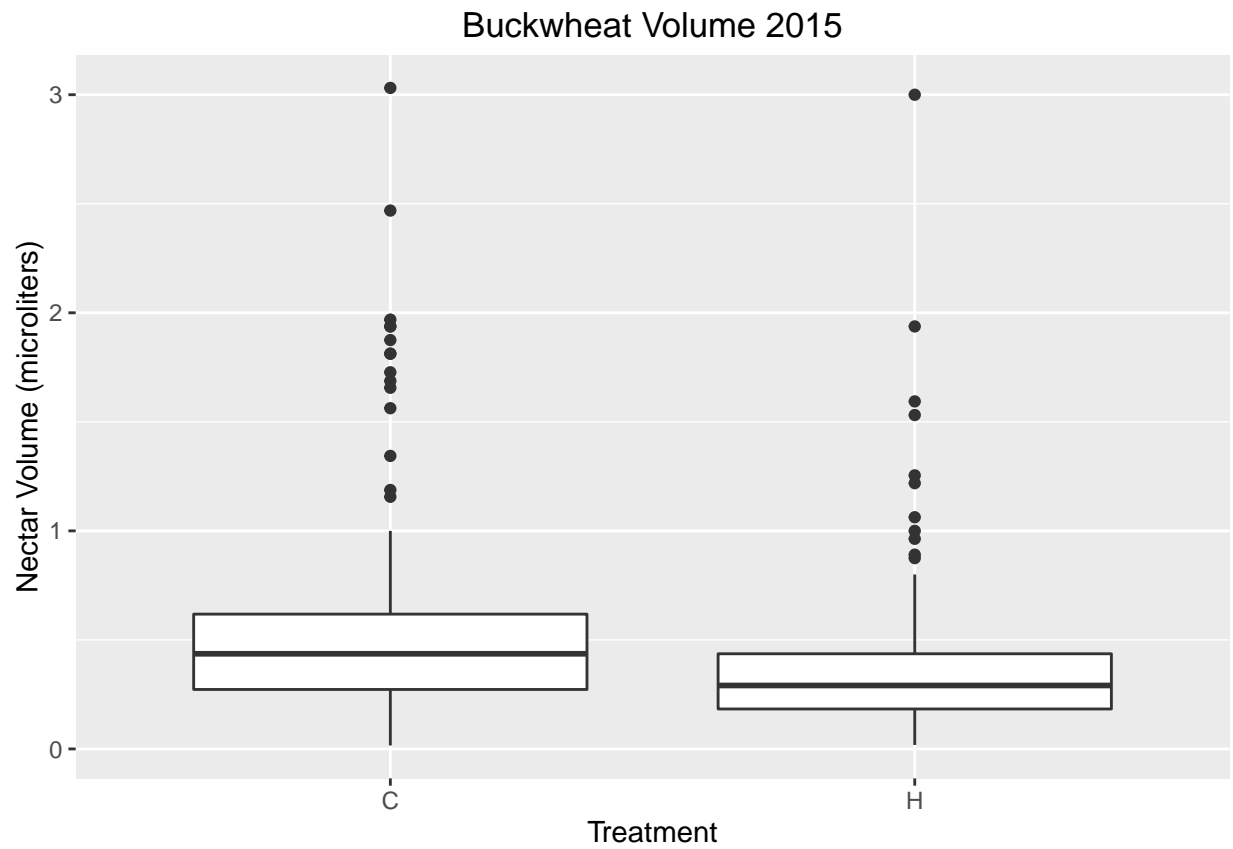
```
qplot(buckvol16$volume, binwidth = .005)
```



