

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

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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>.

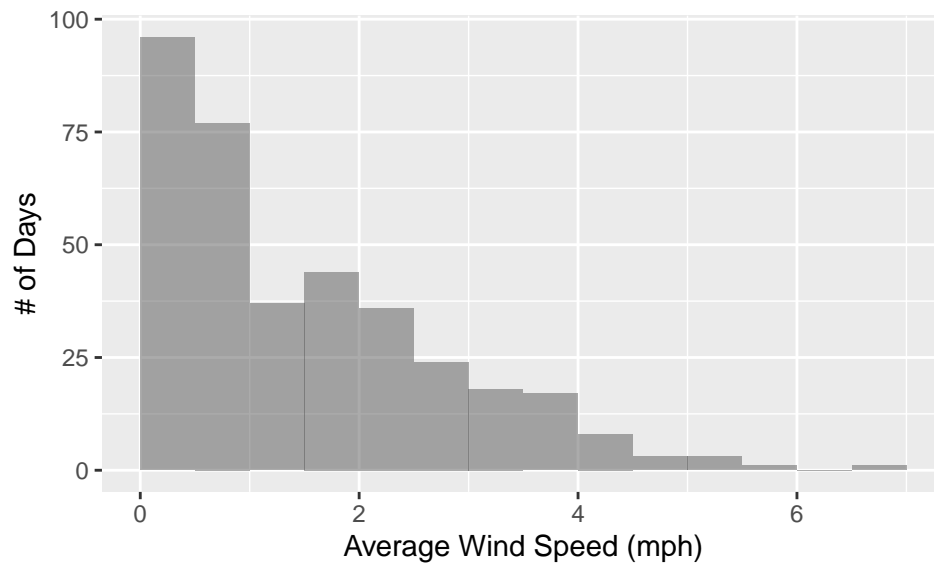
Chapter 4: Understanding and Comparing Distributions

```
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
)
```



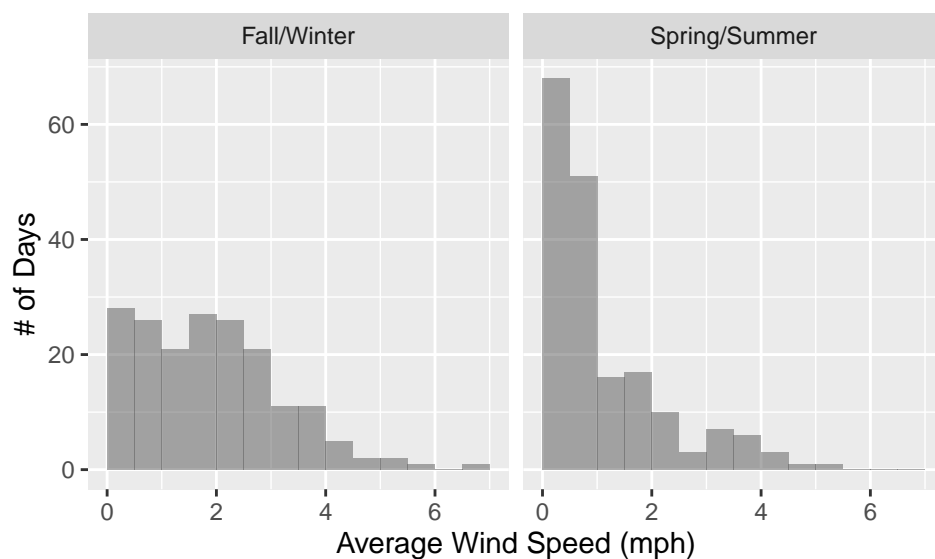
```
df_stats(~avg_wind_mph, data = HopkinsForest) # an improved version of "favstats()"
```

```
##      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

```
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)
```



Histograms

```
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
```

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))
```

