

# L07 Scales, Axes & Legends

Data Visualization (STAT 302)

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```
library(knitr)
opts_chunk$set(warning = FALSE, message = FALSE, comment = FALSE, dpi = 300)
```

## Overview

The goal of this lab is to explore ways to manage and manipulate scales, axes, and legends within `ggplot2`.

## Datasets

We'll be using the `tech_stocks.rda`, `corruption.rda`, `cdc.txt`, and a toy dataset.

```
# Load package(s)
library(tidyverse)
library(ggplot2)
library(skimr)
library(scales)
library(lubridate)
library(ggrepel)

# Load datasets
load(file = "data/tech_stocks.rda")
load(file = "data/corruption.rda")

# Read in the cdc dataset
cdc <- read_delim(file = "data/cdc.txt", delim = "|") %>%
  mutate(genhlth = factor(genhlth,
    levels = c("excellent", "very good", "good", "fair", "poor")
  ))

# Set seed
set.seed(7531246)

# Selecting a random subset of size 100
```

```
cdc_small <- cdc %>% slice_sample(n = 100)

# Generating toy dataset for exercise 2
toy_data <- tibble(
  theta = seq(0, 2 * pi, length.out = 100),
  obs = rnorm(100, sin(theta), 0.1),
  larger_than = ifelse(abs(obs) > abs(sin(theta)), "1", "0")
)
```

## Exercises

Complete the following exercises.

### Exercise 1

Using the `tech_stocks` dataset, recreate the following graphic as precisely as possible.

*Hints:*

- Size of lines in legend is 1.3
- 0.75 and 0.85 will be useful
- Eliminated extra space in horizontal direction
- `key_glyph`

```
tech_stocks %>%
  ungroup() %>%

  ggplot(aes(date, price_indexed
    #order by final values in rainbow order
    , color = fct_reorder2(company, date, price_indexed))) +

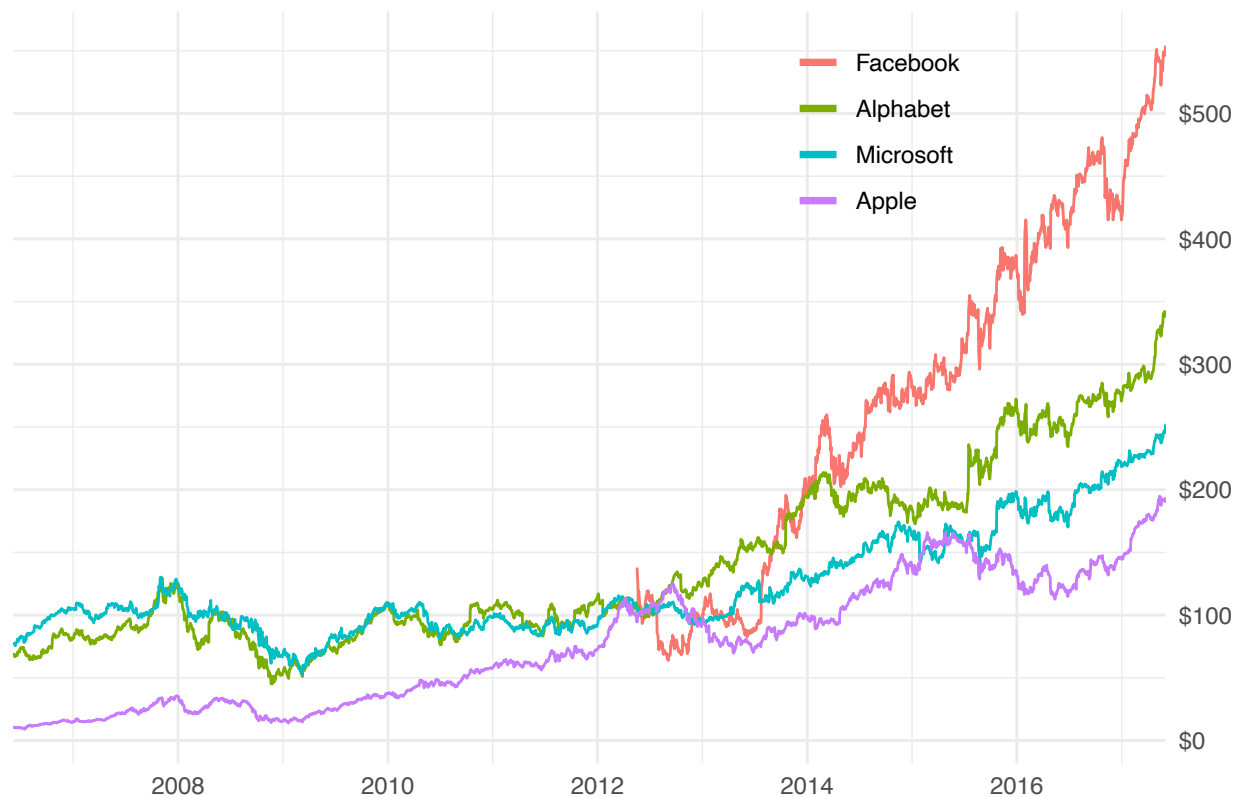
  theme_minimal() +
  geom_line() +
  #legend size is 1.3
  guides(color = guide_legend(override.aes = list(size = 1.3))) +
  scale_y_continuous(
    name = NULL, breaks = seq(0, 500, 100), labels = scales::dollar, position = "right") +

