This is from the fourth chapter of learn.r-journalism.com.

Let’s load some data, create a chart, and break down the layers.

We’ll some data from Vulture.com comparing ages of leading men in movies compared to their love interests.

```r
library(readr)

ages <- read_csv("data/ages.csv")

define(ages)
## # A tibble: 6 x 7
## Movie Genre actor actor_age actress actress_age budget
## <chr> <chr> <chr> <int> <chr> <int> <dbl>
## 1 Mo' Better Blues drama Denzel 35 Joie Lee 28 10
## 2 Malcolm X drama Denzel 37 Angela 34 34
## 3 The Preacher's Wife drama Denzel 41 Whitney 33 40
## 4 He Got Game drama Denzel 43 Milla J 22 25
## 5 Remember the Titans drama Denzel 45 Micole 29 30
## 6 Training Day drama Denzel 46 Eva Men 27 45
```

This is the data we’re working with.

The variables/columns are Movie, Genre, actor, actor_age, actress, actress_age, budget.

Here’s the chart. Run this in your console.

```r
# If you haven't installed the ggplot2 package yet, uncomment and run the line below
#install.packages("ggplot2")

library(ggplot2)

ggplot(data=ages) +
  geom_point(mapping=aes(x=actor_age, y=actress_age)) +
```
What’s this chart show? Men tend to be much older than women in movies.
Let’s break the components of this chart down.
We start with with the data frame. We initialized the data viz function `ggplot()` by passing the `ages` data frame to it.

```r
ggplot(data=ages) +
    geom_point(mapping=aes(x=actor_age, y=actress_age)) +
    expand_limits(x = 0, y = 0) +
    geom_abline(intercept=1, col="light gray")
```
Next, we have the dots representing the data we chose to visualize on the x and y axes. Mappings describe how aesthetics should relate to the variables in the data.

```r
ggplot(data=ages) + geom_point(mapping=aes(x=actor_age, y=actress_age)) + expand_limits(x = 0, y = 0) + geom_abline(intercept=1, col="light gray")
```

Set the aesthetics (`aes()`) visual characteristics that represent your data.

We are using the default geometric object and color but these are customizable:

- position
- size
- color
- shape
- transparency
- fill

For each aesthetic, we can set the scales for how visual characteristic is converted to display values. We’ll practice this later on.

Let’s take a look at the `geom_` function we add to `ggplot()`. 
Notice that \texttt{ggplot2} requires a plus between functions.

This package was created before \texttt{dplyr} and others that implemented the `%>%` piping.

It’s a weird quirk that I find myself messing up often since the shortcut for piping is pretty much muscle memory to me.

If you’re using \texttt{ggplot}: plus it!

For everything else: pipe it!

\begin{verbatim}
\texttt{ggplot(data=\texttt{\langle DATA\rangle}) +}
\texttt{\langle GEOM_FUNCTION\rangle(mapping=aes(\langle MAPPINGS\rangle))}
\texttt{ggplot(data=\texttt{\langle DATA\rangle}) +}
\texttt{geom_point(mapping=aes(x=actor\_age, y=actress\_age))}
\end{verbatim}

So \texttt{geom_point()} is merely one type out of dozens of possible \texttt{geom\_functions}, like \texttt{geom\_bar()} or \texttt{geom\_boxplot()}.

No matter the \texttt{geom\_function} you end up with, it still requires \texttt{mappings} to be passed to it.

In this chart’s instance, it’s \texttt{actor\_age} and \texttt{actress\_age} from the \texttt{ages} data frame.

This is basically all you need to create a chart in R. The \texttt{ggplot()} and the data and the \texttt{geom\_} and mapping from \texttt{aes()}. You don’t have to run the other two lines below from our example.

But we added more instructions for clarity.
The next line refers to some scaling options.

Scaling can also be passed to the `aes()` of the `geom_function` but can also apply to the entire chart.

```r
ggplot(data=ages) +
  geom_point(mapping=aes(x=actor_age, y=actress_age)) +
  expand_limits(x = 0, y = 0) +
  geom_abline(intercept=1, col="light gray")
```

In this chart, we used the `expand_limits()` function to force the x- and y-axis to start at 0.