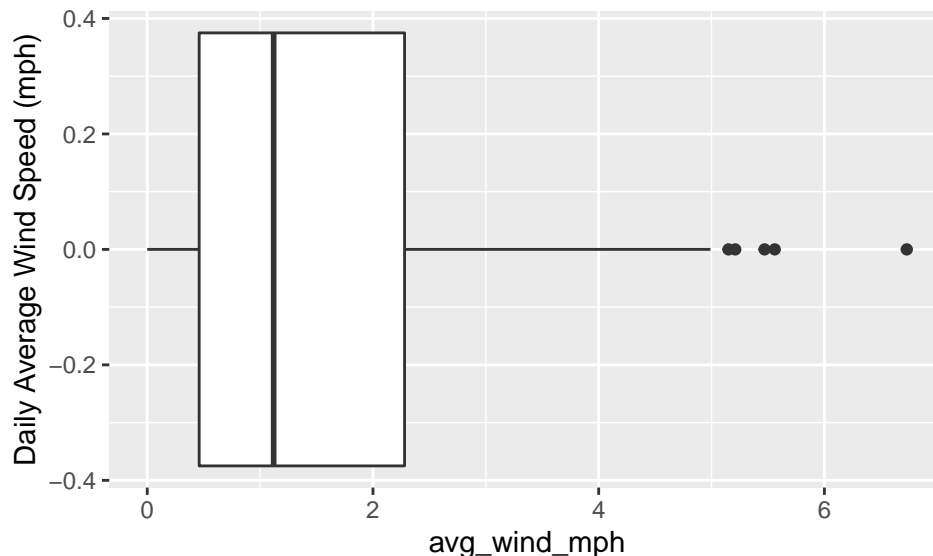
Example 4.1: Comparing Groups with Stem-And-Leaf

```
##
## 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

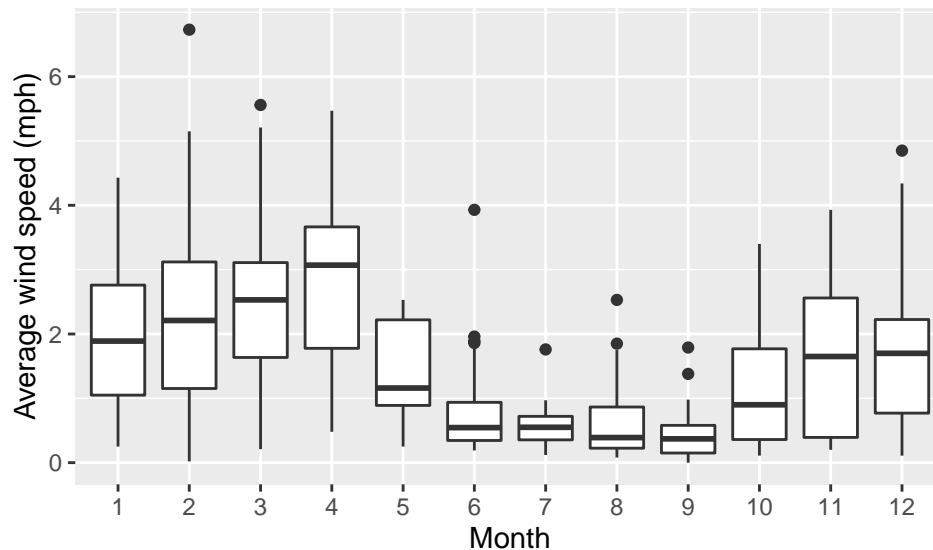
```
gf_boxplot(~avg_wind_mph, data = HopkinsForest) %>% # or gf_boxplot(X ~ 1)
  gf_labs(y = "Daily Average Wind Speed (mph)")
```



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.

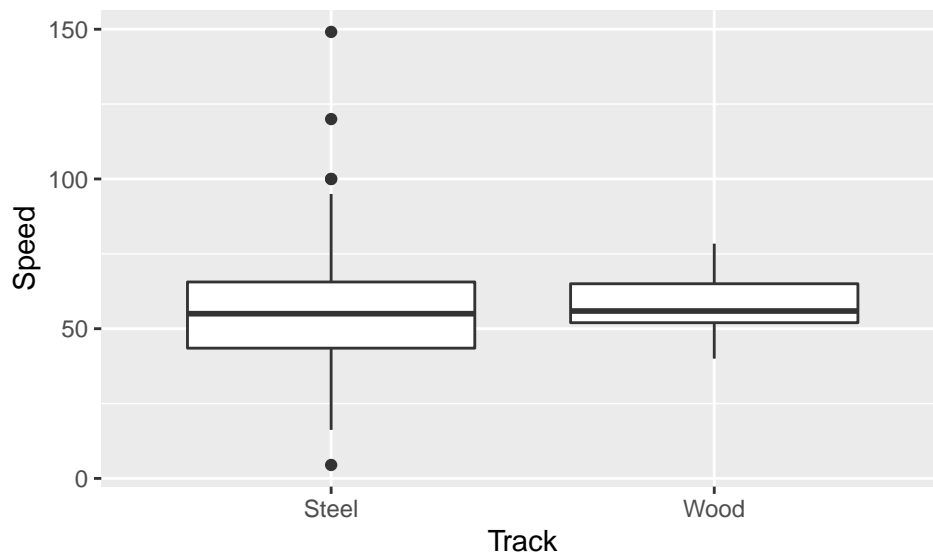
Here we use the mosaic modeling language to specify the variables. As a general form, `GOAL(Y ~ X)` carries out a specific goal for Y as a function of X.

Example 4.2, page 99