  scale_color_discrete(
    name = NULL) +

  theme(legend.position = c(0.75, 0.85)) +
  scale_x_date(
    name = NULL, expand = c(0, 0)) +

  ggtitle("Stock price, indexed")
```

## Stock price, indexed



### Exercise 2

Using `corruption.rda` dataset, recreate the following graphic as precisely as possible.

Hints:

- Transparency is 0.6
- Only use 2015 data
- Point size is 3 in legend
- Color used is `grey40` & color palette "Set1"
- `"y ~ log(x)"` (method "lm")
- Package `ggrepel`
- `box.padding` is 0.6
- Minimum segment length is 0
- seed is 9876

```
#isolate 2015 data
corruption_2015 <- corruption %>%
  #remove countries missing cpi or hdi values
  drop_na(cpi, hdi) %>%
  #filter to 2015 data
  filter(year == 2015)

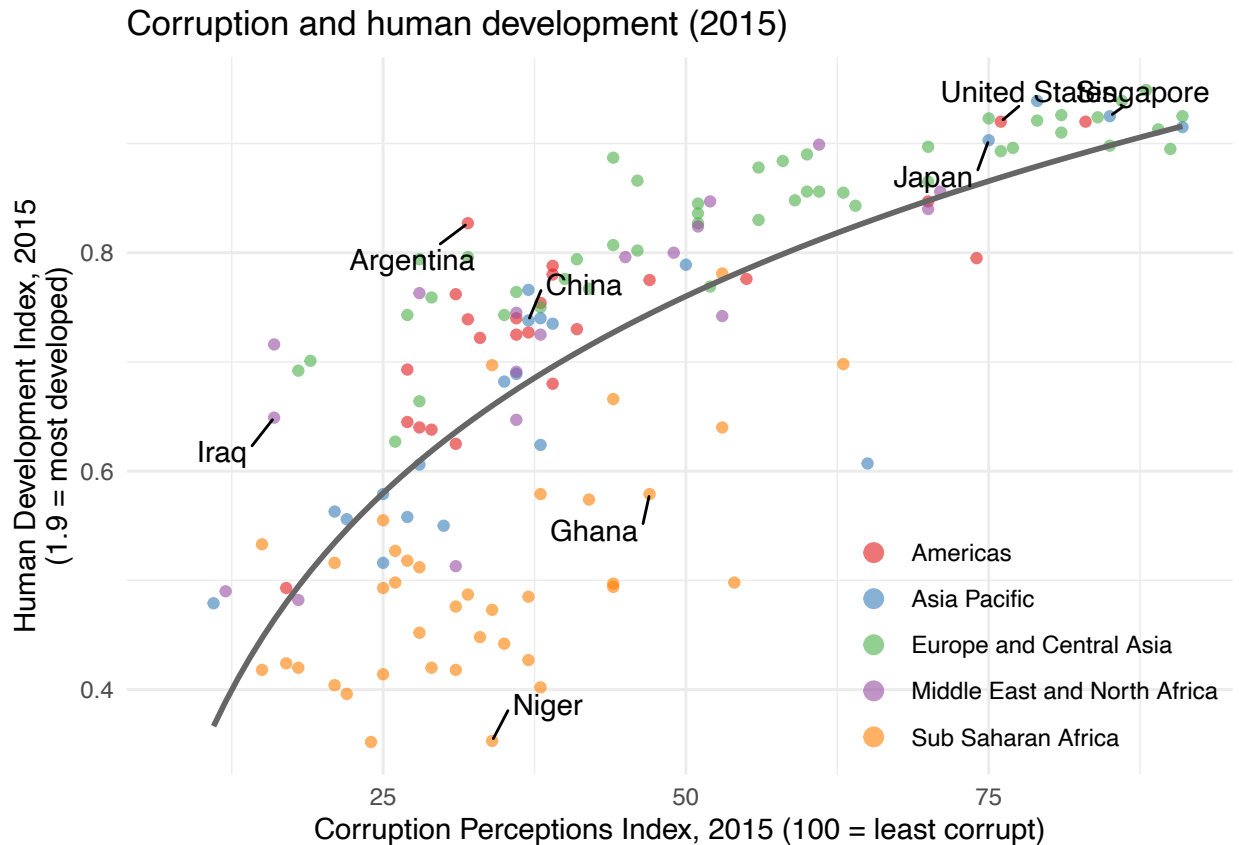
#the countries we want to label
Select_countries <- corruption %>%
  filter(year == 2015) %>%
  filter(country %in% c(
```

