

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

Nicholas Horton (nhorton@amherst.edu)

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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 R Markdown 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>.

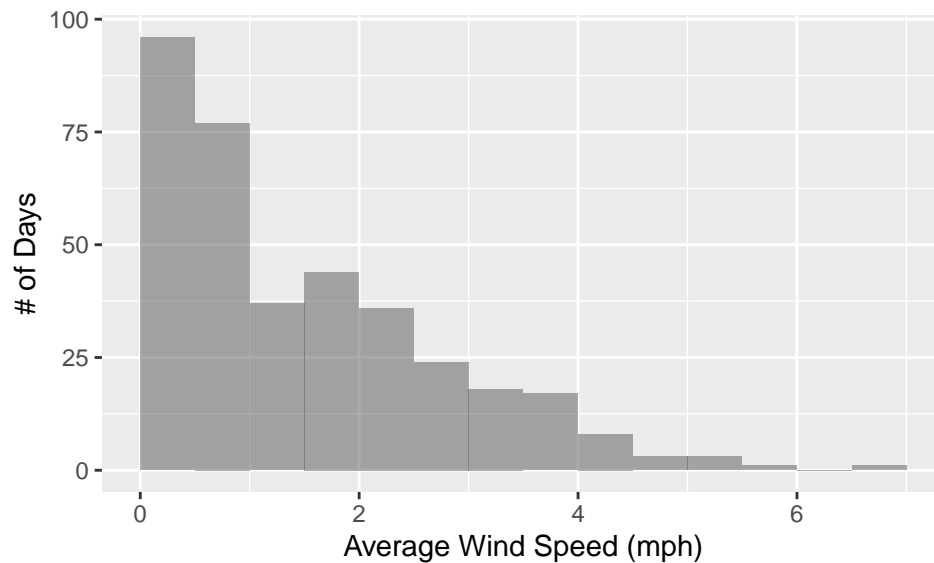
Chapter 4: Understanding and Comparing Distributions

```
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 improved 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
```

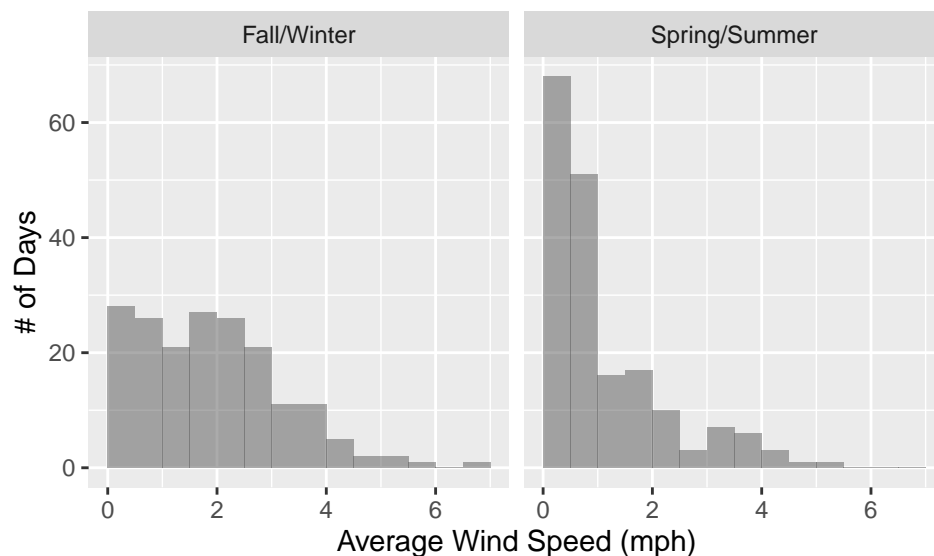
Section 4.1: Displays for Comparing Groups

