Reporting data results #1

Plot guidelines

Guidelines for good plots

There are a number of very thoughtful books and articles about creating graphics that effectively communicate information.

Some of the authors I highly recommend (and from whose work I've pulled the guidelines for good graphics we'll talk about this week) are:

- Edward Tufte
- Howard Wainer
- Stephen Few
- Nathan Yau

You should plan, in particular, to read *The Visual Display of Quantitative Information* by Edward Tufte before you graduate.

Guidelines for good plots

This week, we'll focus on six guidelines for good graphics, based on the writings of these and other specialists in data display.

The guidelines are:

- 1. Aim for high data density.
- 2. Use clear, meaningful labels.
- 3. Provide useful references.
- 4. Highlight interesting aspects of the data.
- 5. Make order meaningful.
- 6. When possible, use small multiples.

Packages for examples

For the examples, I'll use dplyr for data cleaning and, for plotting, the packages ggplot2, gridExtra, and ggthemes.

```
library("dplyr")
library("ggplot2")
library("gridExtra")
library("ggthemes")
```

Example data

You can load the data for today's examples with the following code:

```
library("faraway")
data("worldcup")
```

Example data

```
library("dlnm")
data("chicagoNMMAPS")
chic <- chicagoNMMAPS</pre>
chic july <- chic %>%
  filter(month == 7 & year == 1995)
slice(chic_july, 1:3)
## date time year month doy dow death cvd resp temp dptp
## 1 1995-07-01 3104 1995 7 182 Saturday 109 52 4 18.61111 47.750
## 2 1995-07-02 3105 1995 7 183 Sunday 103 49 4 17.50000 49.375
## 3 1995-07-03 3106 1995 7 184 Monday 112 51 10 21.66667 57.500
## rhum pm10 o3
## 1 55,250 15,64046 27,47199
## 2 47.875 27.12486 24.81275
## 3 51,000 49,11362 43,63302
```

High data density

Guideline 1: Aim for high data density.

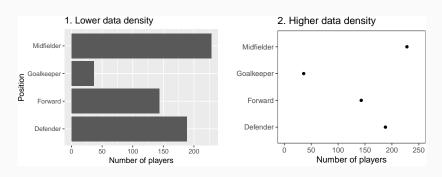
You should try to increase, as much as possible, the **data to ink ratio** in your graphs. This is the ratio of "ink" providing information to all ink used in the figure.

One way to think about this is that the only graphs you make that use up a lot of your printer's ink should be packed with information.

High data density

Guideline 1: Aim for high data density.

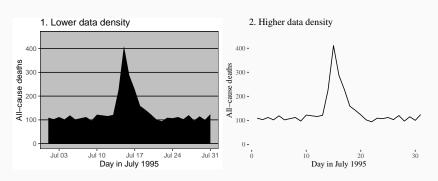
The two graphs below show the same information. Compare the amount of ink used in the left plot to the amount used in the right plot to see how graphs with the same information can have very different data densities.



High data density

Guideline 1: Aim for high data density.

The two graphs below show another example of very different data densities in two plots showing the same information:



Data density

One quick way to increase data density in ggplot2 is to change the *theme* for the plot. This essentially changes the "background" elements to a plot, including elements like the plot grid, background color, and the font used for labeling.

Some themes come with ggplot2, including:

- theme_classic
- theme_bw
- theme_minimal
- theme_void

The ggthemes packages has some excellent additional themes.

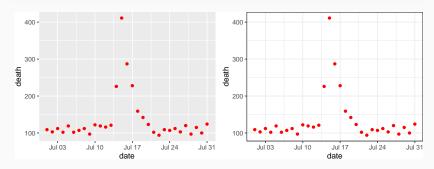
Data density

The following slides show some examples of the effects of using different themes. The following code creates a plot of daily deaths in Chicago in July 1995:

Next, we can see how the graph looks with the default theme and with other themes.