```

    "Singapore"
  , "United States"
  , "Japan"
  , "Chlie"
  , "Ghana"
  , "Argentina"
  , "China"
  , "Iraq"
  , "Niger"))

corruption %>%
  filter(year == 2015) %>%
  ggplot(aes(x = cpi, y = hdi)) +
  geom_point(aes(color = region), alpha = .6) +
  geom_smooth(formula = y ~ log(x), method = 'lm', se = F, color = 'grey40') +
  #labels
  labs(x = 'Corruption Perceptions Index, 2015 (100 = least corrupt)',
       y = 'Human Development Index, 2015\n(1.9 = most developed)',
       title = 'Corruption and human development (2015)') +
  theme_minimal() +
  scale_color_brewer(palette = 'Set1') +
  geom_text_repel(data = Select_countries,
                 aes(label = country),
                 box.padding = 0.6,
                 min.segment.length = 0,
                 seed = 9876) +
  theme(legend.position = c(1, 0),
        legend.justification = c("right", "bottom"),
        legend.title = element_blank()) +
  guides(color = guide_legend(override.aes = list(size = 3)))

```



### Exercise 3

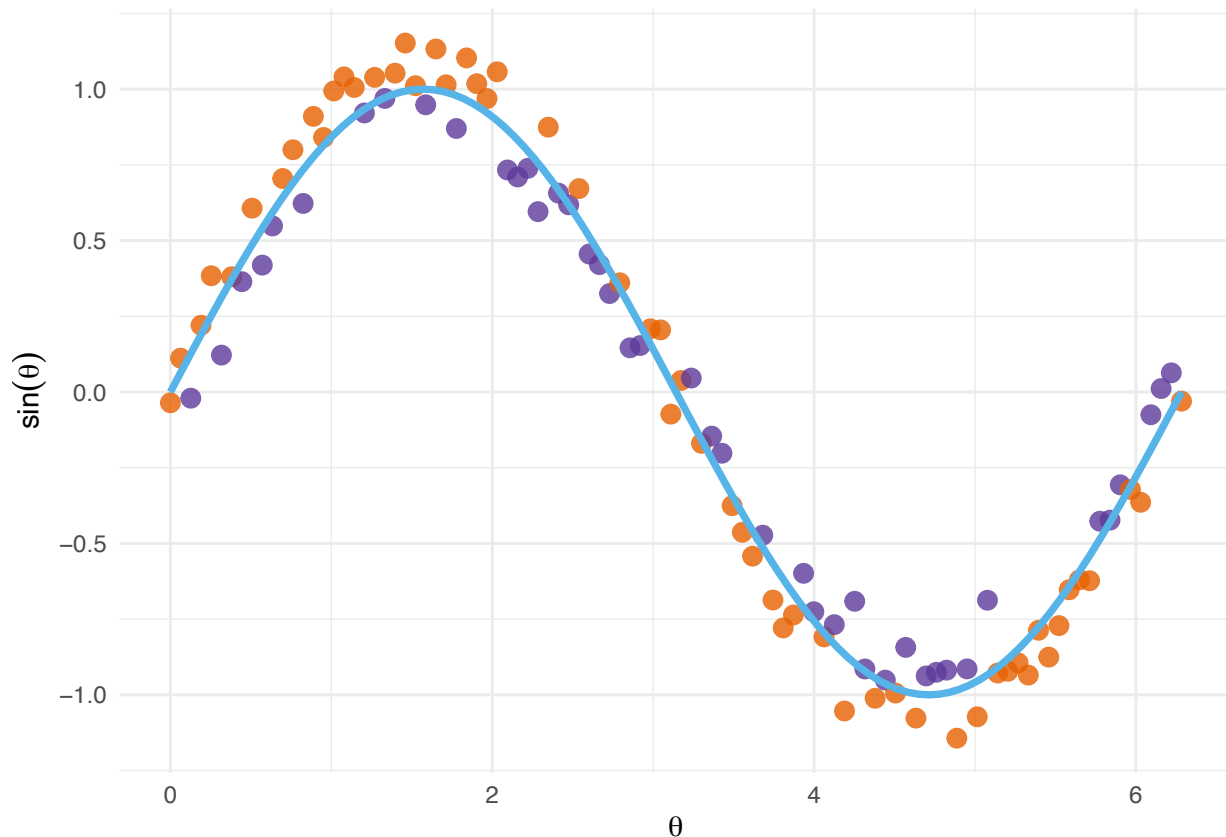
Use `toy_data` to recreate the following graphic as precisely as possible.

*Hints:*

- Sizes used 3 (points) and 1.3 (line)
- Transparency 0.8
- Colors: #56B4E9, #E66100, #5D3A9B
- `stat_function()` will be useful

