IS5 in R: Understanding and Comparing Distributions (Chapter 4)

Nicholas Horton (nhorton@amherst.edu)

2025-01-08

Table of contents

Introduction and background	1
Chapter 4: Understanding and Comparing Distributions	2
Section 4.1: Displays for Comparing Groups	3
Section 4.3: Re-Expressing Data: A First Look	11

Introduction and background

This document is intended to help describe how to undertake analyses introduced as examples in the Fifth Edition of *Intro Stats* (2018) by De Veaux, Velleman, and Bock. This file as well as the associated Quarto reproducible analysis source file used to create it can be found at http://nhorton.people.amherst.edu/is5.

This work leverages initiatives undertaken by Project MOSAIC (http://www.mosaic-web.org), an NSF-funded effort to improve the teaching of statistics, calculus, science and computing in the undergraduate curriculum. In particular, we utilize the mosaic package, which was written to simplify the use of R for introductory statistics courses. A short summary of the R needed to teach introductory statistics can be found in the mosaic package vignettes (https://cran.r-project.org/web/packages/mosaic). A paper describing the mosaic approach was published in the *R Journal*: https://journal.r-project.org/archive/2017/RJ-2017-024.

We begin by loading packages that will be required for our analyses.

```
library(mosaic)
library(tidyverse)
```

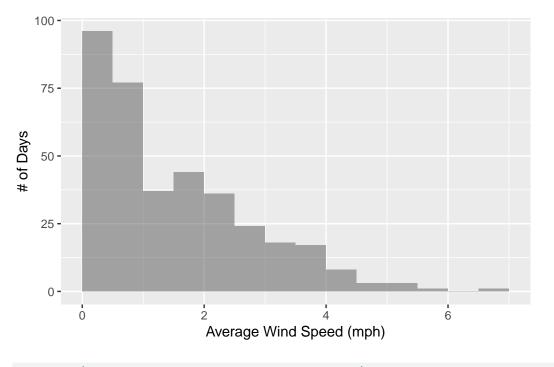
Chapter 4: Understanding and Comparing Distributions

```
library(mosaic)
library(readr)
library(janitor)
HopkinsForest <-
   read_csv("http://nhorton.people.amherst.edu/is5/data/Hopkins_Forest.csv") |>
   janitor::clean_names()
names(HopkinsForest)
```

[1]	"date"	"year"	"month"
[4]	"day"	"day_of_year"	"avg_temp_c"
[7]	"max_temp_c"	"min_temp_c"	"avg_temp_f"
[10]	"max_temp_f"	"min_temp_f"	"avg_rel_hum_percent"
[13]	"max_rel_hum_percent"	"min_rel_hum_percent"	"avg_sol_rad_w_m_2"
[16]	"max_sol_rad_w_m_2"	"min_sol_rad_w_m_2"	"total_sol_rad_w_m_2"
[19]	"avg_wind_mph"	"max_wind_mph"	"min_wind_mph"
[22]	"avg_barom_mb"	"max_barom_mb"	"min_barom_mb"
[25]	"precip_in"	"deep_well_ft"	"shallow_well_ft"
[28]	"x80_cm_soil_c"	"x10_cm_soil_c"	

By default, read_csv() prints the variable names. We suppressed these using the message = FALSE code chunk option to save space and improve readability. Here we use the clean_names() function from the janitor package to sanitize the names of the columns (which would otherwise contain special characters or whitespace). You can use the names() function to check the cleaned names.

```
# Figure 4.1, page 96
gf_histogram(~ avg_wind_mph,
    data = HopkinsForest,
    xlab = "Average Wind Speed (mph)",
    ylab = "# of Days",
    binwidth = 0.5,
    center = 0.25
)
```



df_stats(~ avg_wind_mph, data = HopkinsForest) # an alternative version of "favstats()"

response min Q1 median Q3 max mean sd n missing 1 avg_wind_mph 0 0.46 1.12 2.28 6.73 1.507808 1.260161 365 0