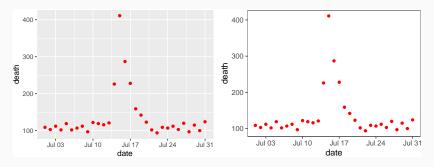
The left graph shows the graph with the default theme, while the right shows the effect of adding the black-and-white theme that comes with ggplot2 as theme_bw:

```
a <- chic_plot
b <- chic_plot + theme_bw()
grid.arrange(a, b, ncol = 2)</pre>
```



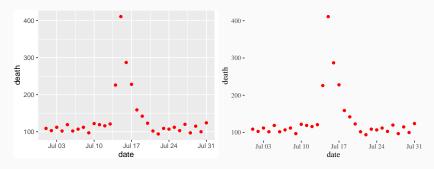
Stephen Few theme:

```
a <- chic_plot
b <- chic_plot + theme_few()
grid.arrange(a, b, ncol = 2)</pre>
```



Edward Tufte theme:

```
a <- chic_plot
b <- chic_plot + theme_tufte()
grid.arrange(a, b, ncol = 2)</pre>
```



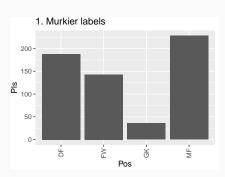
You can even use themes to add some questionable choices for different elements. For example, ggthemes includes an Excel theme:

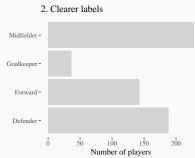
```
a <- chic plot
b <- chic plot + theme excel()
grid.arrange(a, b, ncol = 2)
  400 -
                                                   400
  300 -
                                                   300
leath
                                                death
  200 -
                                                   200
                         Jul 17
                                  Jul 24
       Jul 03
                Jul 10
                                           Jul 31
                                                        Jul 03
                                                                 Jul 10
                                                                          Jul 17
                                                                                   Jul 24
                                                                                           Jul 31
                        date
                                                                         date
```

Meaningful labels

Guideline 2: Use clear, meaningful labels.

Graph defaults often use abbreviations for axis labels and other labeling. Further, text labels can sometimes be aligned in a way that makes them hard to read. The plots below give an example of the same information shown without (left) and with (right) clear, meaningful labels.





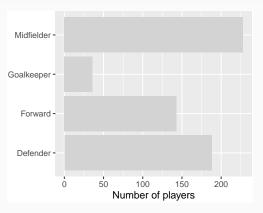
Meaningful labels

There are a few strategies you can use to make labels clearer:

- Add a labs() element to the plot, rather than relying on the column names in the original data. This can also be done with scale elements (e.g., scale_x_continuous), which give you more power to also make other changes to these scales.
- Include units of measurement in axis titles when relevant. If units are dollars or percent, check out the scales package, which allows you to add labels directly to axis elements by including arguments like labels = percent in scale elements.
- If the x-variable requires longer labels (player positions in the example above), consider flipping the coordinates, rather than abbreviating or rotating the labels. You can use coord_flip to do this.

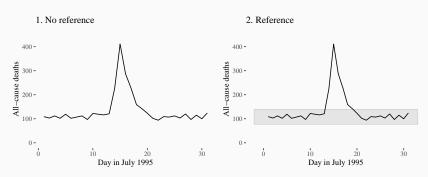
Meaningful labels

```
ggplot(worldcup, aes(Position)) +
    geom_bar(fill = "lightgray") + coord_flip() +
    labs(x = "", y = "Number of players")
```



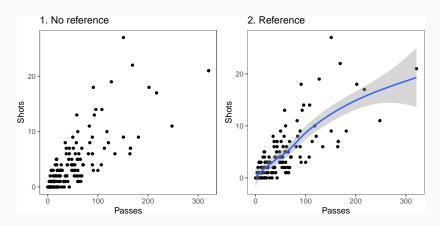
Guideline 3: Provide useful references.

Data is easier to interpret when you add references. For example, if you show what it typical, it helps viewers interpret how unusual outliers are. The graph below on the right has added shading showing the range of daily deaths in July in Chicago for 1990–1994 and 1996–2000, to clarify how unusual July 1995 was.



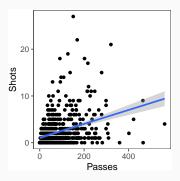
Guideline 3: Provide useful references.

Another useful way to add references is to add a linear or smooth fit to the data, to help clarify trends in the data.