Otherwise, `ggplot()` would shift the scale so that the points in the chart fill up the entire chart. It assumes you don’t want that extraneous white space that doesn’t include data.

But for this particular data, we want to emphasize the disparity in ages by starting at the zeroes. Plus, there is a large contingent of data visualization pros that might yell at you for not starting the base at zero (even though there might be some exceptions to that rule!).
The last line in the stack of functions we have so far is the `ab_line()` function. It’s not necessary, but it’s part of the layering options in `ggplot2`.

```r
ggplot(data=ages) +
  geom_point(mapping=aes(x=actor_age, y=actress_age)) +
  expand_limits(x = 0, y = 0) +
  geom_abline(intercept=1, col="light gray")
```

Each function in `ggplot2` like `geom_abline()` allows for additional variables that can be passed to it. In this example, we specify the intercept should be at 1 and the color of the bar should be “light gray”. We accept the default width of the line by not passing it the variable to the `geom_abline()` at all.
Okay, let’s make some more charts.

But first, consider the data.

**Data**

It’s crucial to understand the structure of the data you have when working with `ggplot2()`. We have six variables (columns) in this data frame. Each row represents data for a single movie. That’s pretty tidy.

It seems like an obvious format, but not all data sets have this structure by default.

Recall the previous chapter when we were dealing with wide versus long data?
The variables for each race are spread out into columns. This looks great in a spreadsheet but it doesn’t work well with `ggplot2` because it isn’t tidy.

<table>
<thead>
<tr>
<th>Metro</th>
<th>Race</th>
<th>Percent_Unsolved</th>
</tr>
</thead>
<tbody>
<tr>
<td>1886</td>
<td>Abilene, TX</td>
<td>15.38</td>
</tr>
<tr>
<td>1887</td>
<td>Abilene, TX</td>
<td></td>
</tr>
<tr>
<td>247</td>
<td>Alexandria, LA</td>
<td></td>
</tr>
</tbody>
</table>

That’s better. Each row is a different variable. This way, we could add another variable for percent solved or total cases per metro and race.

It’s simply more versatile.

Data works best in `ggplot()` in a tidy, long (tall) format.

**Geoms**

Let’s go over some new `geom_` types.

**Simple bar plot**

We’ll start with `geom_bar()` which is the most common type of chart you’ll need to make starting out.

Notice that we are nesting `aes()` into `ggplot()` this time and not using the `method=` or the `data=` because they are not necessary.
Start keeping track of how many variables we’re using in each of these charts.

For the one above, we started with one variable and sort of created a new one by counting the instances they appear in the data frame. We didn’t have to alter the data frame at all or summarize it for this chart to work.

Common `geom_bar()` options:

- `width`
- `fill`
- `color` (border)
- `position_dodge()`

Stacked bar plot

```r
ggplot(data=ages,
    aes(x=actor, fill=Genre)) +
    geom_bar()
```
Aesthetics

Before we move on to other chart types, let’s talk about what just happened in the chart above.

The only difference between the bar plot and the stacked bar plot was that, `fill=Genre` was added to the `aes()` function.

**Variable count:** 2 (actor & genre, if you don’t count the count of actor)

Here’s a reminder that these are the aesthetic properties that can be set:

- position
- size
- color
- shape
- transparency
- fill
What if we changed the `aes()` from `fill` to `color`?

```
ggplot(data=ages,
      aes(x=actor, color=Genre)) +
geom_bar()
```

Yikes, not as effective. But now you know about the option.

When would this be a good option? Line charts or dots.

Notice how the color and legend was added automatically?
This is the default color list. There are ways to customize the colors, but we’ll go over that in the next section.

Let’s go back to `fill=Genre` but this time with a twist.

Take a look at the code and see if you can spot why it didn’t work like it did earlier.

```r
ggplot(data=ages,
       aes(x=actor), fill=Genre) +
       geom_bar()
```
Do you see it?