Section 4.1: Displays for Comparing Groups

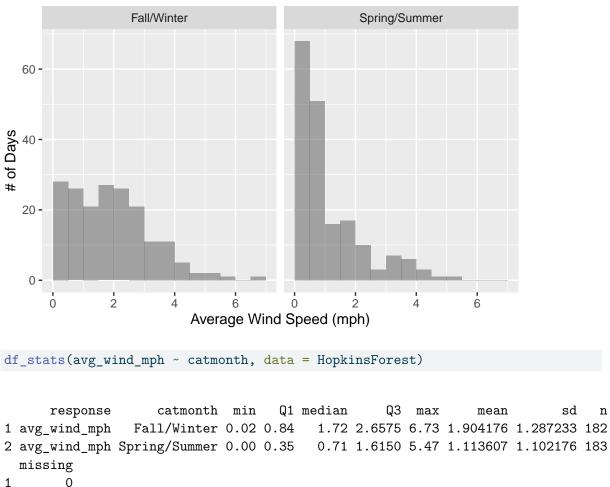
Histograms

We began by creating a new month to categorize the dates.

```
HopkinsForest <- HopkinsForest |>
mutate(catmonth = ifelse(
    month <= 9 & month >= 4,
    "Spring/Summer",
    "Fall/Winter")
)
```

```
# Figure 4.2, page 96
gf_histogram(~ avg_wind_mph,
    data = HopkinsForest, binwidth = 0.5, center = 0.25,
    xlab = "Average Wind Speed (mph)", ylab = "# of Days"
```

) |> gf_facet_wrap(~ catmonth)



1 2 0

Example 4.1: Comparing Groups with Stem-And-Leaf

We begin by reading in the data.

```
# Figure 4.1, page 97
NestEgg <- read_csv("http://nhorton.people.amherst.edu/is5/data/Nest_Egg_Index.csv") |>
  janitor::clean_names()
with(NestEgg, stem(nest_egg_index)) # or stem(NestEgg$nest_egg_index)
```

n

```
The decimal point is 1 digit(s) to the right of the |

8 | 57789

9 | 0123344

9 | 667777888899

10 | 0012233333344

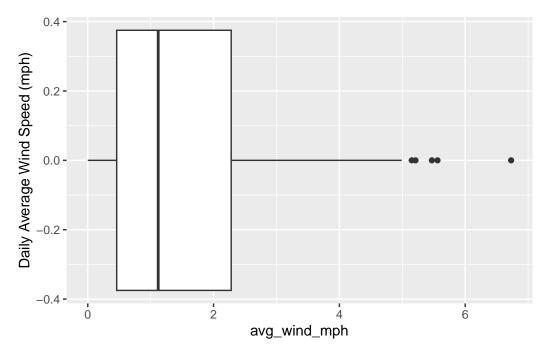
10 | 5566779
```

11 | 122444

Boxplots

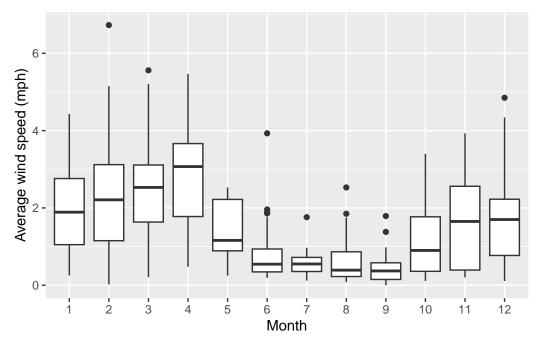
As noted in the book, boxplots are most useful to compare distributions. Below, we have replicated the single boxplot from page 98.

```
# Step 4 on page 98
gf_boxplot(~ avg_wind_mph, data = HopkinsForest) |> # or gf_boxplot(X ~ 1)
gf_labs(y = "Daily Average Wind Speed (mph)")
```



The use of single boxplots isn't recommended. Instead, one can make comparisons more easily by placing boxplots side by side with the following code:

```
# Figure 4.3, page 99
gf_boxplot(avg_wind_mph ~ as.factor(month), data = HopkinsForest) |>
gf_labs(x = "Month", y = "Average wind speed (mph)")
```



We use the as.factor() function to convert a variable into a factor.

