Data Visualization

Fall R'23

Dominic Bordelon, Research Data Librarian, ULS



Agenda

- 1. The grammar of graphics
- 2. Plotting distribution
- 3. Scatter plots
- 4. Comparing categories
- 5. Heat maps of two-way tables
- 6. Time series
- 7. Small multiples
- 8. Labeling, theming, and annotation

About the trainer

Dominic Bordelon, Research Data Librarian
University Library System, University of Pittsburgh
dbordelon@pitt.edu

Services for the Pitt community:

- Consultations
- Training (on-request and via public workshops)
- Talks (on-request and publicly)
- Research collaboration

Support areas and interests:

- Computer programming fundamentals, esp. for data processing and analysis
- Open Science and Data Sharing
- Data stewardship/curation
- Research methods; science and technology studies

Fall R Series

#	Date	Title
1	8/29	Getting Started with Tabular Data
2	9/5	Working with Data Frames
3	9/12	Data Visualization
	0,	Data Visaatization
4		Inference and Modeling Intro

R and RStudio Drop-In Hour

Bring your R questions!

If we don't know the answer,
we'll help you look for it.

Students, post-docs, faculty, and staff are all welcome.

For R users in the Pitt community

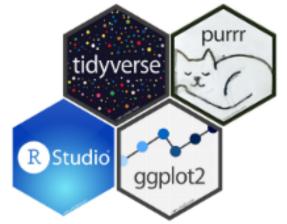
Tuesdays, 4:30–5:30 pm Fall semester 2023

Hillman Library, Rm. 255

https://bit.ly/pitt-r-office-23

Service provided by: University Library System, Digital Scholarship Services













The ggplot2 package

"ggplot2 is a system for declaratively creating graphics, based on The Grammar of Graphics. You provide the data, tell ggplot2 how to map variables to aesthetics, what graphical primitives to use, and it takes care of the details."



...and using penguins examples

```
1 install.packages("palmerpenguins")

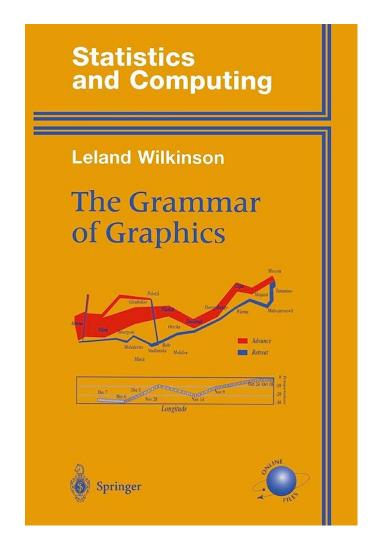
1 library(palmerpenguins)
2
3 # load palmerpenguins' data into your environment:
4 data(penguins)
5 names(penguins)

[1] "species" "island" "bill_length_mm"
[4] "bill_depth_mm" "flipper_length_mm" "body_mass_g"
[7] "sex" "year"
```

palmerpenguins is one of many examples of an R package which functions as a downloadable data set.

The grammar of graphics

- A plot is constructed in layers:
 - data
 - aesthetics (axes, encodings)
 - scale (axis labels, color coding)
 - geometric objects (bar, scatter, heatmap tiles, etc.)
 - facets
 - statistical summaries (e.g., highlighted mean; smoother)
 - annotations
 - coordinate system (Cartesian, polar, or map projection)
 - theme