```
toy_data %>%
  ggplot(aes(x = theta, y = obs, color = larger_than)) +
  geom_point(size = 3, alpha = 0.8) + #given size and transparency
  geom_function(fun = function(x) sin(x), color = '#56B4E9', size = 1.3) +
  labs(x = quote(theta), y = quote(sin(theta))) + #label axes

theme_minimal() +
  theme(legend.position = 'none') +
  #set given colors
  scale_color_manual(values = c('0' = '#5D3A9B', '1' = '#E66100'))
```



#### Exercise 4

Using `cdc_small` construct a scatterplot of `weight` by `height` with the following requirements:

- Size of plotting characters should be 3.
- Color and shape should both identify `genhlth`.
- One legend for both color and shape.
- Legend title should be “General Health?” with a newline starting after general.
- Legend categories should be ordered from excellent (top) to poor (bottom) with each word in category capitalized in the legend.
- Legend should be placed in the lower right-hand corner of the plotting area.
- Color should follow the “Set1” palette.
- Shape should have a solid triangle (17) for excellent, solid circle (19) for very good, an x (4) for poor, an hollow rotated square with an x in it (9) for fair, and a solid square (15) for good.
- `height` values should be limited between 55 and 80.
- `height` axis should display every 5th number between 55 and 80 and be appropriately labeled (i.e. 55 in, 60 in, ..., 80 in). No axis title is necessary.
- `weight` values should be limited between 100 and 300.
- `weight` axis should be on log base 10 scale, but still display weights in pounds starting at 100 and displaying every 25 pounds until 300. Must be appropriately labeled (i.e. 100 lbs, 125 lbs, ..., 300 lbs). No axis title is necessary.
- Graph title should be CDC BRFSS: Weight by Height.
- Minimal theme.

```
cdc_small %>%
  ggplot(aes(x = height, y = weight)) + #plot height by weight
  geom_point(aes(color = genhlth, shape = genhlth), size = 3) +
  theme_minimal() +
```

```

scale_shape_manual(values = c(17, 19, 15, 9, 4),
  labels = c('Excellent', 'Very Good', 'Good', 'Fair', 'Poor'),
  name = 'General\nHealth?') +

theme(#legend.title = 'General\nHealth?',
  legend.position = c(1, 0),
  legend.justification = c(1, 0)) +

scale_color_brewer(palette = 'Set1',
  name = 'General\nHealth?',
  labels = c('Excellent', 'Very Good', 'Good', 'Fair', 'Poor')) +

labs(x = NULL, y = NULL, title = 'CDC BRFSS: Weight by Height') +
scale_y_log10(breaks = c(100, 125, 150, 175, 200, 225, 250, 275, 300),
  limits = c(100, 300),
  labels = c('100 lbs', '125 lbs', '150 lbs', '175 lbs', '200 lbs', '225 lbs', '250 lbs', '275 lbs', '300 lbs'),

scale_x_continuous(breaks = c(55, 60, 65, 70, 75, 80),
  limits = c(55, 80),
  labels = c('55 in', '60 in', '65 in', '70 in', '75 in', '80 in'))

```

CDC BRFSS: Weight by Height