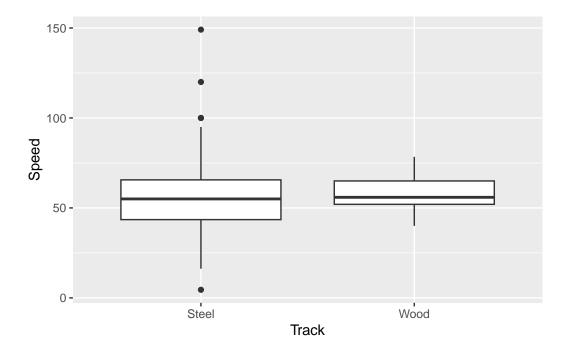
We also use gf_labs() to clean up the code for the first line and improve readability.

Here we use the mosaic modeling language to specify the variables. The ~ symbol is used to separate the response variable from the explanatory variable.

As a general form, GOAL(Y ~ X) carries out a specific goal for Y as a function of X.

Example 4.2: Comparing Groups with Boxplots

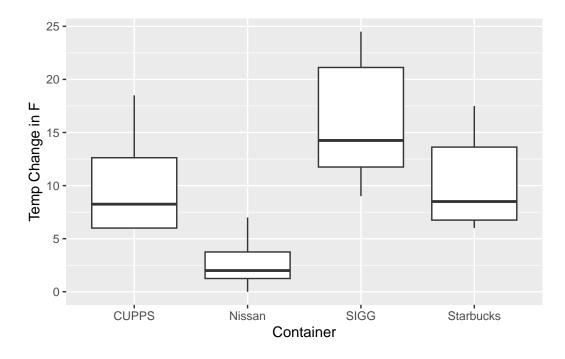
```
# Example 4.2, page 99
Coasters <- read_csv("http://nhorton.people.amherst.edu/is5/data/Coasters_2015.csv")
gf_boxplot(Speed ~ Track, data = Coasters)</pre>
```



Step-By-Step Example: Comparing Groups

```
Cups <- read_csv("http://nhorton.people.amherst.edu/is5/data/Cups.csv")
df_stats(Difference ~ Container, data = Cups)</pre>
```

```
Q1 median
    response Container min
                                            Q3 max
                                                                  sd n missing
                                                       mean
                                   8.25 12.625 18.5 10.1875 5.202592 8
1 Difference
                 CUPPS
                         6
                            6.00
                                                                             0
2 Difference
                         0 1.25
                                   2.00 3.750 7.0 2.7500 2.507133 8
                                                                             0
                Nissan
3 Difference
                  SIGG
                         9 11.75 14.25 21.125 24.5 16.0625 5.900590 8
                                                                             0
4 Difference Starbucks
                         6 6.75
                                   8.50 13.625 17.5 10.2500 4.551295 8
                                                                             0
# Mechanics, page 101
gf_boxplot(Difference ~ Container, data = Cups, ylab = "Temp Change in F")
```



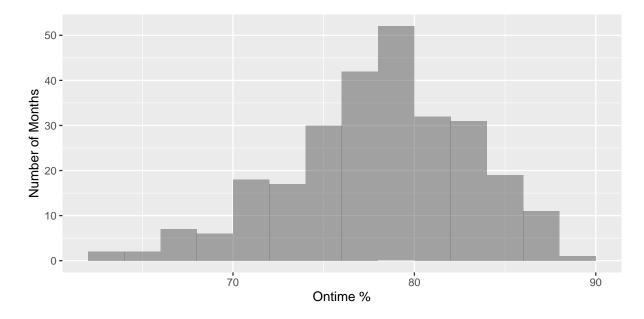
Just Checking

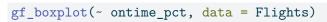
We begin by reading in the data.