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)
```



```
df_stats(avg_wind_mph ~ catmonth, data = HopkinsForest)
```

```
##      response      catmonth min   Q1 median   Q3 max   mean      sd   n
## 1 avg_wind_mph  Fall/Winter 0.02 0.84   1.72 2.6575 6.73 1.904176 1.287233 182
## 2 avg_wind_mph Spring/Summer 0.00 0.35   0.71 1.6150 5.47 1.113607 1.102176 183
## missing
## 1      0
## 2      0
```

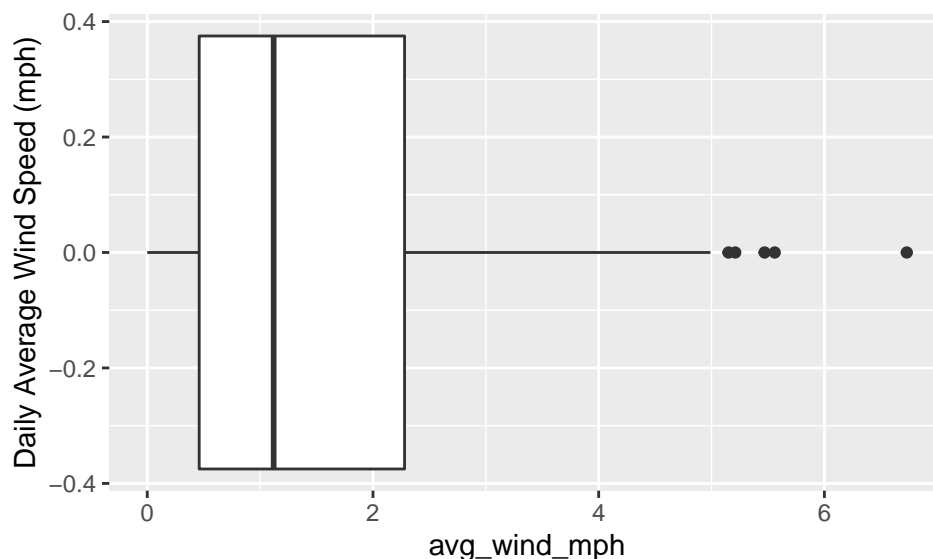
Example 4.1: Comparing Groups with Stem-And-Leaf We begin by reading in the data.

```
# 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))
```

```
##
## 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.

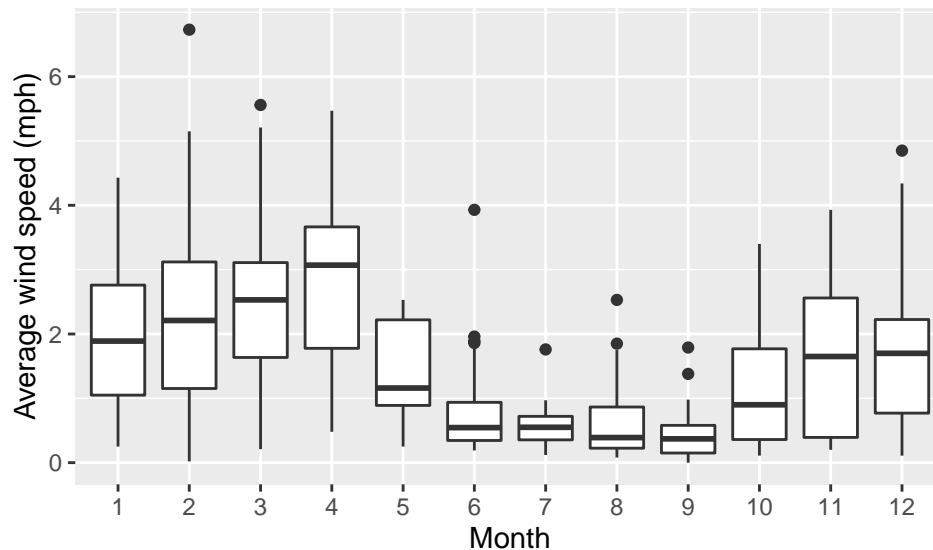
```
# Step 4 on page 98
gf_boxplot(~avg_wind_mph, data = HopkinsForest) %>% # or gf_boxplot(X ~ 1)
  gf_labs(y = "Daily Average Wind Speed (mph)")
```



I don't recommend the use of single boxplots. 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.

