

(there should probably be a figures title page here)

```
library(tidyverse)
library(readxl)
library(smmr)
```

Figure 1: Packages

```
treatment height
low      10.1
low      9.7
medium   4.8
medium   11.3
high     10.3
high     9.8
```

Figure 2: Data file stored in `biscuit1.txt`

```
treatment length
low 6.1
low 6.5
medium 8.7
medium 10.3
high 3.1
high 13.8
```

Figure 3: Data file stored in `biscuit2.txt`

```
test3 <- read_delim("biscuit2.txt")

## Rows: 6 Columns: 2
## -- Column specification -----
## Delimiter: " "
## chr (1): treatment
## dbl (1): length
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
The datafram as read in:
## # A tibble: 6 x 2
##   treatment length
##   <chr>     <dbl>
## 1 low       6.1
## 2 low       6.5
## 3 medium    8.7
## 4 medium   10.3
## 5 high      3.1
## 6 high     13.8
```

Figure 4: Alternative way of reading data file stored in `biscuit2.txt`

The screenshot shows a LibreOffice Calc spreadsheet window titled "precious.xlsx - LibreOffice Calc". The menu bar includes File, Edit, View, Insert, Format, Styles, Sheet, Data, and a toolbar with various icons. The font is set to "Liberation Sans" at 10 pt, and the bold and italic buttons are selected. The active cell is A6. The spreadsheet contains the following data:

	A	B	C
1	x	y	
2		1	2
3		3	4
4		5	6
5			
6			
7			
8			

The bottom navigation bar shows "Sheet1" and "Sheet2".

Figure 5: Spreadsheet to be read into R

```
eggs %>% slice(1:20)

## # A tibble: 20 x 3
##   day    weight day_number
##   <chr>   <dbl>     <dbl>
## 1 A         55       1
## 2 A         53       1
## 3 A         56       1
## 4 A         63       1
## 5 A         66       1
## 6 A         58       1
## 7 A         53       1
## 8 A         57       1
## 9 A         61       1
## 10 A        53       1
## 11 B        59       2
## 12 B        62       2
## 13 B        56       2
## 14 B        51       2
## 15 B        61       2
## 16 B        75       2
## 17 B        57       2
## 18 B        60       2
## 19 B        55       2
## 20 B        74       2
```

Figure 6: Egg weight data (first 20 rows)

