

Exploring data with R

IMT 573A - Data Science 1 - Theoretical Foundations

6-Oct-2020 (Week 2, Day 3)

Our Zoom class sessions will be recorded.

Today's topic

- Review - checklist of exploratory data analysis
- Markdown and Rmarkdown
- Data visualization for exploring data
- ggplot basics
- Lab2

DUE

- Lab2: upload by tomorrow (Oct 9) 9am PT
- PS1: deadline next Tuesday (Oct 13) 9am PT

Exploring Data: A checklist

1. Interpret your data
2. Formulate your question
3. Read in your data
4. Examine your data, look at the top and bottom of your data, look at structure
5. Tidy data (Data cleaning)
6. Try the easy solution first
7. Challenge your solution, validate with external data
8. Follow up with new interesting questions/directions

Think about your data! Think about your question!

RMarkdown

Writing in Markdwon

Markdown is a lightweight [markup language](#) that is used to format and structure text. It is a kind of “code” that you write in order to *annotate* plain text

```
This is a paragraph in which we'll add
**bold text**, _italicized text_, and `code`
into the middle of a sentence
```

This is a paragraph in which we'll add **bold text**, *italicized text*, and `code` into the middle of a sentence

Syntax	Formatting
<code>_text_</code>	<i>italicized</i> using underscores (<code>_</code>)
<code>**text**</code>	bolded using two asterisks (<code>*</code>)
<code>`text`</code>	inline <code>code</code> with backticks (<code>`</code>) text ~~strike-through~~ using tildes (<code>~`</code>)

Markdown

Writing in Markdown

Text Blocks

```
# Top Level header
## Second Level Header

Here is a normal paragraph

- List item 1
- List item 2
- List item 3

...
block of code
across multiple lines
...

> Here is a block quote
```

Top Level header

Second Level Header

Here is a normal paragraph

- List item 1
- List item 2
- List item 3

```
block of code
across multiple lines
```

Here is a block quote

Syntax	Formatting
#	Header (use ## for 2nd-level, ### for 3rd, etc.)
````	Code section (3 back ticks) that encapsulate the code
-	Bulleted/unordered lists (hyphens)
>	Block quote

```
```{r import titanic data}
# Let's write some R code here
```
```

Markdown can even make [tables](#), create [hyperlinks](#), and include [images](#)!

# Rmarkdown (review)

## .Rmd Content

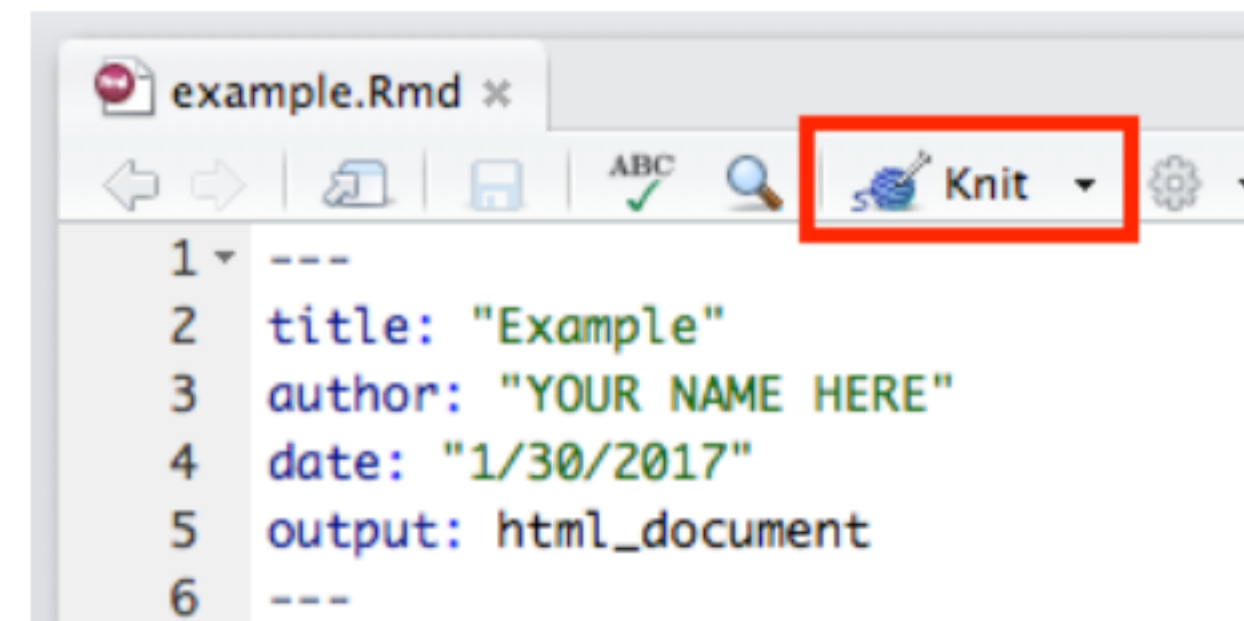
### Header information