```
Coasters <- read_csv("http://nhorton.people.amherst.edu/is5/data/Coasters_2015.csv")
```

Example 4.2: Comparing Groups with Boxplots

```
##
## -- Column specification -----
## cols(
##   Name = col_character(),
##   Park = col_character(),
##   Track = col_character(),
##   Speed = col_double(),
##   Height = col_double(),
##   Drop = col_double(),
##   Length = col_double(),
##   Duration = col_double(),
##   Inversions = col_double()
## )
gf_boxplot(Speed ~ Track, data = Coasters)
```



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

Step-By-Step Example: Comparing Groups

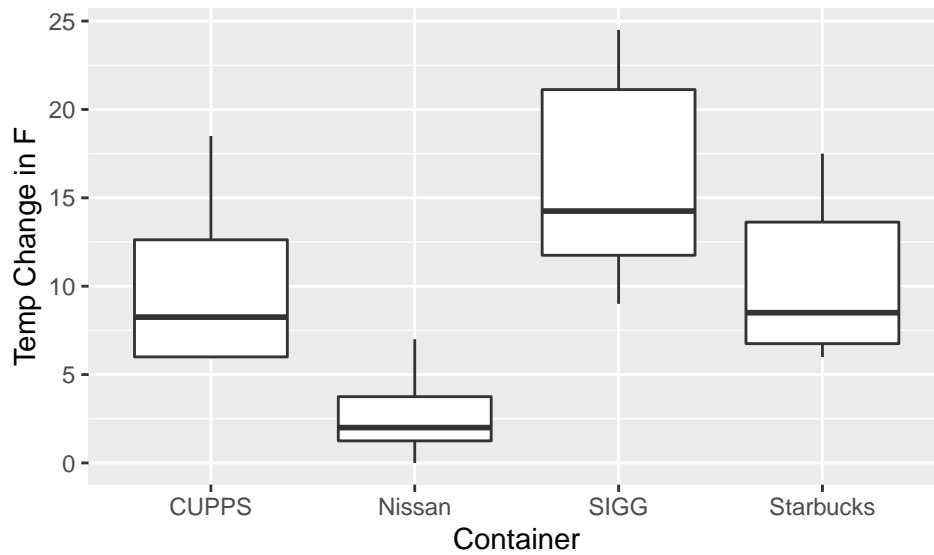
```
##
## -- Column specification -----
## cols(
##   Difference = col_double(),
##   Container = col_character()
## )
```