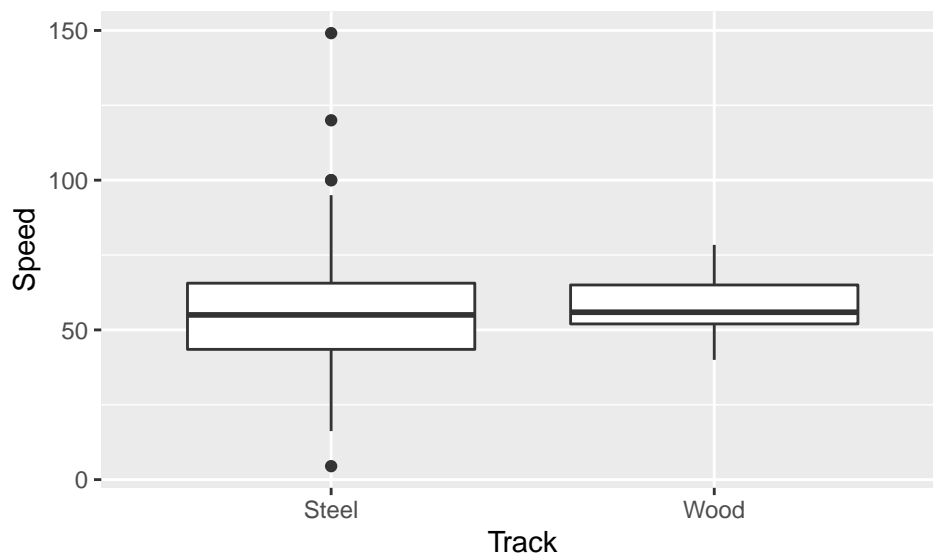
We also use `gf_labs()` to clean up the code for the first line and improve readability.

Here we use the mosaic modeling language to specify the variables. 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 We begin by reading in the data.

Example 4.2, page 99

```
Coasters <- read_csv("http://nhorton.people.amherst.edu/is5/data/Coasters_2015.csv")
gf_boxplot(Speed ~ Track, data = Coasters)
```



Step-By-Step Example: Comparing Groups We begin by reading in the data.

```

Cups <- read_csv("http://nhorton.people.amherst.edu/is5/data/Cups.csv")

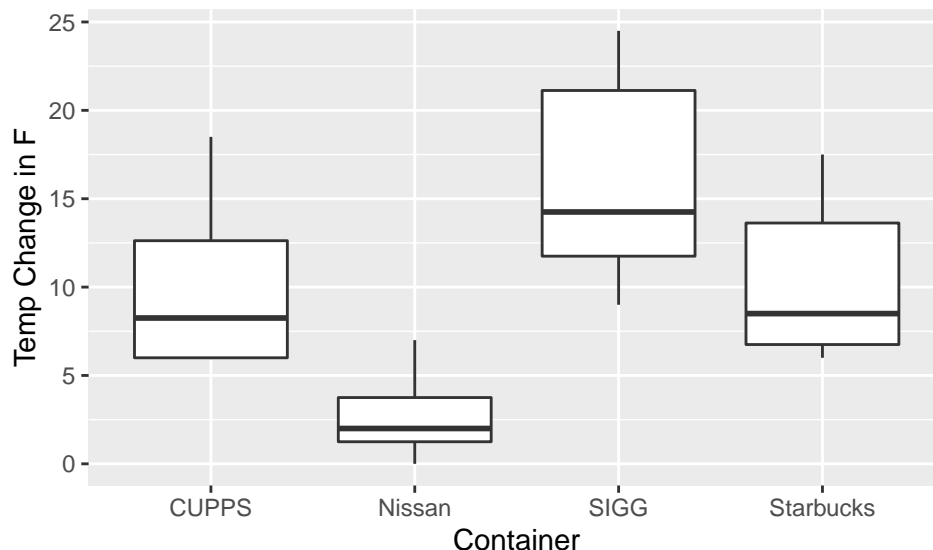
##
## -- Column specification -----
## cols(
##   Difference = col_double(),
##   Container = col_character()
## )

df_stats(Difference ~ Container, data = Cups)

##      response Container min    Q1 median    Q3 max    mean    sd n missing
## 1 Difference    CUPPS   6  6.00   8.25 12.625 18.5 10.1875 5.202592 8      0
## 2 Difference   Nissan   0  1.25   2.00  3.750   7.0  2.7500 2.507133 8      0
## 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")