It’s the parenthesis in `aes()` – it closed off before including `fill=Genre` so it wasn’t included. Try it this time correctly.

```r
ggplot(data=ages, 
      aes(x=actor, fill=Genre)) + 
      geom_bar()
```
Grouped bar plot

```r
ggplot(data=ages,
    aes(x=actor, fill=Genre)) +
geom_bar(position="dodge")
```

So this chart was similar to the stacked bar plot above, but this time `position="dodge"` was passed to the `geom_bar()` function.

This tells ggplot to group the bar plot.

This is useful for easier comparison within groups.

Spinogram

It’s simple to turn a stacked bar plot into a percent-based chart.

```r
ggplot(data=ages,
    aes(x=actor, fill=Genre)) +
geom_bar(position="fill")
```
This type of chart allows for a greater understanding of the proportion compared to other groups.

**Box plot**

You’ll see this type of chart often in research papers.

It succinctly summarizes the distribution of numbers in each category we’ve set: actors.

```r
ggplot(ages, aes(x=actor, y=actress_age)) + geom_boxplot()
```
The height of the line is the max and min of the numbers in the y axis: actress ages.

The top of the box is the top quartile and the bottom of the box is the bottom quartile. The line in the middle is the median age. That floating dot? Considered an outlier.

Common `geom_boxplot()` options:

- fill
- color
- notch (=TRUE or FALSE)
- outlier. color shape size

Violin plot

This is another way to show distribution of numbers.

```r
ggplot(ages, aes(x=actor, y=actress_age)) + geom_violin()
```
The wider the diameter, the more numbers there are. It’s a bit more intuitive to follow.

**Variable count:** 2 - *actress_age* and *actor*

**Scaling**

Useful to transform or rescale the data with.

**geom_histogram**

```r
ggplot(data=ages, aes(x=actor_age)) + geom_histogram(binwidth=1)
```
Common `geom_histogram()` options:

- `binwidth`
- `color` (border)
- `fill`

**Applying a logarithmic scale**

It makes more sense to apply scalar transformations to while plotting rather than altering the data set itself.

```r
 ggplot(data=ages, aes(x=actor_age)) + 
    geom_histogram() + scale_x_log10()
```

## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
Kernel density plot

```r
ggplot(data=ages,
      aes(x=actress_age)) +
geom_density(fill="red")
```
Common `geom_density()` options:

- fill
- color
- alpha

```r
ggplot(data=ages,
       aes(x=actress_age, fill=actor)) +
  geom_density(alpha=.3)
```
Line plot

Let’s start by summarizing the ages of the actresses per actor.

```r
library(dplyr)

avg_age <- ages %>%
```
group_by(actor) %>%
mutate(age_diff = actor_age - actress_age) %>%
summarize(average_age_diff = mean(age_diff))

ggplot(avg_age, aes(x=actor, y=average_age_diff, group=1)) +
  geom_line()

Variable count: 2 - the new mutated variable `average_age_diff` and `actor`

Don’t forget, we can add more layers.

Let’s make the same chart above but with dots, too.

ggplot(avg_age, aes(x=actor, y=average_age_diff, group=1)) +
  geom_line() +
  geom_point()
Does this make sense as a visualization? Mmm... nah.

Never use a line chart to chart anything across an axis that doesn’t represent something continuous.

Common `geom_point()` options:

- color
- fill
- alpha
- shape
- size

Scatterplot with fit

We’ve made a scatter plot before.

This time, let’s add a `geom_smooth()` layer.

```r
ggplot(ages,
      aes(x=actor_age,
           y=actress_age)) +
geom_point() +
geom_smooth()
```

```r
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
```
We can customize it and throw in a linear regression line. Notice the ~ which is used often in statistical analysis formulas.

```r
ggplot(ages,
      aes(x=actor_age,
          y=actress_age)) +
  geom_point() +
  geom_smooth(method="lm",
              formula=y~x)
```
Grouping

ggplot(data=ages, 
    aes(x=actor_age, 
        y=actress_age, 
        color=actor)) + 
geom_point()
Variable count: 3! We’re working with `actor_age`, `actress_age`, and `actor`.