```
df_stats(Difference ~ Container, data = Cups)
```

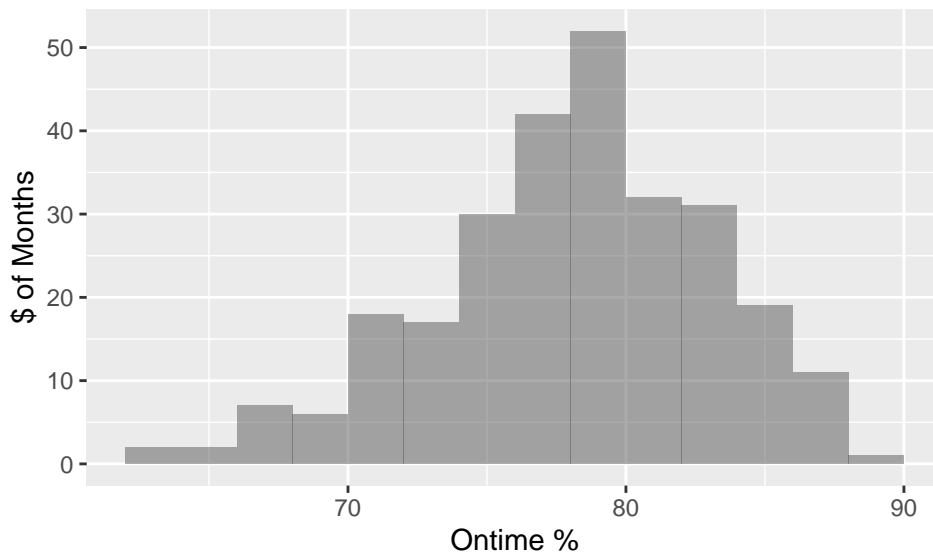
##	response	Container	min	Q1	median	Q3	max	mean	sd	n	missing
## 1	Difference	CUPPS	6	6.00	8.25	12.625	18.5	10.1875	5.202592	8	0
## 2	Difference	Nissan	0	1.25	2.00	3.750	7.0	2.7500	2.507133	8	0
## 3	Difference	SIGG	9	11.75	14.25	21.125	24.5	16.0625	5.900590	8	0
## 4	Difference	Starbucks	6	6.75	8.50	13.625	17.5	10.2500	4.551295	8	0

```
# Mechanics, page 101
```

```
gf_boxplot(Difference ~ Container, data = Cups, ylab = "Temp Change in F")
```

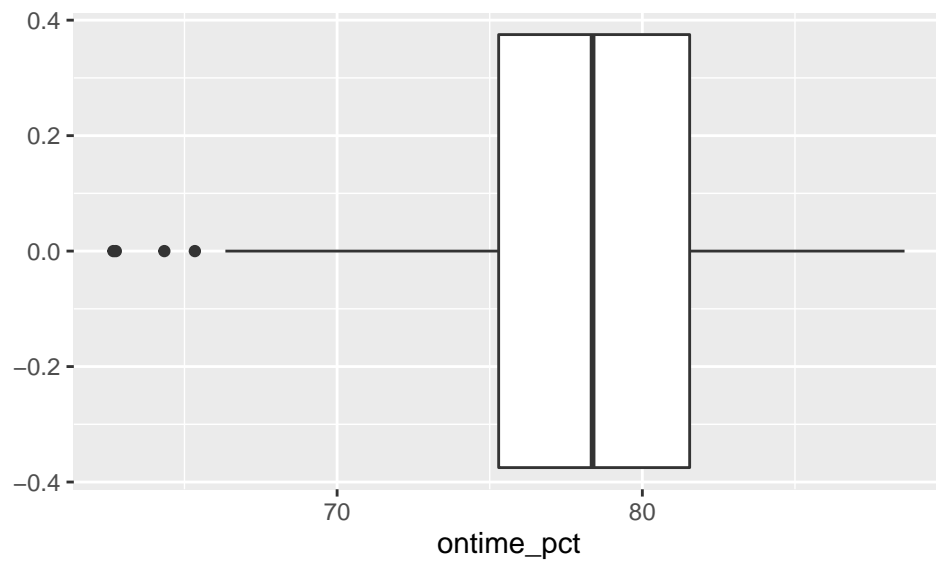