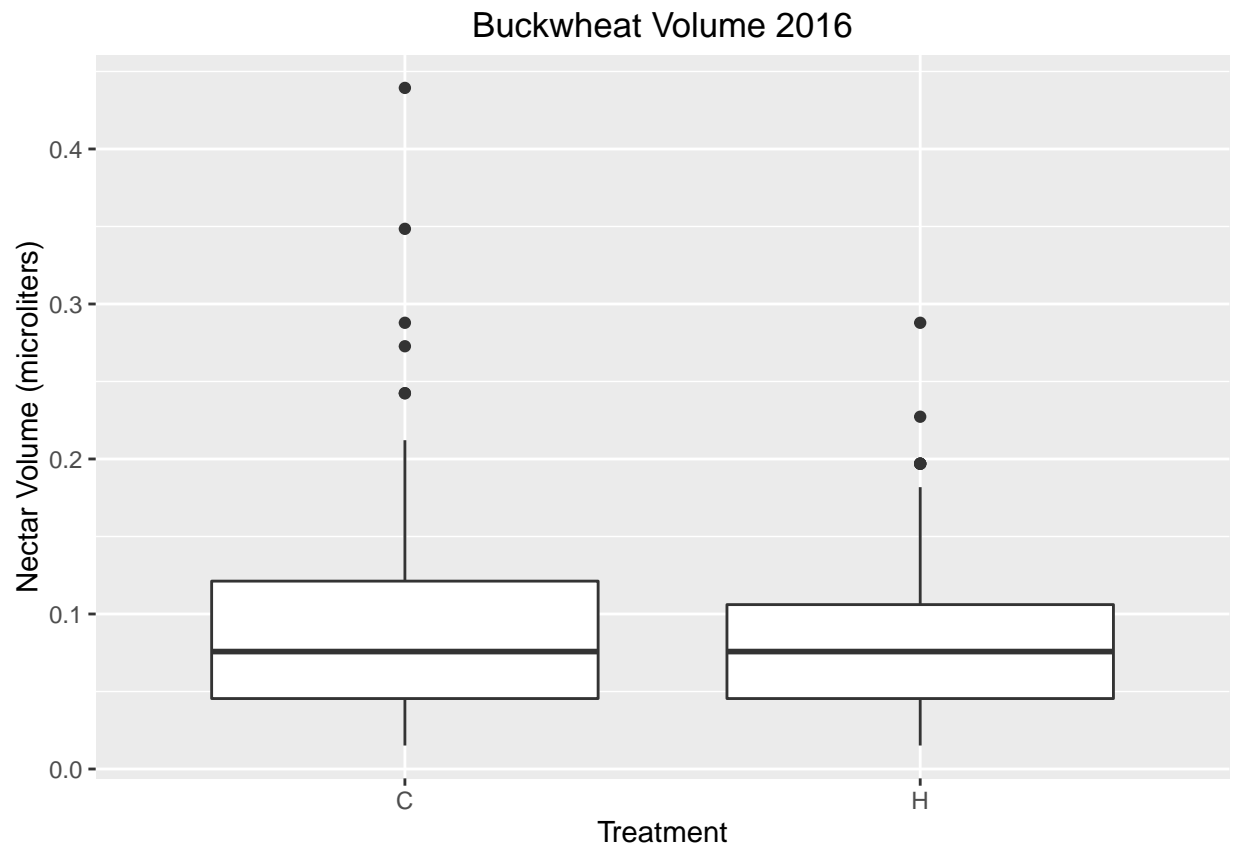
```
qplot(buckvolboth$volume, binwidth = .025)
```