You can use the function geom_smooth to add a smooth or linear reference line:

```
ggplot(worldcup, aes(x = Passes, y = Shots)) +
geom_point() + theme_few() +
geom_smooth(method = "lm")
```

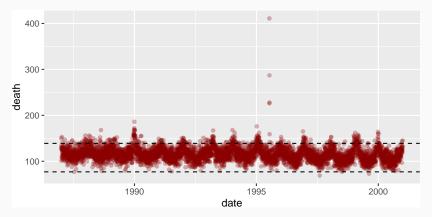


The most useful geom_smooth parameters to know are:

- method: The default is to add a loess curve if the data includes less than 1000 points and a generalized additive model for 1000 points or more. However, you can change to show the fitted line from a linear model using method = "lm" or from a generalized linear model using method = "glm".
- span: How wiggly or smooth the smooth line should be (smaller value: more wiggly; larger value: more smooth)
- se: TRUE or FALSE, indicating whether to include shading for 95% confidence intervals.
- level: Confidence level for confidence interval (e.g., 0.90 for 90% confidence intervals)

Lines and polygons can also be useful for adding references. Useful geoms include:

- geom hline, geom vline: Add a horizontal or vertical line
- geom_abline: Add a line with an intercept and slope
- geom_polygon: Add a filled polygon
- geom_path: Add an unfilled polygon



When adding these references:

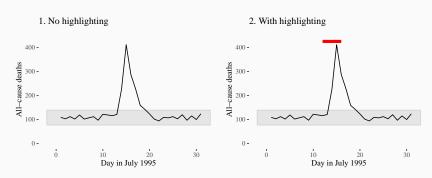
- If you want them to go behind the main points, add reference elements first, so they will be plotted under the data, instead of on top of it.
- Use alpha to add transparency to these elements.
- Use colors that are unobtrusive (e.g., grays)
- For lines, consider using non-solid line types (e.g., linetype = 3)

In-course exercise

We'll take a break now to do Sections 1 and 2 of the In-Course Exercise (Chapter 4).

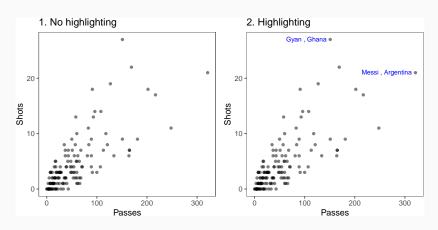
Guideline 3: **Highlight interesting aspects.**

Consider adding elements to highlight noteworthy elements of the data. For example, in the graph on the right, the days of a major heat wave have been highlighted with a red line.



Guideline 3: **Highlight interesting aspects.**

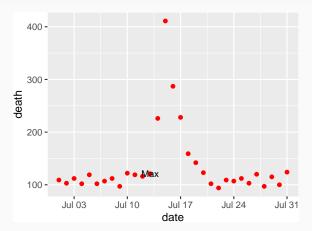
In the below graphs, the names of the players with the most shots and passes have been added to highlight these unusual points.



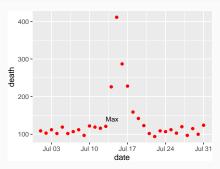
One helpful way to annotate is with text, using geom_text(). For this, you'll first need to create a dataframe with the hottest day in the data:

```
hottest_day <- chic_july %>%
filter(temp == max(temp))
hottest_day %>% select(date:dow)
```

```
## date time year month doy dow
## 1 1995-07-13 3116 1995 7 194 Thursday
```



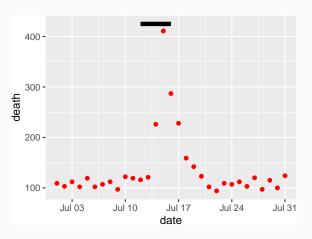
With geom_text, you'll often want to use position adjustment (the position parameter) to move the text so it won't be right on top of the data points:



You can also use lines to highlight. For this, it is often useful to create a new dataframe with data for the reference. To add a line for the Chicago heat wave, I've added a dataframe called hw with the relevant date range. I'm setting the y-value to be high enough (425) to ensure the line will be placed above the mortality data.

```
library(lubridate)
hw <- data.frame(date = c(ymd("1995-07-12")),
                          ymd("1995-07-16")),
                 death = c(425, 425))
b <- ggplot(data = chic_july, aes(x = date, y = death)) +
        geom point(color = "red") +
        geom_line(data = hw,
                  aes(x = date, y = death),
                  size = 2)
```

b



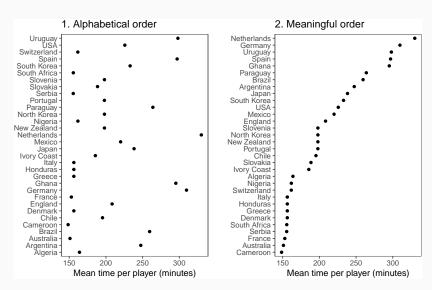
Order

Guideline 4: Make order meaningful.

You can make the ranking of data clearer from a graph by using order to show rank. Often, factor or categorical variables are ordered by something that is not interesting, like alphabetical order.