```

title: "Example"
author: "YOUR NAME HERE"
date: "1/30/2017"
output: html_document

```

### RStudio knitting



RStudio's Knit button

### R code chunks

```
```{r}
# Execute R code in here
x <- 201
```
```

### Markdown lists

```
```{r echo=FALSE}
markdown.list <- "
- Lions
- Tigers
- Bears
"
```

`r markdown.list`
```

Would output a list that looks like:

- Lions
- Tigers
- Bears

### Rendering Tables

```
Including Tables
```{r echo=FALSE}
library(knitr) # make sure you load this library (once per doc)

# make a data frame
letters <- c("a", "b", "c")
numbers <- 1:3
df <- data.frame(letters = letters, numbers = numbers)

# render the table
kable(df)
```
```

| letters | numbers |
|---------|---------|
| a       | 1       |
| b       | 2       |
| c       | 3       |



# R Markdown Cheat Sheet

learn more at [rmarkdown.rstudio.com](http://rmarkdown.rstudio.com)



**.Rmd files**  
An R Markdown (.Rmd) file is a record of your research. It contains the code that a scientist needs to reproduce your work along with the narration that a reader needs to understand your work.

**Reproducible Research**  
At the click of a button, or the type of a command, you can rerun the code in an R Markdown file to reproduce your work and export the results as a finished report.

**Dynamic Documents**  
You can choose to export the finished report as a html, pdf, MS Word, ODT, RTF, or markdown document; or as a html or pdf based slide show.

### Workflow

- 1 Open a new .Rmd file** at File ► New File ► R Markdown. Use the wizard that opens to pre-populate the file with a template
- 2 Write document** by editing template
- 3 Knit document to create report** Use knit button or `render()` to knit
- 4 Preview Output** in IDE window
- 5 Publish (optional)** to web or server  
Synch publish button to accounts at  
• [rpubs.com](http://rpubs.com),  
• [shinyapps.io](http://shinyapps.io)  
• RStudio Connect  
Reload document  
Find in document  
File path to output document
- 6 Examine build log** in R Markdown console
- 7 Use output file** that is saved alongside .Rmd