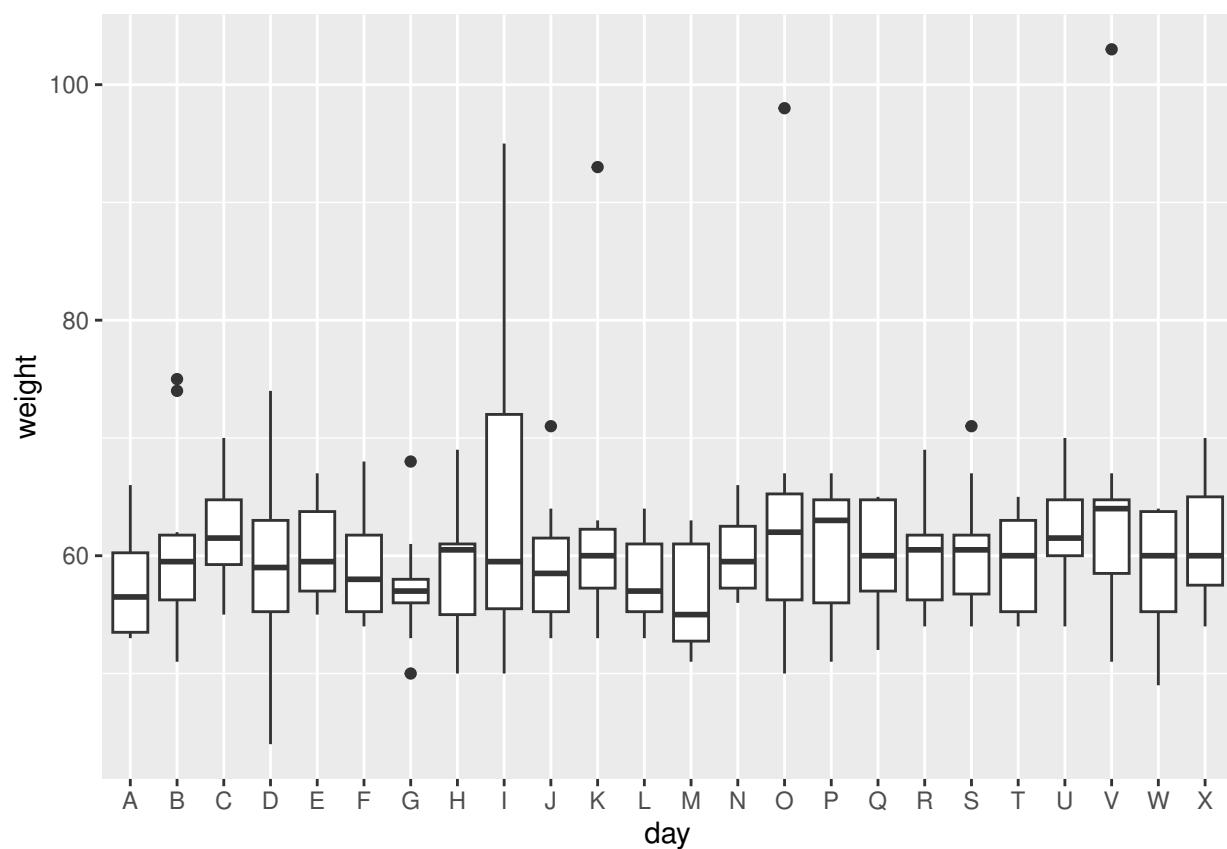


Figure 7: Graph of egg weight data

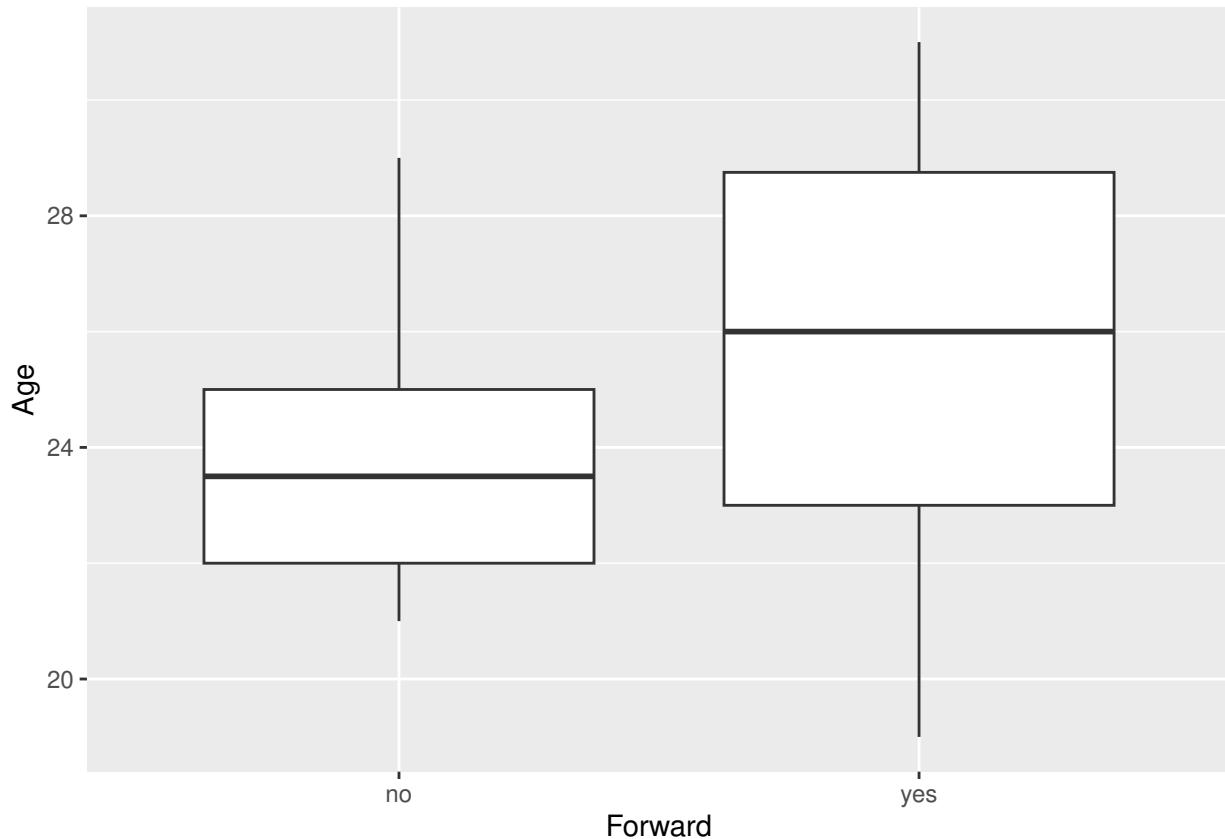
```
seattlepets

## # A tibble: 52,519 x 7
##   license_issue_date license_number animal_name species primary_breed
##   <date>           <chr>          <chr>        <chr>      <chr>
## 1 2018-11-16       8002756     Wall-E       Dog      Mixed Breed, Medium (u-
## 2 2018-11-11       S124529    Andre        Dog      Terrier, Jack Russell
## 3 2018-11-21       903793     Mac         Dog      Retriever, Labrador
## 4 2018-11-23       824666     Melb        Cat      Domestic Shorthair
## 5 2018-12-30       S119138    Gingersnap   Cat      Domestic Shorthair
## 6 2018-12-16       S138529    Cody         Dog      Retriever, Labrador
## 7 2017-10-04       580652     Millie       Dog      Terrier, Boston
## 8 2018-08-09       S142558    Sebastian    Cat      Domestic Shorthair
## 9 2018-08-20       S142546    Madeline     Cat      Domestic Shorthair
## 10 2018-12-08      S123830    Cleo         Cat      Domestic Shorthair
## # ... with 52,509 more rows, and 2 more variables: secondary_breed <chr>,
## #   zip_code <chr>
```