See how we’re able to start packing more information into a chart with a line of code?

When you’re exploring data visually, this might help surface insights you wouldn’t notice by just looking at the raw data.

Everything’s still pretty clustered in this chart so you might only be able to notice outliers like Tom Cruise and Johnny Depp on the bottom left. But everything in the middle is kinda just too much.

Let’s go crazy and add another variable.

```r
ggplot(data=ages,
       aes(x=actor_age,
           y=actress_age,
           color=actor,
           shape=Genre)) +
geom_point()
```
Variable count: 4! We’re working with `actor_age`, `actress_age`, `actor`, and now `Genre`.

This looks not good.

Why did we do this?? I just wanted to show you that the shapes option exist. You’ll probably never use it.

Okay, what if we try keeping the color and shapes to one variable: `actor`?

```r
ggplot(data=ages,
       aes(x=actor_age,
           y=actress_age,
           color=actor,
           shape=actor)) +
geom_point()
```

## Warning: The shape palette can deal with a maximum of 6 discrete values
## because more than 6 becomes difficult to discriminate; you have 7.
## Consider specifying shapes manually if you must have them.

## Warning: Removed 10 rows containing missing values (geom_point).
Nope, still not great. Plus, there are only a limited amount of shapes to pull from.

Did you see that warning?

```r
## Warning: Removed 10 rows containing missing values (geom_point).
```

That’s because it couldn’t find a shape for Tom Hanks. Hanks was left off the chart.

Don’t you feel horrible? Tom Hanks is the nicest guy ever.

Like colors, you should limit the palette so it doesn’t confuse readers.

**Scatterplot with color and size**

Let’s swap out *Genre* for another variable that might make more sense to add to the visualization: *budget*.

```r
ggplot(data=ages,
      aes(x=actor_age,
           y=actress_age,
           color=actor,
           size=budget)) +
geom_point()
```
Variable count: 4! We’re working with actor_age, actress_age, actor, and now Genre.
Lots of data being communicated to readers here. The legends were auto-generated.

Coords
You’ll see stats in the geom_bar() function. This is the statistical transformations that summarize data. In this instance we’re setting it to "identity" because we want the bars to represent the numbers in the y in aes(), not the count of it.

- counts
- means
- trend lines

We’re going to alter the coordinates of the chart so that we get a horizontal chart instead of the default vertical one.

The coord_flip() option flips a chart you’ve been working on from vertical to horizontal or horizontal to vertical.

```r
ggplot(data = avg_age, aes(x = actor, y = average_age_diff)) + geom_bar(stat = "identity") + coord_flip()
```
Facets

Remember how that big scatter plot above had all the different colors on one viz and it was all clustered together?

It’s simple to break those out into individual charts.

This is sometimes referred to as “small multiples”.

```r
ggplot(data=ages) +
  geom_point(mapping=aes(x=actor_age, y=actress_age)) +
  expand_limits(x = 0, y = 0) +
  geom_abline(intercept=0, col="light gray") +
  facet_grid(Genre ~ actor)
```
Variable count: 4! - We’re working with `actor_age`, `actress_age`, `actor`, and also `Genre`.

Note: Take a moment to appreciate how easy this was. This is a pain in the ass in other languages, like D3.

Want to add another variable? Let’s do it. Add `size=budget` the `geom_point()`.

```r
ggplot(data=ages) +
  geom_point(mapping=aes(x=actor_age, y=actress_age, size=budget)) +
  expand_limits(x = 0, y = 0) +
  geom_abline(intercept=0, col="light gray") +
  facet_grid(Genre~actor)
```
**Variable count:** 5! - We're working with `actor_age`, `actress_age`, `actor`, and also `Genre` AND `budget`. The sizing kinda jumbles up the circles in some, but keep playing around with alpha or change the size of the chart, maybe. You can see it's quite easy to play around with the options until you find the right balance.