```

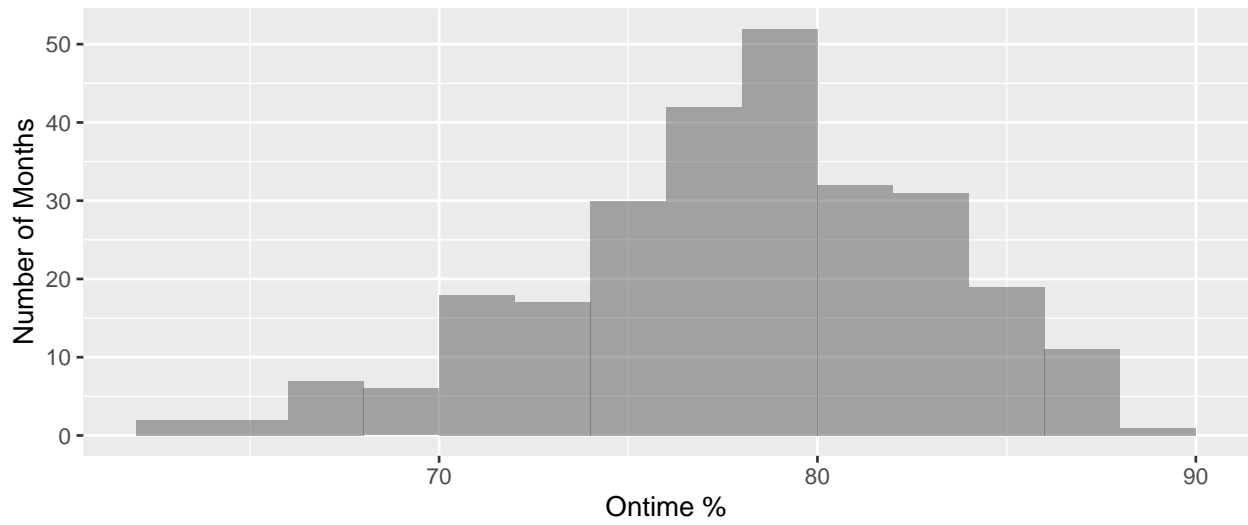


Just Checking We begin by reading in the data.