### .Rmd structure

**YAML Header**  
Optional section of render (e.g. pandoc) options written as key:value pairs (YAML).  
• At start of file  
• Between lines of ---

**Text**  
Narration formatted with markdown, mixed with:

**Code chunks**  
Chunks of embedded code. Each chunk:  
• Begins with `{r}`  
• ends with `}`  
R Markdown will run the code and append the results to the doc.  
It will use the location of the .Rmd file as the **working directory**

**Workflow Diagram:** Open in window → Save → Spell Check → Find and replace → Publish → Show outline → R Markdown report.html

**Code chunks actions:** Set preview location, Insert code chunk, Go to code chunk, Run code chunk(s), Modify chunk options, Run all previous chunks, Run current chunk

**render()**  
Use `rmarkdown::render()` to render/knit at cmd line.  
Important args:  
**input** - file to render  
**output_format**  
**output_options** - List of render options (as in YAML)  
**output_file**  
**output_dir**  
**params** - list of params to use  
**envir** - environment to evaluate code chunks in  
**encoding** - of input file

### Interactive Documents

Turn your report into an interactive Shiny document in 4 steps

- 1 Add runtime: shiny** to the YAML header.
- 2 Call Shiny input functions** to embed input objects.
- 3 Call Shiny render functions** to embed reactive output.
- 4 Render with `rmarkdown::run`** or click **Run Document** in RStudio IDE

Embed a complete app into your document with `shiny::shinyAppDir()`

* Your report will be rendered as a Shiny app, which means you must choose an html output format, like `html_document`, and serve it with an active R Session.

**Shiny Example:** How many cars? (Input: 5) → Table with speed and dist columns

### Embed code with knitr syntax

**Inline code**  
Insert with ``r <code>``. Results appear as text without code.  
Built with ``r getRversion()`` → Built with 3.2.3

**Code chunks**  
One or more lines surrounded with `{r}` and `}`. Place chunk options within curly braces, after `r`. Insert with `{ }`  
`{r echo=TRUE} getRversion() }` → `getRversion()` → `[1] '3.2.3'`

**Global options**  
Set with `knitr::opts_chunk$set()`, e.g.  