Figure 8: Seattle pets data (some)

```
## # A tibble: 26 x 8
##   Player            Position  Age Goals Assists PlusMinus PenMins Forward
##   <chr>           <chr>    <dbl> <dbl>  <dbl>    <dbl>    <dbl> <chr>
## 1 Chris Tierney    C          24    9     39     -22     26 yes
## 2 Magnus Paajarvi LW         27   11     8     -14      6 yes
## 3 Bobby Ryan       LW         31   15     27     -29     35 yes
## 4 Dylan DeMelo     D          25    4     18      -1     32 no
## 5 Cody Ceci        D          25    7     19     -22     18 no
## 6 Brady Tkachuk    LW         19   22     23     -10     75 yes
## 7 Colin White      C          22   14     27     -24     24 yes
## 8 Mikkel Boedker   RW         29    7     28     -23      6 yes
## 9 Thomas Chabot    D          22   14     41     -12     32 no
## 10 Zack Smith      C          30    9     19      -6     81 yes
## # ... with 16 more rows
```

Figure 9: Ottawa Senators data (some)



```
OttawaSenators2019 %>%
  count(Forward)

## # A tibble: 2 x 2
##   Forward     n
##   <chr>   <int>
## 1 no          8
## 2 yes         18
```

Figure 10: Boxplot by position and age, with other information

```
##
## Welch Two Sample t-test
##
## data: Age by Forward
## t = -1.4615, df = 19.779, p-value = 0.1596
## alternative hypothesis: true difference in means between group no and group yes is not equal to 0
## 95 percent confidence interval:
## -4.4855683 0.7911238
## sample estimates:
## mean in group no mean in group yes
## 23.87500 25.72222
```

Figure 11: *t*-test for Ottawa Senators data

```
tibble(sim = 1:1000) %>%
  rowwise() %>%
  mutate(my_sample = list(rnorm(15, 50, 9))) %>%
  mutate(t_test = list(t.test(my_sample, mu = 55))) %>%
  mutate(p_value = t_test$p.value) %>%
  count(p_value <= 0.05)

## # A tibble: 2 x 2
## # Rowwise:
##   `p_value <= 0.05`     n
##   <lg1>             <int>
## 1 FALSE              508
## 2 TRUE               492
```

Figure 12: A power analysis

```
tibble(sim = 1:1000) %>%
  rowwise() %>%
  mutate(my_sample = list(rnorm(30, 50, 9))) %>%
  mutate(t_test = list(t.test(my_sample, mu = 55))) %>%
  mutate(p_value = t_test$p.value) %>%
  count(p_value <= 0.05)

## # A tibble: 2 x 2
## # Rowwise:
##   `p_value <= 0.05`     n
##   <lg1>             <int>
## 1 FALSE              159
## 2 TRUE               841
```