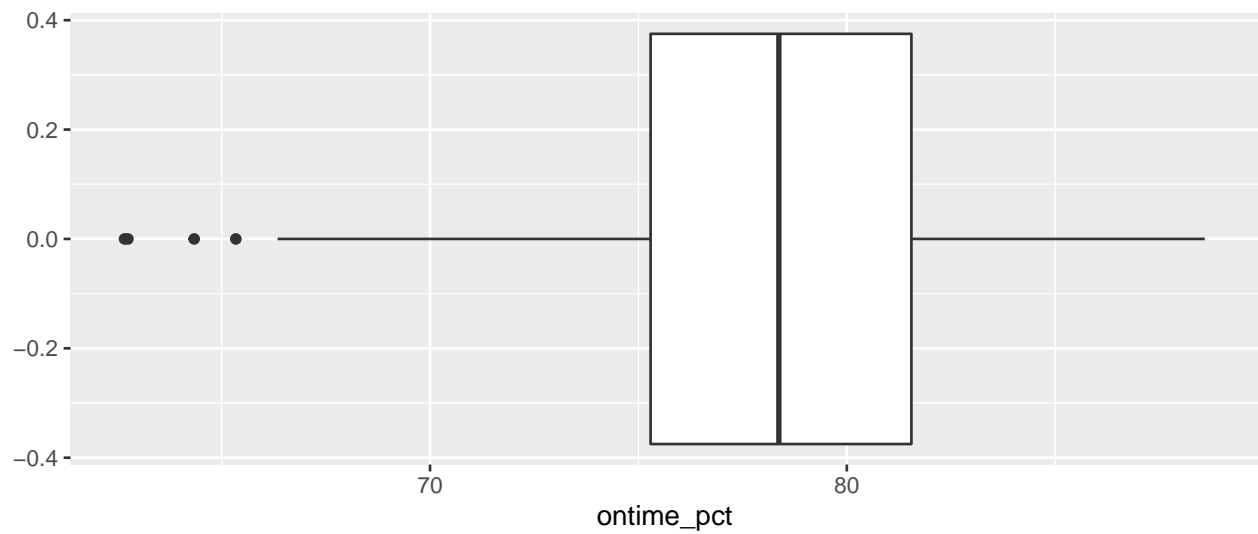
```

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

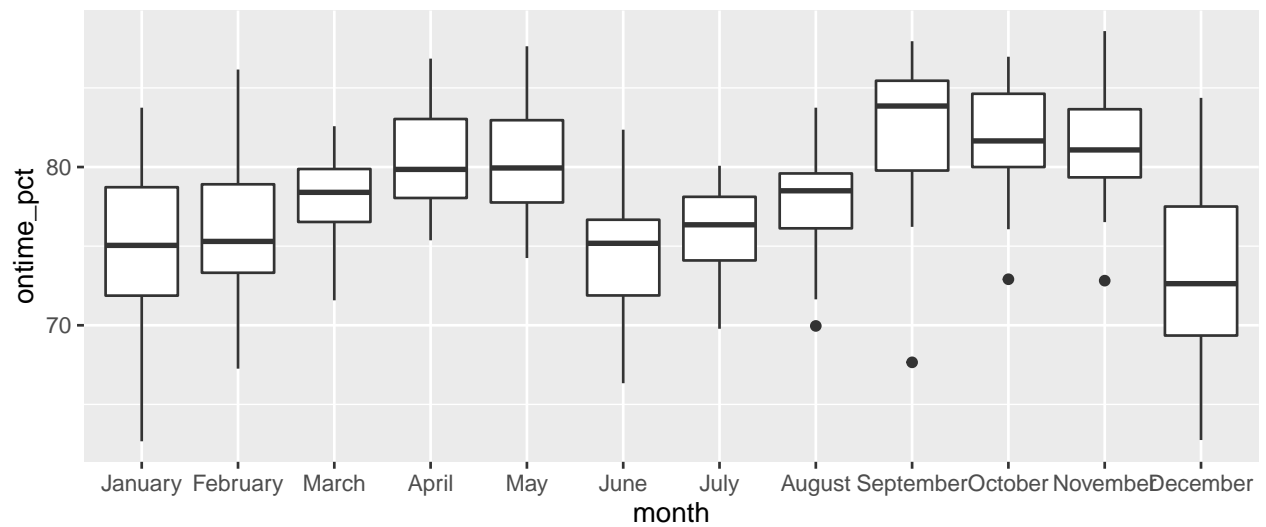
```



```
gf_boxplot(~ontime_pct, data = Flights)
```



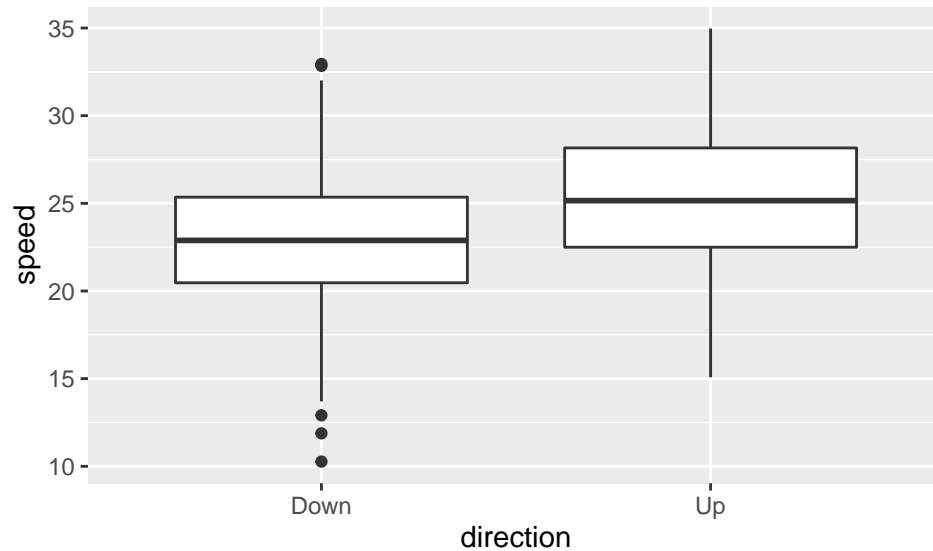
```
gf_boxplot(ontime_pct ~ month, data = Flights) # should be in order!
```



Random Matters We begin by reading in the data.