```
Flights <- read_csv("http://nhorton.people.amherst.edu/is5/data/Flights_on_time_2016.csv") %>%
  janitor::clean_names()
# Bureau of Transportation Statistics, page 101
gf_histogram(~ontime_pct, data = Flights, binwidth = 2, center = 1) %>%
  gf_labs(x = "On-time %", y = "$ of Months")
```



Just Checking

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



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

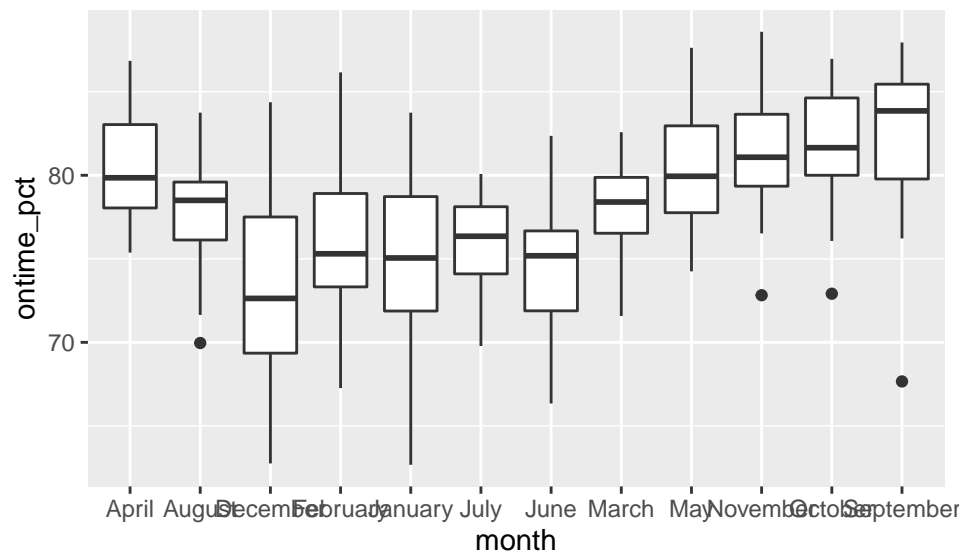


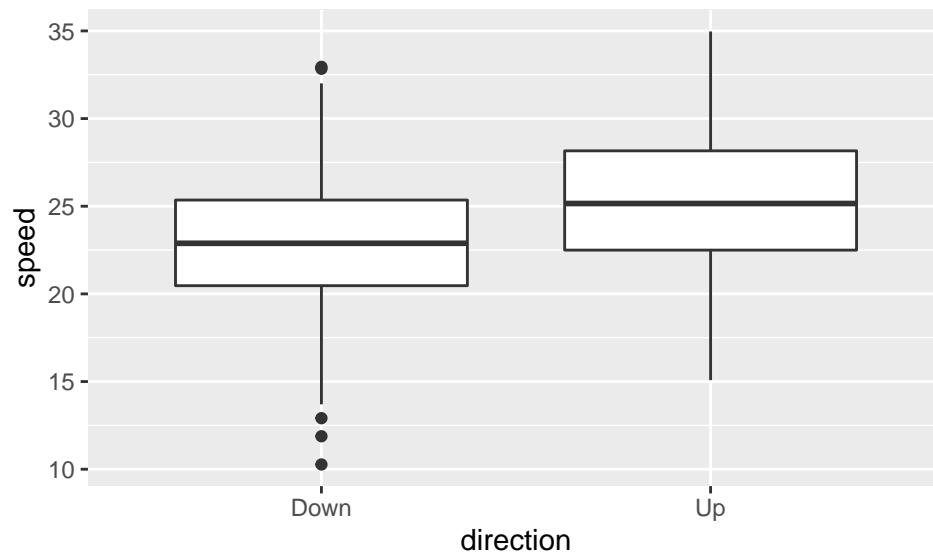
Figure 4.4, page 102

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

Random Matters

