

Lab 02: Data Visualization

To run this lab yourself you should create a blank quarto document and copy and paste the included code over, to then complete the required tasks.

HIV prevalence from WHO

- Estimated HIV prevalence was obtained from the `gapminder` website <https://www.gapminder.org/data/>
 - Estimated number of people living with HIV per 100 population of age group 15-49.
 - Original data source is the UNAIDS online database at <http://www.aidsinfoonline.org>
- A spreadsheet of the data, `HIVprev.csv`, is necessary for this lab.

We can read in these data as follows (we'll learn about reading in data later in STAT 260):

```
library(tidyverse)

# you must have already installed the tidyverse package

hiv <- read.csv("HIVprev.csv", stringsAsFactors = FALSE)
hiv <- select(hiv, Country, year, prevalence)
```

Take a look at the top and bottom few lines of raw data.

```
head(hiv)
```

	Country	year	prevalence
1	Algeria	1990	0.06
2	Algeria	1991	0.06
3	Algeria	1992	0.06
4	Algeria	1993	0.06
5	Algeria	1994	0.06
6	Algeria	1995	0.06

```
tail(hiv)
```

	Country	year	prevalence
1601	Zimbabwe	1995	25.1
1602	Zimbabwe	1996	26.2
1603	Zimbabwe	1997	26.5
1604	Zimbabwe	1998	26.3
1605	Zimbabwe	1999	25.7
1606	Zimbabwe	2000	24.8

```
summary(hiv)
```

	Country	year	prevalence
Length:	1606	Min. :1990	Min. : 0.060
Class :	character	1st Qu.:1992	1st Qu.: 0.060
Mode :	character	Median :1995	Median : 0.200
		Mean :1995	Mean : 1.575
		3rd Qu.:1998	3rd Qu.: 1.100
		Max. :2000	Max. :26.500

Exercises:

1. Plot the time series of HIV prevalence by year for each country using `geom_line()`.
2. Redo the above plot but experiment with different `alpha` values. What problem does setting a small `alpha` overcome? What feature of the graph is hidden when we do not set `alpha`?
3. In the following code chunk we create a new dataset comprised of countries that had HIV prevalence greater than 10% in one or more of the years monitored (we will learn about this kind of “data wrangling” in future lectures of STAT 260).

```
cc <- c(
  "Botswana", "Central African Republic", "Congo", "Kenya", "Lesotho", "Malawi",
  "Namibia", "South Africa", "Swaziland", "Uganda", "Zambia", "Zimbabwe"
)
hihiv <- filter(hiv, Country %in% cc)
```

Add red lines for the above countries to your time series plot from Exercise 2.

4. Redo the time series plot from Exercise 1, with the following modifications. Color the time series for all but the countries in the `hihiv` data frame (i.e., those with high HIV prevalence) **grey** and with `alpha=0.3`. For the high-HIV-prevalence countries, color them **red**, also using `alpha=0.3`. Next, add two smoothers: (i) for all the data, i.e. all the countries in the `hiv` data frame, colored **black**, and (ii) for the countries with a high prevalence of HIV, i.e. those in the `hihiv` data frame, colored **red**. Your final plot should look like this (do not worry about axis labels or title):