Figure 4.4, page 102

```
CarSpeeds <- read_csv("http://nhorton.people.amherst.edu/is5/data/Car_speeds.csv")
gf_boxplot(speed ~ direction, data = CarSpeeds)
```



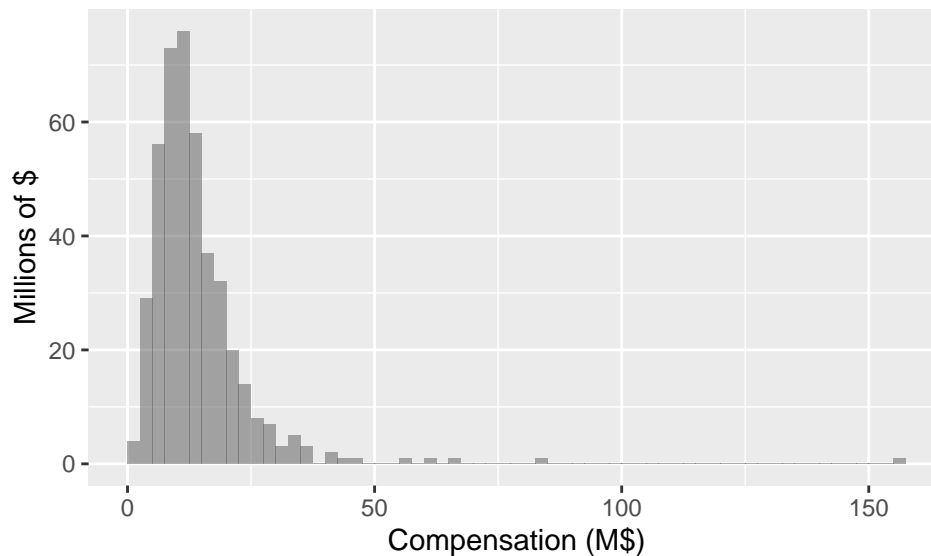
Section 4.3: Re-Expressing Data: A First Look

Re-Expressing to Improve Symmetry We begin by reading in the data.

```
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 $")
```



```
gf_boxplot(~ceo_compensation_m, data = CEOComp) %>%
  gf_labs(x = "Compensation (M$)", y = "Millions of $")
```

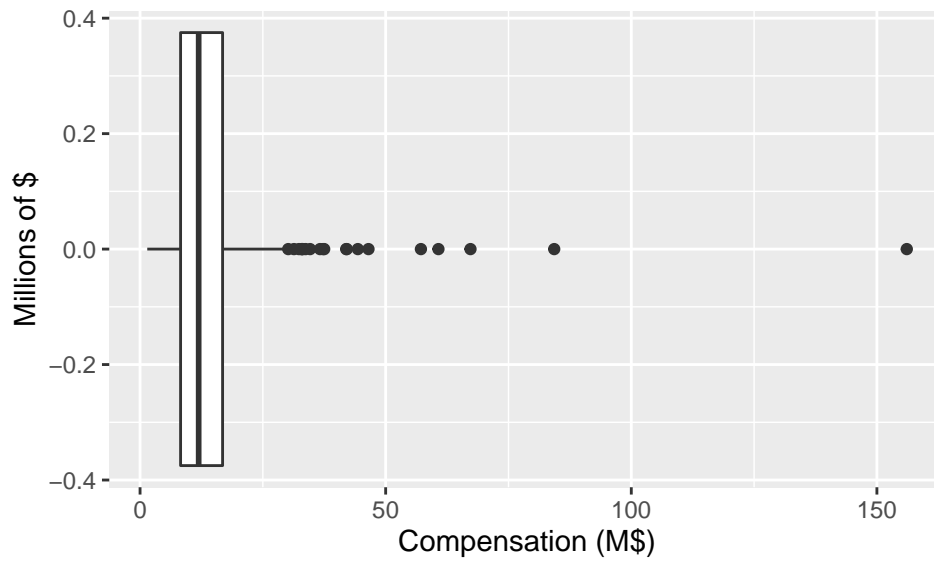
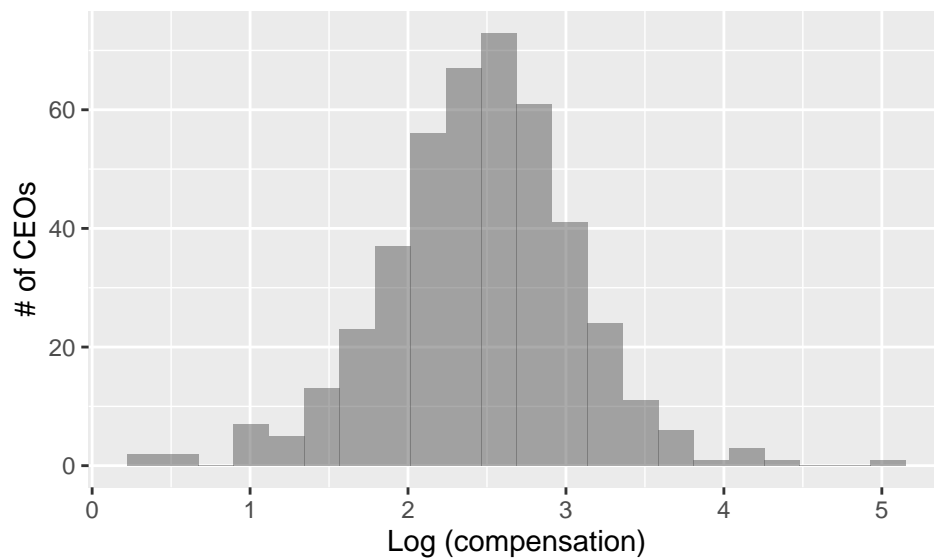