```
##
## -- Column specification -----
## cols(
##   direction = col_character(),
##   speed = col_double()
## )
```

```
gf_boxplot(speed ~ direction, data = CarSpeeds)
```

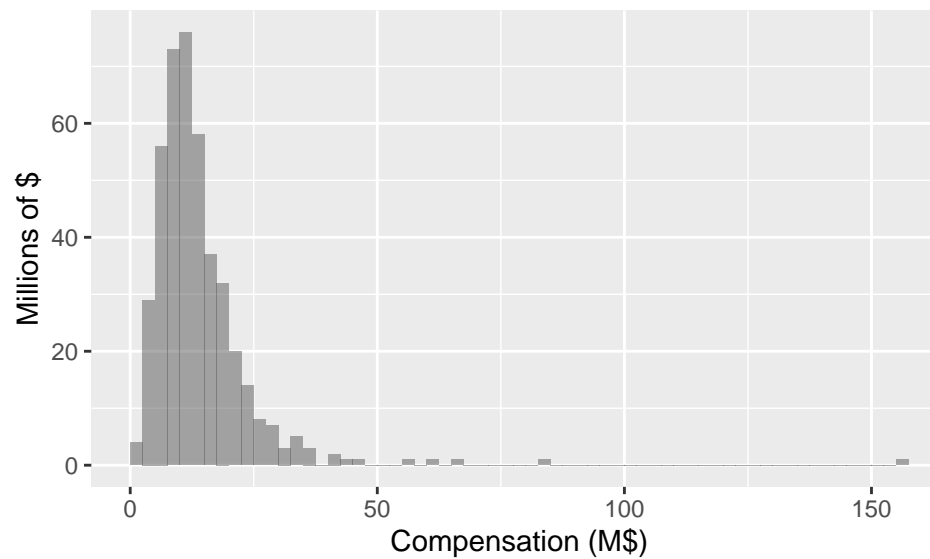


Section 4.3: Re-Expressing Data: A First Look