`{r include=FALSE} knitr::opts_chunk$set(echo = TRUE) }`

**Important chunk options**

|                                                                              |                                                                    |                                                                       |                                                                                                        |
|------------------------------------------------------------------------------|--------------------------------------------------------------------|-----------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------|
| <b>cache</b> - cache results for future knits (default = FALSE)              | <b>dependson</b> - chunk dependencies for caching (default = NULL) | <b>fig.align</b> - 'left', 'right', or 'center' (default = 'default') | <b>message</b> - display code messages in document (default = TRUE)                                    |
| <b>cache.path</b> - directory to save cached results in (default = "cache/") | <b>echo</b> - Display code in output document (default = TRUE)     | <b>fig.cap</b> - figure caption as character string (default = NULL)  | <b>results</b> (default = 'markup')<br>'asis' - passthrough results<br>'hide' - do not display results |
| <b>child</b> - file(s) to knit and then include                              | <b>engine</b> - code language used in chunk                        | <b>fig.height</b> , <b>fig.width</b> - Dimensions of                  |                                                                                                        |

### Pandoc's Markdown

Write with syntax on the left to create effect on right (after render)

**Plain text**  
End a line with two spaces to start a new paragraph.  
*italics* and **bold**  
`verbatim code`  
sub/superscript²  
~~strikethrough~~  
escaped: * _ \\  
endash: --, emdash: ---  
equation:  $A = \pi * r^2$   
equation block:  
$$E = mc^2$$

**Block quote**  
> block quote

**Header**  
# Header1 {#anchor}  
## Header 2 {#css_id}  
### Header 3 {css_class}  
#### Header 4  
##### Header 5  
##### Header 6  
<!--Text comment-->  