Order

Guideline 4: Make order meaningful.



Order



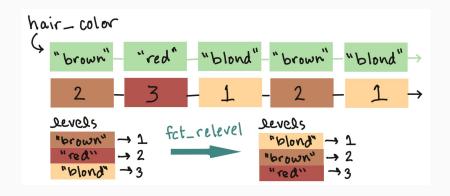
Order

You can re-order factor variables in a graph by resetting the factor using several functions from the forcats package.

The simplest is fct_relevel, which lets you change the levels "by hand".

See the "factors" cheatsheet for more forcats functions.

Changing factor levels



Order

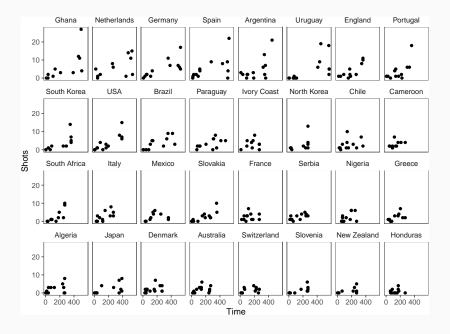
"And if you think that I'm going to miss even one opportunity to pick up half-a-mile boat speed, you're absolutely out of your mind. When it costs us nothing, when we give up nothing?! You're out of your mind."

Bruno, The West Wing, "The Indians in the Lobby"

Guideline 5: When possible, use small multiples.

Small multiples are graphs that use many small plots showing the same thing for different facets of the data. For example, instead of using color in a single plot to show data for males and females, you could use two small plots, one each for males and females.

Typically, in small multiples, all plots with use the same x- and y-axes. This makes it easier to compare across plots, and it also allows you to save room by limiting axis annotation.



You can use the facet functions to create small multiples. This separates the graph into several small graphs, one for each level of a factor.

The facet functions are:

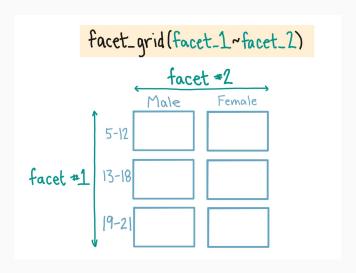
- facet_grid()
- facet_wrap()

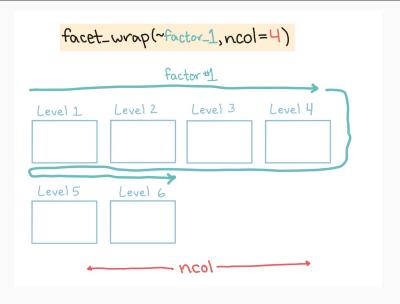
The facet_grid function can facet by one or two variables. One will be shown by rows, and one by columns:

```
## Generic code
facet_grid([factor for rows] ~ [factor for columns])
```

The facet_wrap() function can only facet by one variable, but it can "wrap" the small graphs for that variable, so the don't all have to be in one row or column:

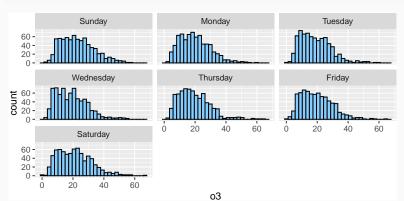
```
## Generic code
facet_wrap(~ [factor for faceting], ncol = [# of columns])
```





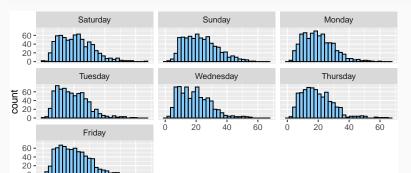
For example, to create small multiples of ozone concentration by day of week for the Chicago data, you can run:

```
chic %>%
  ggplot(aes(x = o3)) +
  geom_histogram(fill = "skyblue1", color = "black") +
  facet_wrap(~ dow)
```



To change the order of the facets, change the order of the factor levels for the factor you're using to facet. For example, to move weekend days together, you could run:

```
library(forcats)
chic %>%
  mutate(dow = fct_relevel(dow, "Saturday")) %>%
  ggplot(aes(x = o3)) +
  geom_histogram(fill = "skyblue1", color = "black") +
  facet_wrap(~ dow)
```



In-course exercise

We'll now take a break to do Section 3 of the In-Course Exercise (Chapter 4).