Wilkinson 1999

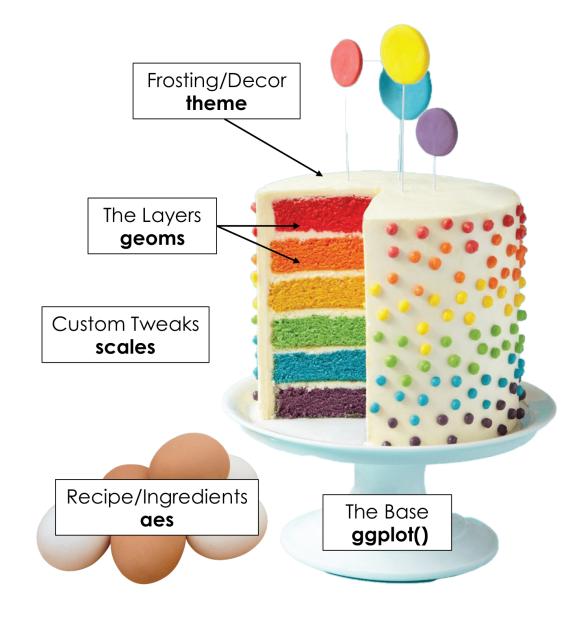
ggplot is a little bit like cake...

We always start by setting up the foundation with **ggplot()**

We specify our ingredients (data variables) with an **aes mapping**

We can create *layers* to our plot with **geoms**

We can style our cake ggplot with **themes.** We have out-of-the-box options, or we can go totally custom!



Source: Tanya Shapiro, https://twitter.com/tanya_shapiro/status/1576935152575340544

Aesthetic mapping

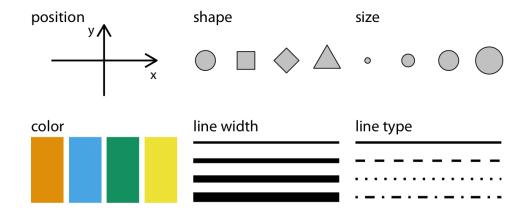
An aesthetic mapping, created with aes(), associates an aesthetic property of the plot with a variable in your data.

penguins example: I want a scatter plot of flipper length on x and bill length on y. The aesthetic mapping would be:

```
aes(x = flipper_length_mm, y =
bill length mm)
```

Common aesthetics:

- x, y
- color, fill
- linetype
- size
- shape (of points)



Plotting distribution

Histograms and density plots

geom_histogram(mapping, binwidth) and geom_density(mapping)

A histogram "bins" a variable's values and charts how many values are in each bin, giving a sense of central tendency and spread. A density plot works similarly, but it produces a smooth curve while sacrificing countable units.

```
1 # distribution of body mass:
 2 penguins %>%
     gaplot() +
     geom\ histogram\ (mapping = aes\ (x = body\ mass\ g))
   # remove NA values, and specify bin width:
   penguins %>%
     drop na(body mass g) %>%
     ggplot() +
 9
     geom histogram (mapping = aes (x = body mass g),
10
11
                     binwidth = 100)
12
   # density plot:
   penguins %>%
15
     gaplot() +
     geom density (aes (x = body mass g))
16
17
18 # layering histogram and density plot
   # note that 1) mapping is specified in ggplot(), and inherited by geoms; 2) histogram's y mapping needs adjus
   penguins %>%
     drop na(body mass g) %>%
     ggplot(aes(x = body mass g)) +
                                                                                                        Pittsburgh Library System
                                                R 3: Data Visualization
```

Comparing distributions: violin, box

geom_violin(), geom_boxplot(). Mapping a categorical variable onto an axis will
make a violin or box plot for each level of the variable.

A violin plot mirrors a density plot across the axis. A box plot marks interquartile range (white box), median (line inside the box), tails, and outliers (individual points). (where outlier has a distance \(\\gt 1.5 \times \mathrm{IQR}\) from the median)

Scatter plots

geom_point() plots points

A scatter plot compares continuous variables (x) and (y) on a Cartesian plane. geom_point() can take several aesthetics (size, shape, color), but be careful not to overload your plot with information.

geom_smooth() adds a smoother to a scatter plot

A natural next step to a scatter plot is to fit a regression model. <code>geom_smooth()</code> does this job; specify the modeling function you want with the <code>method</code> argument. The default methods are <code>loess</code>, local polynomial regression fitting (when \(n \t 1000\)), or <code>gam</code>, generalized additive model with restricted maximum likelihood (when \(n \geq 1000\)). See <code>?stats::loess</code> and <code>?mgcv::gam</code> for more info on these modeling functions.