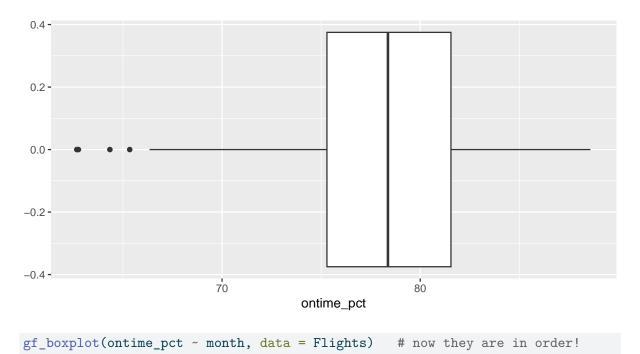
```
Flights <-
  read_csv("http://nhorton.people.amherst.edu/is5/data/Flights_on_time_2016.csv") |>
  janitor::clean_names()
# Let's improve the ordering of the months (by default they are alphabetical!)
Flights <- Flights |>
  mutate(month = forcats::fct_relevel(
    month,
    "January", "February", "March", "April",
    "May", "June", "July", "August",
    "September", "October", "November", "December"
    )
)
```

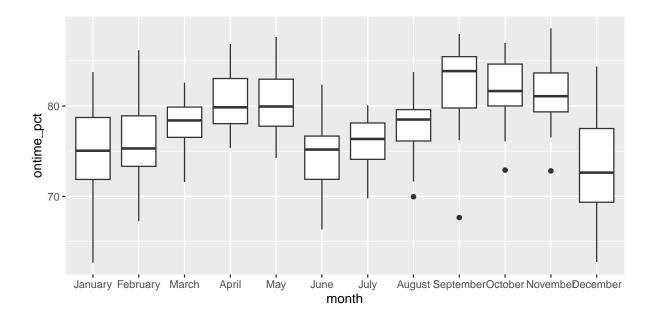
Here we use the fct_relevel() function from the forcats package to reorder the months in the dataset (the default is that the months are ordered alphabetically, which isn't very helpful). This function is a very useful idiom to remember when you want to reorder factor levels.

```
# Bureau of Transportation Statistics, page 101
gf_histogram(~ ontime_pct, data = Flights, binwidth = 2, center = 1) |>
gf_labs(x = "Ontime %", y = "Number of Months")
```



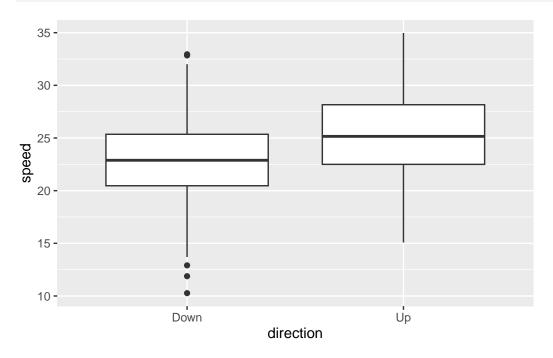






Random Matters

```
# Figure 4.4, page 102
CarSpeeds <- read_csv("http://nhorton.people.amherst.edu/is5/data/Car_speeds.csv")
gf_boxplot(speed ~ direction, data = CarSpeeds)</pre>
```

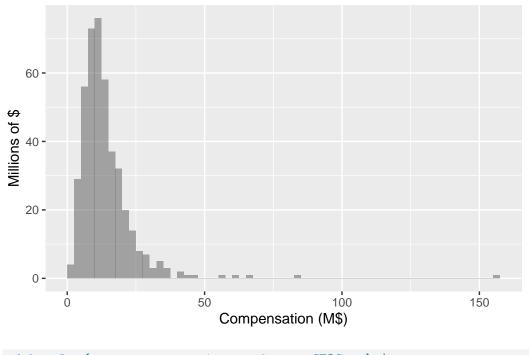


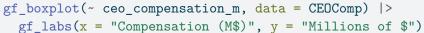
Section 4.3: Re-Expressing Data: A First Look