Figure 13: A second power analysis

```
abbey
## # A tibble: 31 x 1
##       nickel
##   <dbl>
## 1     5.2
## 2     6.5
## 3     6.9
## 4      7
## 5      7
## 6      7
## 7    7.4
## 8      8
## 9      8
## 10     8
## # ... with 21 more rows
```

Figure 14: Nickel content data (some)

```
ggplot(abby, aes(x = nickel)) + geom_histogram(bins = 10)
```

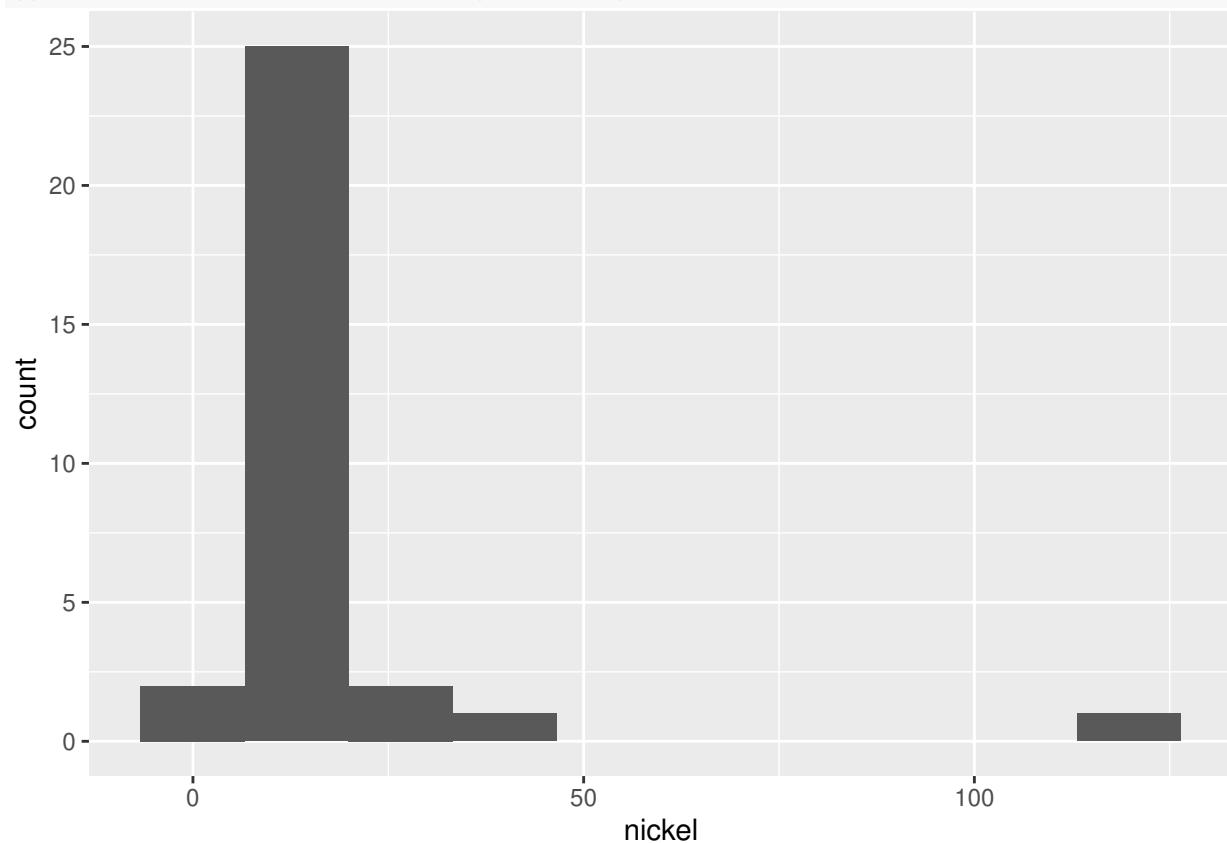


Figure 15: Nickel content histogram

```
tibble(sim = 1:1000) %>%
  rowwise() %>%
  mutate(my_sample = list(sample(abbey$nickel, replace = TRUE))) %>%
  mutate(my_mean = mean(my_sample)) %>%
  ggplot(aes(x = my_mean)) + geom_histogram(bins = 10)
```