(If you have fit your own model outside of ggplot, you should instead use predict() to generate points to plot with geom_line().)

Comparing categories

geom_bar() makes bar charts of counts

```
penguins %>%
    ggplot(aes(x = species)) +
    geom_bar()

# note 2 variables, y axis, and dodge position
# we also reorder the factors for aesthetic reasons
# ...and note unexpected colors!
penguins %>%

mutate(species = fct_rev(fct_infreq(species)),
    sex = fct_infreq(sex)) %>%

ggplot(aes(y = species, fill = sex)) +
geom_bar(position = position_dodge2(reverse=TRUE, preserve="single"))
```

Bar charts of calculated statistics

Perform summary calculations using dplyr, then use geom_bar() with stat = "identity" to plot the numbers as-is (default is "count"). \Delta When plotting a summary statistic, one should also include error! geom_errorbar(), geom_linerange()

Example: mean body mass for each species and gender, with standard error

```
penguins %>%
     drop na(body mass g) %>%
     group by(species) %>%
     summarize (mean mass g = mean (body mass g),
                standard error = sd(body mass g)/sqrt(n())
 6
     ggplot(aes(x = species, y = mean mass g)) +
     geom bar(stat = "identity") +
     geom errorbar(aes(ymax = mean mass g + standard error,
                        ymin = mean mass g - standard error), width = 0.5, color="blue")
11
   # note 2 variables, y axis, and dodge position
     we also reorder the factors for aesthetic reasons
   # ...and note unexpected colors!
   penguins %>%
     drop na(body mass g) %>%
16
     group by (species, sex) %>%
17
18
     summarize (mean mass g = mean (body mass g),
                standard error = sd(body mass g)/sqrt(n())
19
                ) 응>응
     ggplot(aes(y = species, fill = sex, x = mean_pmassig)alization
                                                                                                       Pittsburgh | Library System
22
     geom bar(stat="identity",
```

Stacked proportional bars and pies

Sometimes we want to compare *proportions* of category membership. With **position** = "fill", each bar has the same height, and the fill aesthetic works proportionately.

```
penguins %>%
ggplot(aes(x = species, fill=sex)) +
geom_bar(position = "fill") +
scale_y_continuous(labels = scales::percent)
```

A pie chart in ggplot2 terms is a proportional bar cast onto a polar coordinate system.

Heat maps of two-way tables

Two-way tables

The two-way table or contingency table format shows the conditional distribution of observations among two categorical variables. In penguin terms, how many penguins of each species were observed on each of three islands?

```
1 penguins %>%
2  janitor::tabyl(species, island)

species Biscoe Dream Torgersen
Adelie 44 56 52
Chinstrap 0 68 0
Gentoo 124 0 0
```

Heat maps of two-way tables

A graphical version of the previous table is given by geom_tile(). However, you will need to group and summarize the data first.

```
penguins %>%
group_by(species, island) %>%
summarize(n = n()) %>%
ggplot() +
geom_tile(aes(y=species, x=island, fill=n))
```

Time series

geom_line() connects point observations

Time series data are the most common application for line graphs. Since line graphs expect a single (y) for each (x), I recommend using dplyr to generate the table that you need. The examples use n() as the summarizing function, but one could also use sum() of a variable (for example).

```
# observations over time
2 penguins %>%
   group by (year) %>%
   summarize(n = n()) %>%
   ggplot() +
     geom line(aes(x=year, y=n))
   # a more interesting example, using tropical storm/hurricane data
   ?storms
10 storms %>%
     group by (year) %>%
12
     summarize (n = n()) \%
13
     ggplot() +
     geom line(aes(x=year, y=n)) +
14
     labs(v = "Storm observations")
15
```