Figure 4.7, page 106

```
gf_histogram(~ log(ceo_compensation_m), data = CEOComp, binwidth = .224, center = .112) %>%
  gf_labs(x = "Log (compensation)", y = "# of CEOs")
```



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")
```

```
##
## -- Column specification -----
## cols(
##   cotinine = col_double(),
##   smoke_exposure = col_character()
## )
```

Figure 4.8, page 107

```
gf_boxplot(cotinine ~ smoke_exposure, data = PassiveSmoke) %>%
  gf_labs(x = "Smoke Exposure", y = "Cotinine (ng/ml)")
```

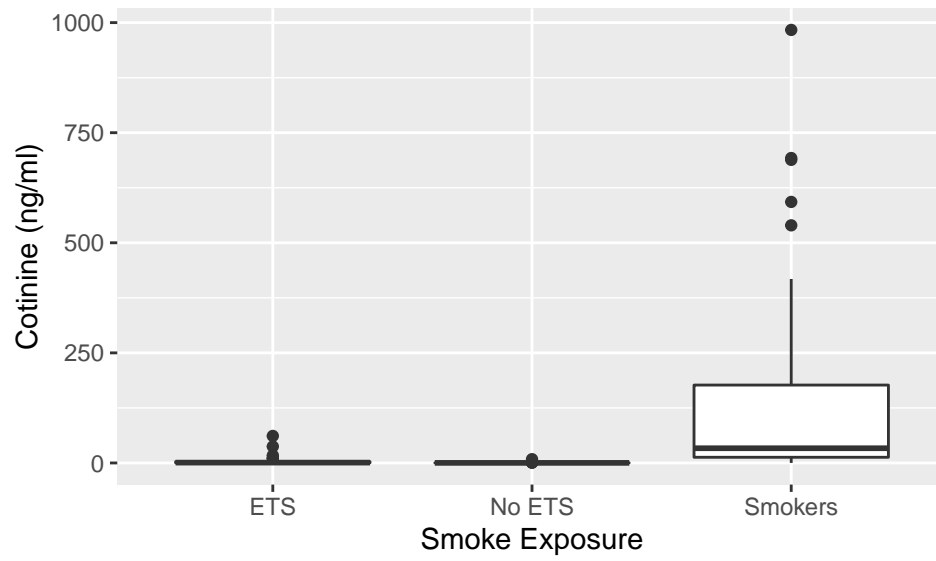



Figure 4.9

```
gf_boxplot(log(cotinine) ~ smoke_exposure, data = PassiveSmoke) %>%  
  gf_labs(x = "Smoke Exposure", y = "Log(cotinine)")
```