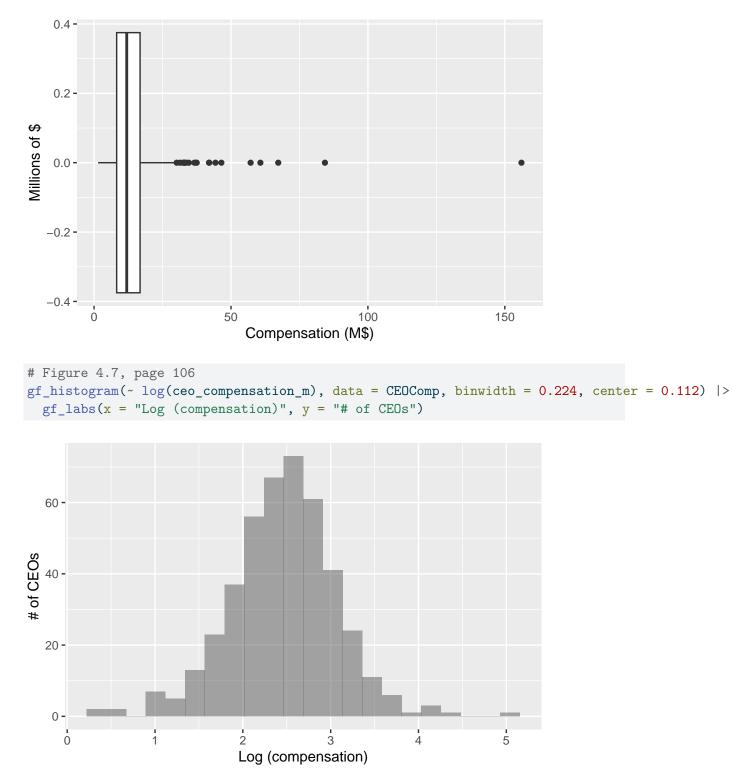
Re-Expressing to Improve Symmetry

```
CEOComp <-
read_csv("http://nhorton.people.amherst.edu/is5/data/CEO_Compensation_2014.csv") |>
janitor::clean_names()
```

```
# Figure 4.6, page 105
gf_histogram(~ ceo_compensation_m, data = CEOComp, binwidth = 2.5, center = 2.5 / 2) |>
gf_labs(x = "Compensation (M$)", y = "Millions of $")
```







Here we needed to pick magic numbers for the binwidth (e.g., 0.224) and centering of the

histogram so that it matched the results from the book.

Re-Expression to Equalize Spread Across Groups

We begin by reading in the data.

PassiveSmoke <- read_csv("http://nhorton.people.amherst.edu/is5/data/Passive_smoke.csv")</pre>

```
# Figure 4.8, page 107
gf_boxplot(cotinine ~ smoke_exposure, data = PassiveSmoke) |>
gf_labs(x = "Smoke Exposure", y = "Cotinine (ng/ml)")
```

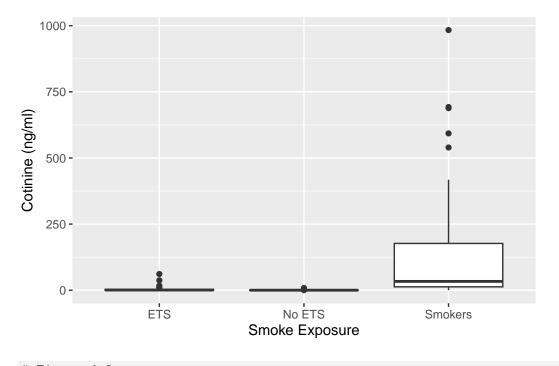


Figure 4.9
gf_boxplot(log(cotinine) ~ smoke_exposure, data = PassiveSmoke) |>
gf_labs(x = "Smoke Exposure", y = "Log(cotinine)")