Categories over time

Consider geom_line() for comparison, or geom_area() for showing cumulative distribution.

```
1 # observations over time
 2 penguins %>%
     group by (year, species) %>%
     summarize (n = n()) \%
     gaplot() +
     geom line(aes(x=year, y=n, linetype=species))
8 # storms by hurricane category:
   # note: ~5-7 is the greatest number of categories you can color code before readers start having difficulty
10 storms %>%
     mutate(category = as factor(category)) %>%
     drop na(category) %>%
     group by (year, category) %>%
14
     summarize (n = n()) \%>\%
15
     ggplot() +
     geom line(aes(x=year, y=n, color=category)) +
16
17
     labs(y = "Storm observations")
18
19 storms %>%
     mutate(category = as factor(category)) %>%
     drop na(category) %>%
     group by (year, category) %>%
23
     summarize (n = n()) \%
24
     gaplot() +
25
     geom area(aes(x=year, y=n, fill=category)) +
26
     labs(y = "Storm observations")
```

Small multiples

Small multiples

Small multiples AKA faceting replicate a plot in columns and/or rows, with one plot for each level of some categorical variable.

Want to combine multiple plots into one figure? Check out the patchwork package.

```
1 # one line graph for each storm category
 2 storms %>%
     mutate(category = as factor(category)) %>%
     drop na(category) %>%
     group by (year, category) %>%
     summarize (n = n()) \%
     ggplot() +
     geom line(aes(x=year, y=n, color=category)) +
 8
     facet wrap(vars(category), ncol=3, nrow=2) +
     labs(y = "Storm observations")
10
11
   # a scatter for each island, with groups colored
   penguins %>%
14
     ggplot(aes(x=flipper length mm,
15
                y=bill length mm,
                color=species)) +
16
     geom point() +
17
18
     facet wrap(vars(island), nrow=2, ncol=2)
```

Labeling, theming, and annotation

labs() sets labels

```
penguins %>%
     drop na(flipper length mm, bill length mm) %>%
     ggplot(aes(x = flipper length mm,
                y = bill length mm,
                color = species)) +
 5
     geom point() +
 6
     geom smooth(method="lm") +
     labs(title = "Gentoo tend to have the longest flippers",
          x = "Flipper length (mm)",
9
     y = "Bill length (mm)",
10
     color = "Species",
11
12
          caption = "Source: Gorman KB, Williams TD, Fraser WR (2014)")
```

theme_ layers change the plot's look

Start typing theme_ in RStudio to see available options, or install and attach the ggthemes package for more. There is also a theme() function for adjusting specific aspects of the plot.

Lastly, annotate() can add annotations to the plot, using the coordinate system.

```
penguins %>%
     drop na (body mass g) %>%
     gaplot() +
     geom histogram (mapping = aes(x = body mass g),
                    binwidth = 100) +
     annotate (geom="text", x = 6200, y = 7,
 6
7
              label="Biggest\n penguin") +
     annotate (geom="segment", x = 6200, y = 5,
8
9
              xend = max(penguins$body mass g, na.rm=TRUE),
              yend = 1.5) +
10
     theme bw()
11
```

We can't forget to save!

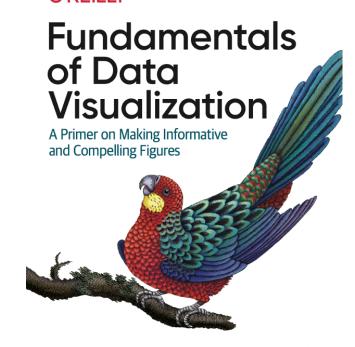
ggsave(filename, plot) where plot is a ggplot() object you have assigned..png,
.svg, or .pdf in the filename determines the output type.

Other arguments to use: width, height, units

Visualize responsibly

- Consider the validity and reliability of your measures
- Always cite any data you visualize which you did not collect yourself
- Design for accessibility
- Study best practices, and some of the pitfalls/abuses of viz, to keep your plot honest

O'REILLY®



Claus O. Wilke

Wilke 2019, available free-to-read at https://clauswilke.com/dataviz/



Wrap up

Session in review

Today we learned about:

- the grammar of graphics and aesthetic mapping
- lots of plot types! for a variety of applications
- some easy ways of customizing plots

Join us next week for inference and modeling!