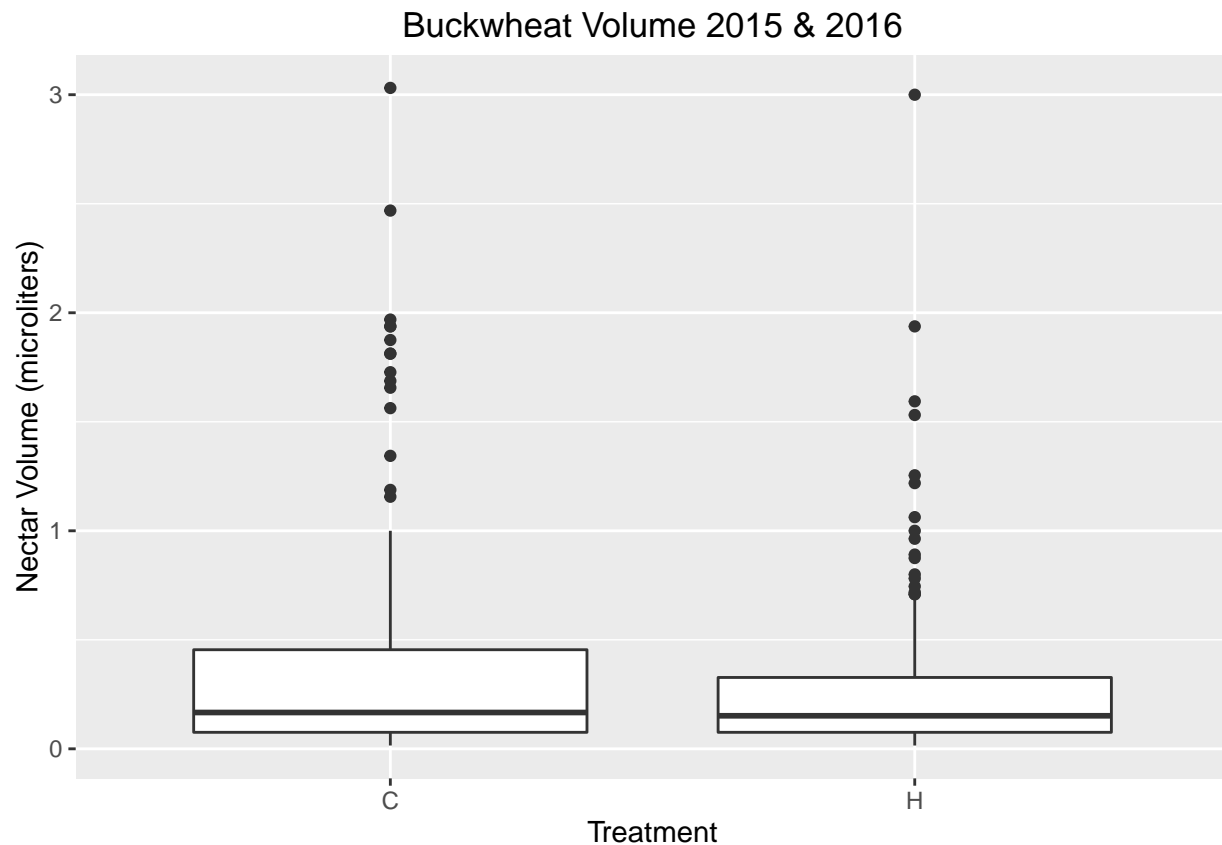
```
ggplot(buckvol15, aes(x=treatment, y=volume)) + geom_boxplot() +  
  xlab("Treatment") +  
  ylab("Nectar Volume (microliters)") + ggtitle("Buckwheat Volume 2015")
```



```
ggplot(buckvol16, aes(x=treatment, y=volume)) + geom_boxplot() +  
  xlab("Treatment") +  
  ylab("Nectar Volume (microliters)") + ggtitle("Buckwheat Volume 2016")
```



```
ggplot(buckvolboth, aes(x=treatment, y=volume)) + geom_boxplot() +  
  xlab("Treatment") +  
  ylab("Nectar Volume (microliters)") + ggtitle("Buckwheat Volume 2015 & 2016")
```



*# Homoscedastic?*

```
var15C <- sd(buckvol15$volume[buckvol15$treatment=="C"])^2
var15H <- sd(buckvol15$volume[buckvol15$treatment=="H"])^2
ratio15 <- var15C/var15H
ratio15
```

```
## [1] 1.849601
```

```
var16C <- sd(buckvol16$volume[buckvol16$treatment=="C"])^2
var16H <- sd(buckvol16$volume[buckvol16$treatment=="H"])^2
ratio16 <- var16C/var16H
ratio16
```

```
## [1] 1.586239
```

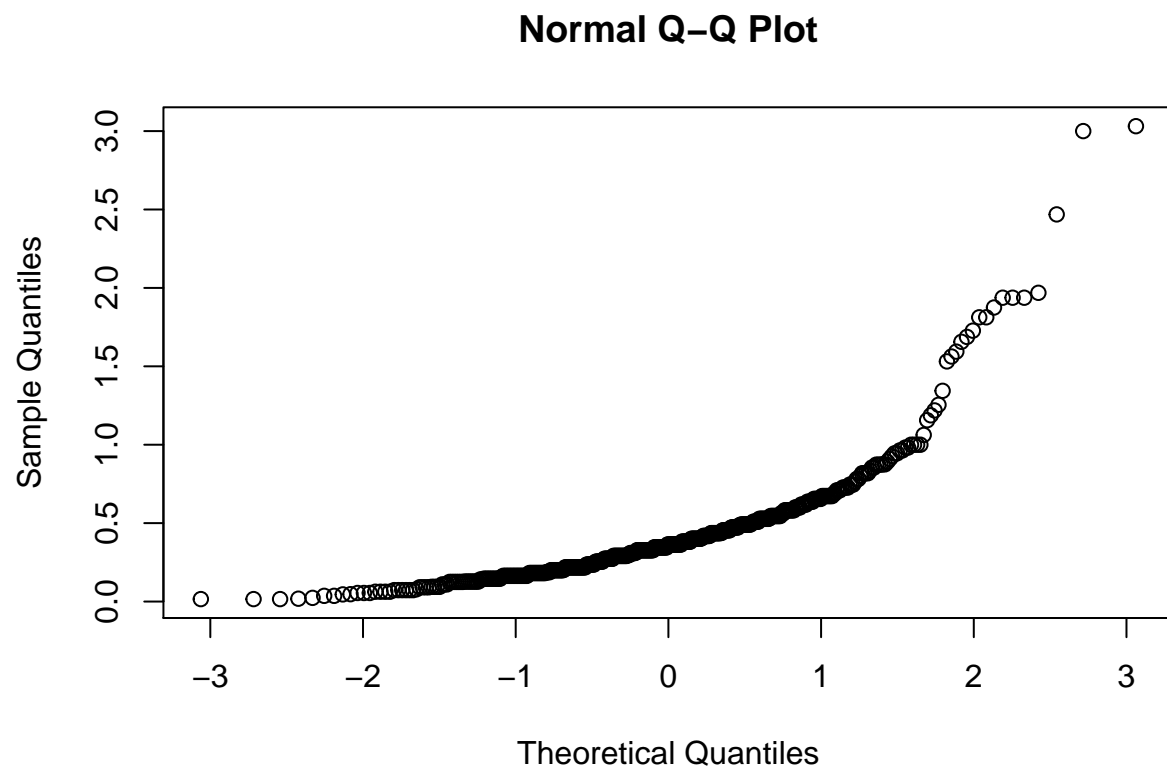
```
varbothC <- sd(buckvolboth$volume[buckvolboth$treatment=="C"])^2
varbothH <- sd(buckvolboth$volume[buckvolboth$treatment=="H"])^2
ratioboth <- varbothC/varbothH
ratioboth
```

```
## [1] 1.96817
```