Advanced customization

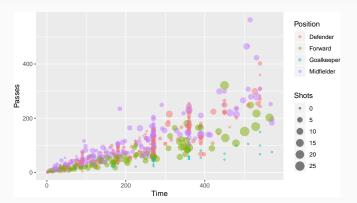
There are a number of different functions for adjusting scales. These follow the following convention:

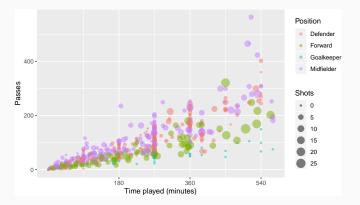
```
## Generic code
scale_[aesthetic]_[vector type]
```

For example, to adjust the x-axis scale for a continuous variable, you'd use scale_x_continuous.

You can use a scale function for an axis to change things like the axis label (which you could also change with xlab or ylab) as well as position and labeling of breaks.

For example, here is the default for plotting time versus passes for the worldcup dataset, with the number of shots taken shown by size and position shown by color:



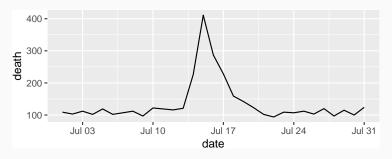


Parameters you might find useful in scale functions include:

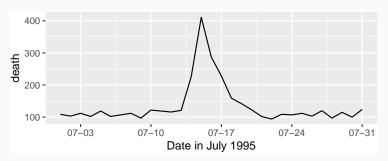
Parameter	Description
name	Label or legend name
breaks	Vector of break points
minor_breaks	Vector of minor break points
labels	Labels to use for each break
limits	Limits to the range of the axis

For dates, you can use scale functions like scale_x_date and scale_x_datetime. For example, here's a plot of deaths in Chicago in July 1995 using default values for the x-axis:

```
ggplot(chic_july, aes(x = date, y = death)) +
  geom_line()
```

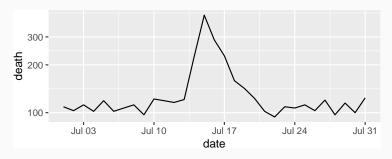


And here's an example of changing the formatting and name of the x-axis:



You can also use the scale functions to transform an axis. For example, to show the Chicago plot with "deaths" on a log scale, you can run:

```
ggplot(chic_july, aes(x = date, y = death)) +
  geom_line() +
  scale_y_log10()
```



For colors and fills, the conventions for the names of the scale functions can vary.

For example, to adjust the color scale when you're mapping a discrete variable (i.e., categorical, like gender or animal breed) to color, you'd use scale_color_hue. To adjust the color scale for a continuous variable, like age, you'll use scale_color_gradient.

For any color scales, consider starting with brewer first (e.g., scale_color_brewer).

Scale functions from brewer allow you to set colors using different palettes. You can explore these palettes at http://colorbrewer2.org/.

The Brewer palettes fall into three categories: sequential, divergent, and qualitative. You should use sequential or divergent for continuous data and qualitative for categorical data. Use display.brewer.pal to show the palette for a given number of colors.

```
library("RColorBrewer")

display.brewer.pal(name = "Set1", n = 8)

display.brewer.pal(name = "PRGn", n = 8)

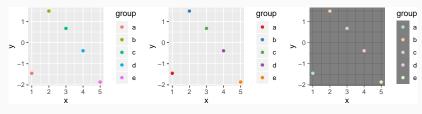
display.brewer.pal(name = "PuBuGn", n = 8)

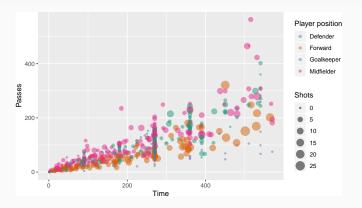
Set1 (qualitative)

PRGn (divergent)

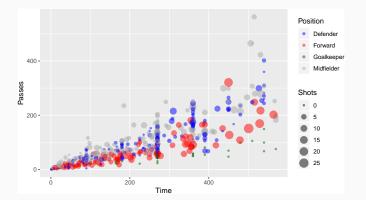
PuBuGn (sequential)
```

Use the palette argument within a scales function to customize the palette:





You can also set colors manually:



Excellent references

Chapter 3 of "R for Data Science", http://r4ds.had.co.nz/

Some excellent further references for plotting are:

- R Graphics Cookbook (book and website)
- Google images

For more technical details about plotting in R:

- ggplot2: Elegant Graphics for Data Analysis, Hadley Wickham
- R Graphics, Paul Murrell