This is all possible thanks to the tidiness of the data structure.

The variables might contain more than one group but it's all tied back to rows with each individual movie.

Here are other types of ways to create “small multiples” through facets.

- `facets_grid(genre~actor)`
- `facets_grid(. ~ actor)` - just columns
- `facets_grid(actor ~ .)` - just rows
- `facets_wrap(~ var, ncol=#)` - one classification variable wrapped to fill page

```r
ggplot(data=ages) +
  geom_point(mapping=aes(x=actor_age, y=actress_age)) +
  expand_limits(x = 0, y = 0) +
  geom_abline(intercept=0, col="light gray") +
  facet_grid(.-actor)
```
You don’t have to use two variables in `facet_grid()` – you can just use one and it’ll break out the charts based on where the ~ is in relation to the variable and the ..

The chart above had `facet_wrap(.~actor)` so the grid is split vertically.

If it was swapped to `facet_wrap(actor~.)` then the grid will split by row.

```r
ggplot(data=ages) +
  geom_point(mapping=aes(x=actor_age, y=actress_age)) +
  expand_limits(x = 0, y = 0) +
  geom_abline(intercept=0, col="light gray") +
  facet_grid(actor~.)
```
I find myself using `facet_wrap()` more often because I can customize the number of columns in the small multiples with `ncol=`.

```r
ggplot(data=ages) +
  geom_point(mapping=aes(x=actor_age, y=actress_age)) +
  expand_limits(x = 0, y = 0) +
  geom_abline(intercept=0, col="light gray") +
  facet_wrap(~actor, ncol=4)
```
Global versus Local

The layers in your `ggplot()` chart can be customized in nuanced ways. Everything nested in the initial `ggplot()` call will be considered global and apply to the rest of the layers.

```r
ggplot(ages,
      aes(x=actor_age,
          y=actress_age)) +
geom_point()
```
So in the chart above, `aes()` with `actor_age` and `actress_age` are in the x and y globally.

But I can also pass `aes()` to the `geom_point()` layer, which will make that local and apply to that layer only.

```r
ggplot(ages,
      aes(x=actor_age,
          y=actress_age)) +
      geom_point(aes(color=actor))
```
You can also set data locally. This is very useful if you want to highlight a specific subset from the data.

```r
# Let's make a copy of the minus the actor variable so it can't be faceted
ages_copy <- ages %>% select(-actor)

ggplot(data = ages,
       aes(x = actor_age,
           y = actress_age,
           color = actor)) +
    geom_point(data = ages_copy, color = "grey") +
    geom_point() +
    facet_wrap(~ actor) +
    theme(legend.position = "none") # This removes the legend
```
We were able to use different data sets. Globally, we wanted to use the ages data frame because we were going to facet it.

We needed a second data set without the actor variable to plot in a background layer and set the color manually to “grey” (go back to the code above and this time put color="grey" in aes() and see the difference). This was the first geom_point() layer referenced locally.

The second geom_point() built a scatter plot based on the global data set in ggplot().

This type of chart really improves on that first grouped scatter plot because it makes it easier to see each individual group in the context to the rest of the data.

Your turn

Challenge yourself with these exercises so you’ll retain the knowledge of this section.

Instructions on how to run the exercise app are on the intro page to this section.

Next steps

With some practice and understanding of basic geom-types you can quickly iterate through different ways to visualize your data.

Once your data is structured correctly, you can use ggplot2 to slice, group, and facet data visually multiple ways until you find something worth elaborating on.
What is being communicated? What information is left out or obscured?

Going through these examples, you saw how shapes, layout, the number of variables, etc, all matter. Swapping out geoms and aesthetics. Different combinations communicated better than others. It all depends on the structure of the data and what it contains.

By visualizing data with the grammar of graphics, you’ll approach data in a flexible new way.

Great, now let’s do more with charts including how to move beyond exploring the data visually and how to turn get these charts to publication quality.