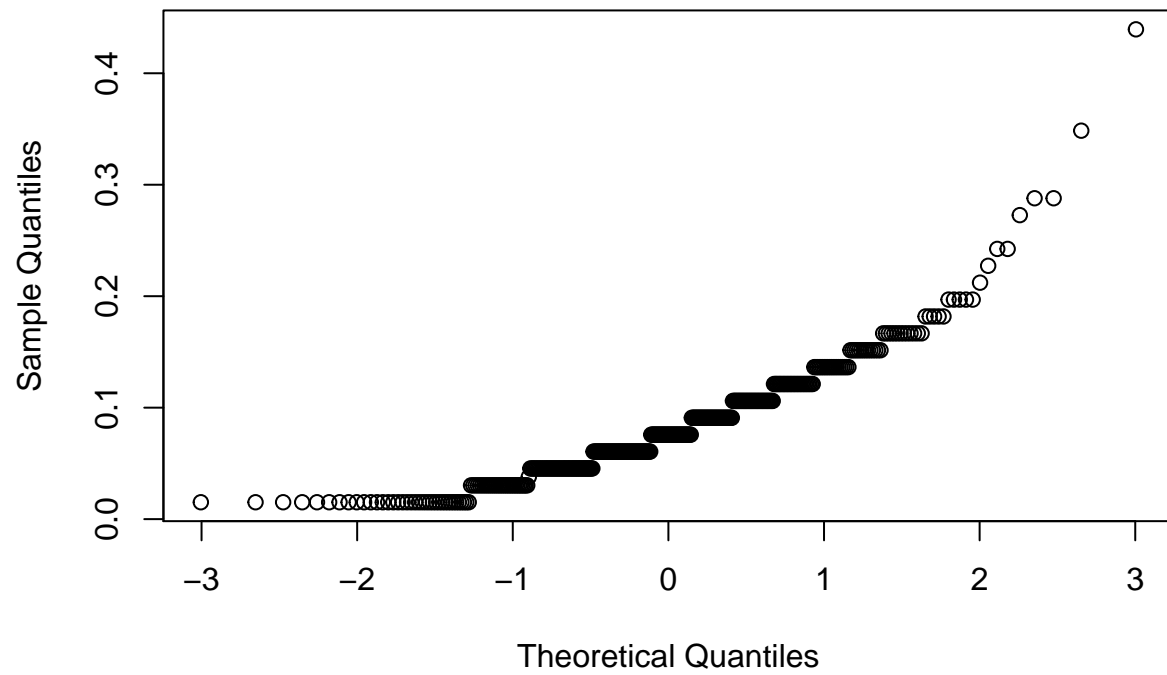
```
# Q-Q plots
```

```
qqnorm(buckvol15$volume)
```



```
qqnorm(buckvol16$volume)
```

Normal Q-Q Plot



```
qqnorm(buckvolboth$volume)
```

Normal Q-Q Plot