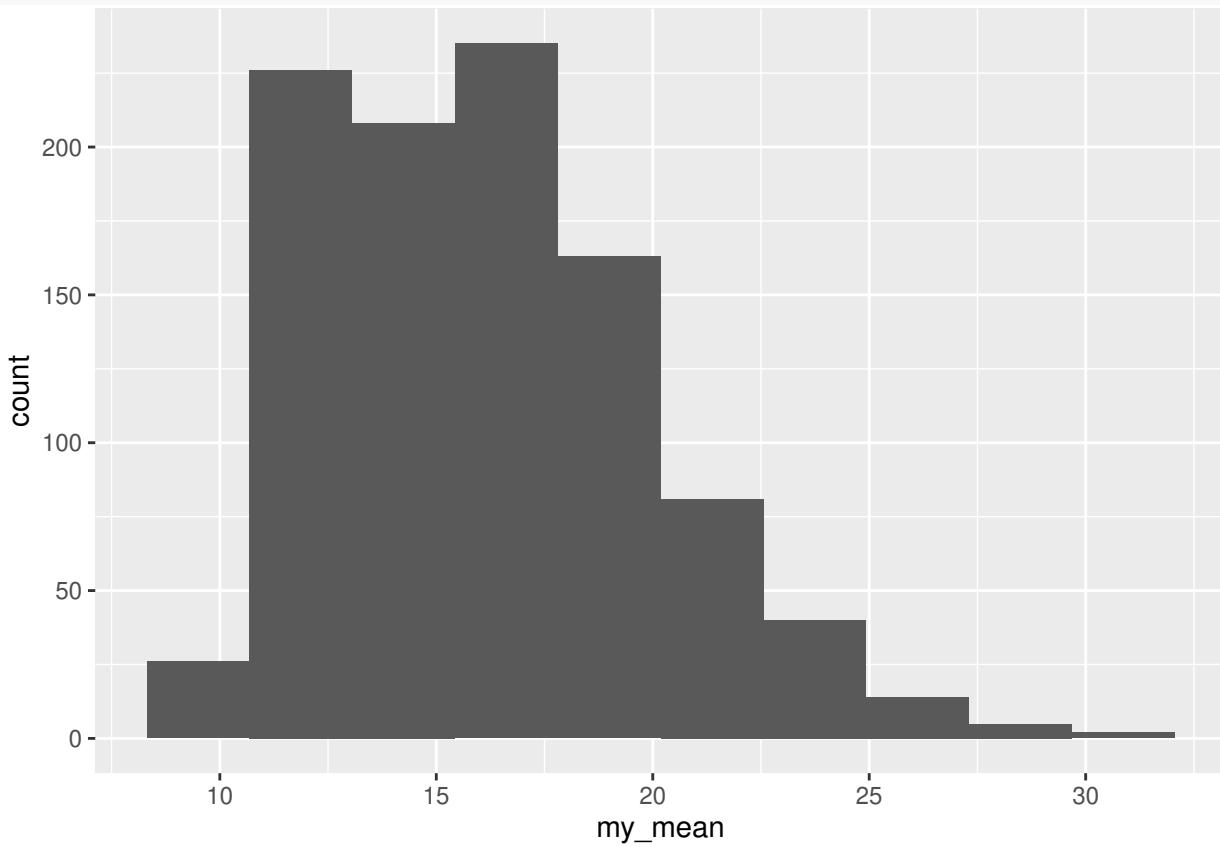


Figure 16: Nickel content further analysis

```
with(abbey, t.test(nickel))

##
##  One Sample t-test
##
## data: nickel
## t = 4.1901, df = 30, p-value = 0.0002259
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
##  8.204894 23.808009
## sample estimates:
## mean of x
## 16.00645
```

Figure 17: Nickel content confidence interval 1

```
ci_median(abbey, nickel)

## [1] 8.005078 13.997131
```

Figure 18: Nickel content confidence interval 2

```
sign_test(abbey, nickel, 15)

## $above_below
## below above
##      23      8
##
## $p_values
##   alternative   p_value
## 1          lower 0.00533692
## 2          upper 0.99833655
## 3 two-sided 0.01067384
```

Figure 19: A test