```
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 $")
```



Re-Expressing to Improve Symmetry

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

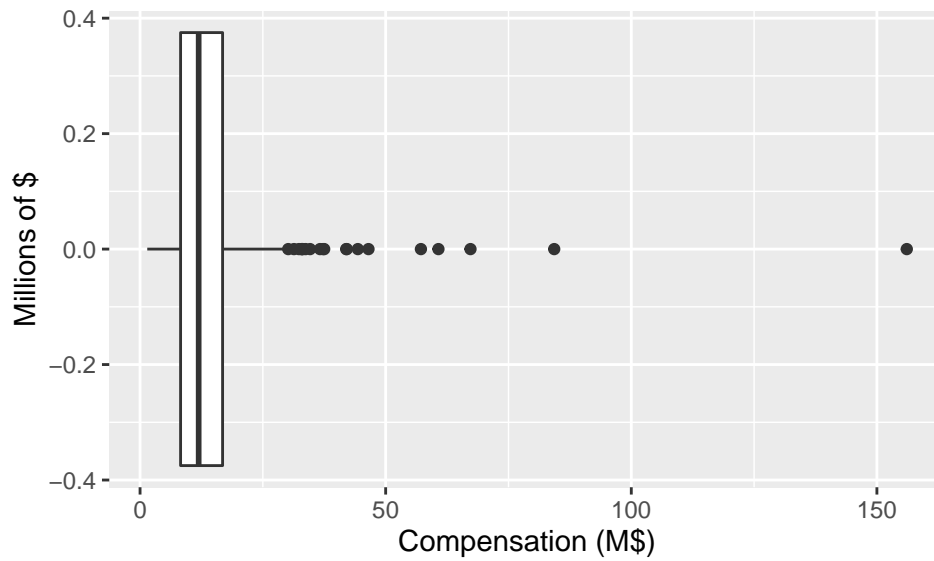
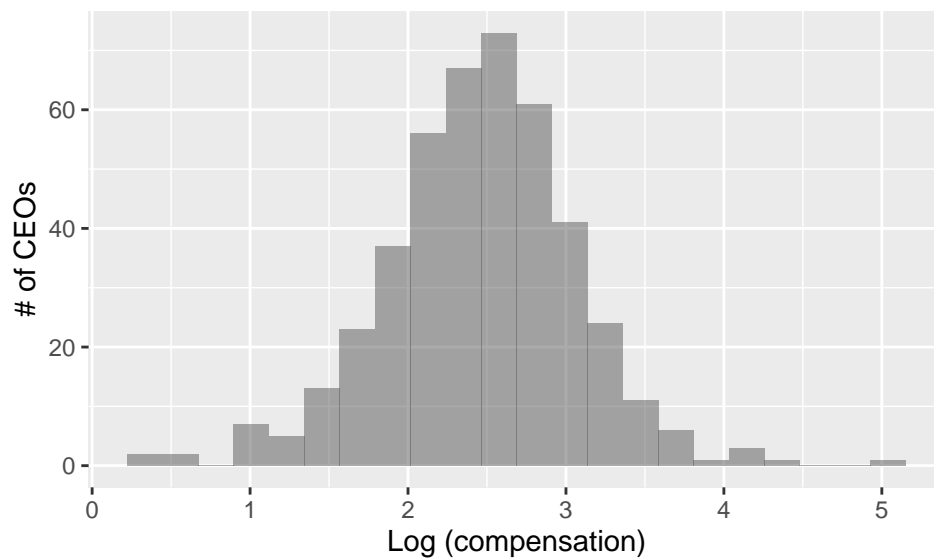



Figure 4.7, page 106

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



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

Re-Expression to Equalize Spread Across Groups

```
##
## -- 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)")
```

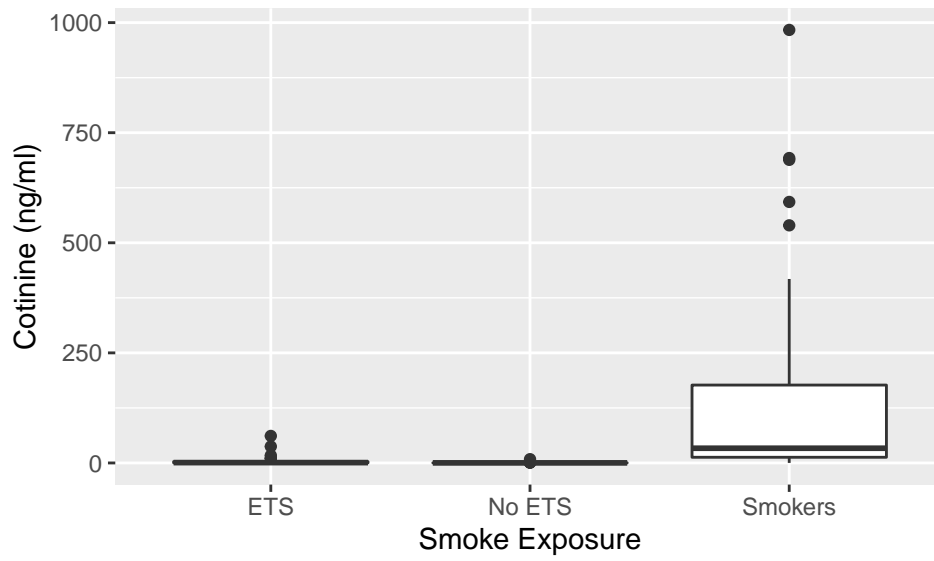


Figure 4.9

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