\textbf{Tex ignored in HTML}  
<em>HTML ignored in pdfs</em>  
<http://www.rstudio.com>  
[link](www.rstudio.com)  
Jump to [Header 1]({#anchor})  
image:  
![Caption](smallorb.png)

**Lists**  
* unordered list  
+ sub-item 1  
+ sub-item 2  
- sub-sub-item 1  
* item 2  
Continued (indent 4 spaces)  
1. ordered list  
2. item 2  
i) sub-item 1  
A. sub-sub-item 1  
(@) A list whose numbering continues after  
(@) an interruption  
Term 1  
: Definition 1  
Right Left Default Center  
12 12 12 12  
123 123 123 123  
1 1 1 1  
- slide bullet 1  
- slide bullet 2  
(>- to have bullets appear on click)  
horizontal rule/slide break:  
***  
A footnote [^1]  
[^1]: Here is the footnote.

**Parameters**  
Parameterize your documents to reuse with different inputs (e.g., data sets, values, etc.)  
1 **Add parameters**  
Create and set parameters in the header as sub-values of **params**  
params:  
n: 100  
d: !r Sys.Date()  
2 **Call parameters**  
Call parameter values in code as `params$name`  
Today's date is `{r params$d}`

When you render, 1. runs the R code, 2. then converts the

Set a document default output in the YAML

**output v**  
html_document  
pdf_document  
word_document  
odt_document  
rtf_document  
md_document  
github_document  
ioslides_presentation  
slidy_presentation  
beamer_presentation

Customize output with sub-options (listed at right)

**html tabs**  
Use .tabset class to place sub-into tabs  
# Tabset {.tabset}  
## Tab 1  
text 1  
## Tab 2  
text 2  
### End tabs

### Create

- 1 Create inst/r
- 2 In the
- 3 Install
- 4 Access template wizard New File R Mark

template.yaml  
name: My Temp



# Data visualization basics (needed for exploring data)

Human brain is exceptionally good at visual pattern

Map dimensions of data to visual characteristics

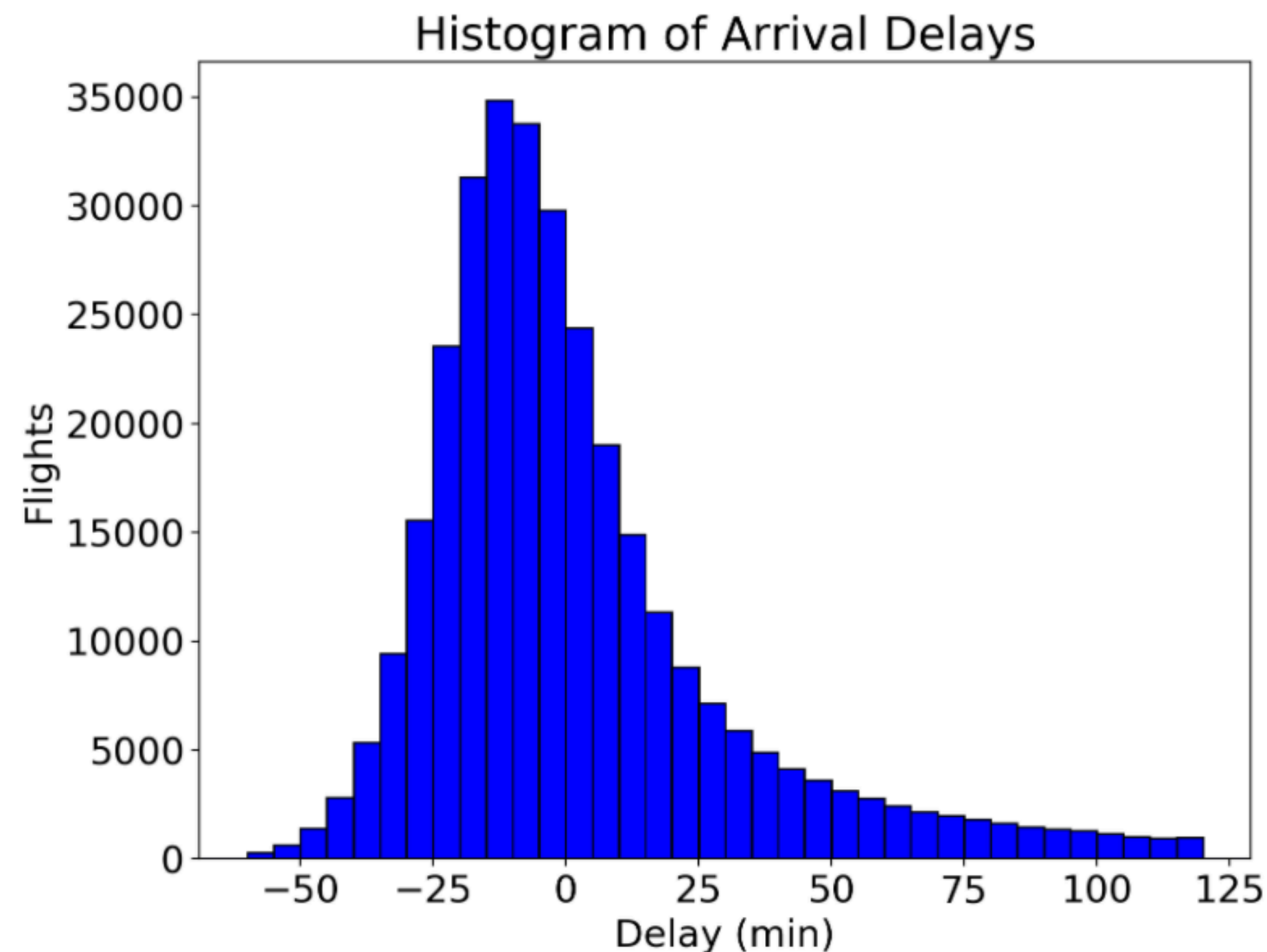
- Location
- Size/Length
- Color
- Shape



# Common visualizations for low dimensional data (1D/2D)

**Univariate** - single variable, 1D data

Visualization — Histograms/KDE

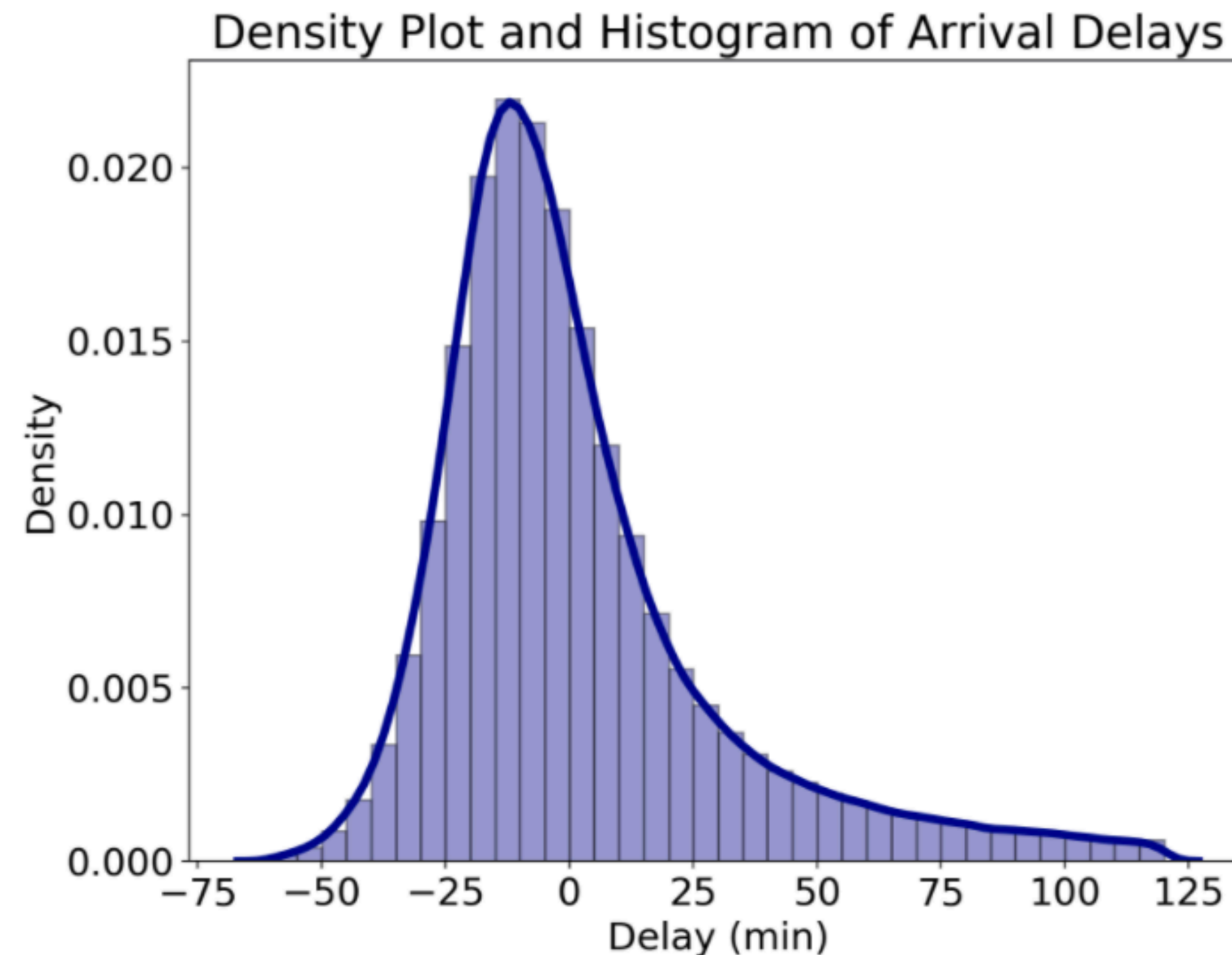


# Density plot (KDE)

A smoothed, continuous version of a histogram estimated from data.

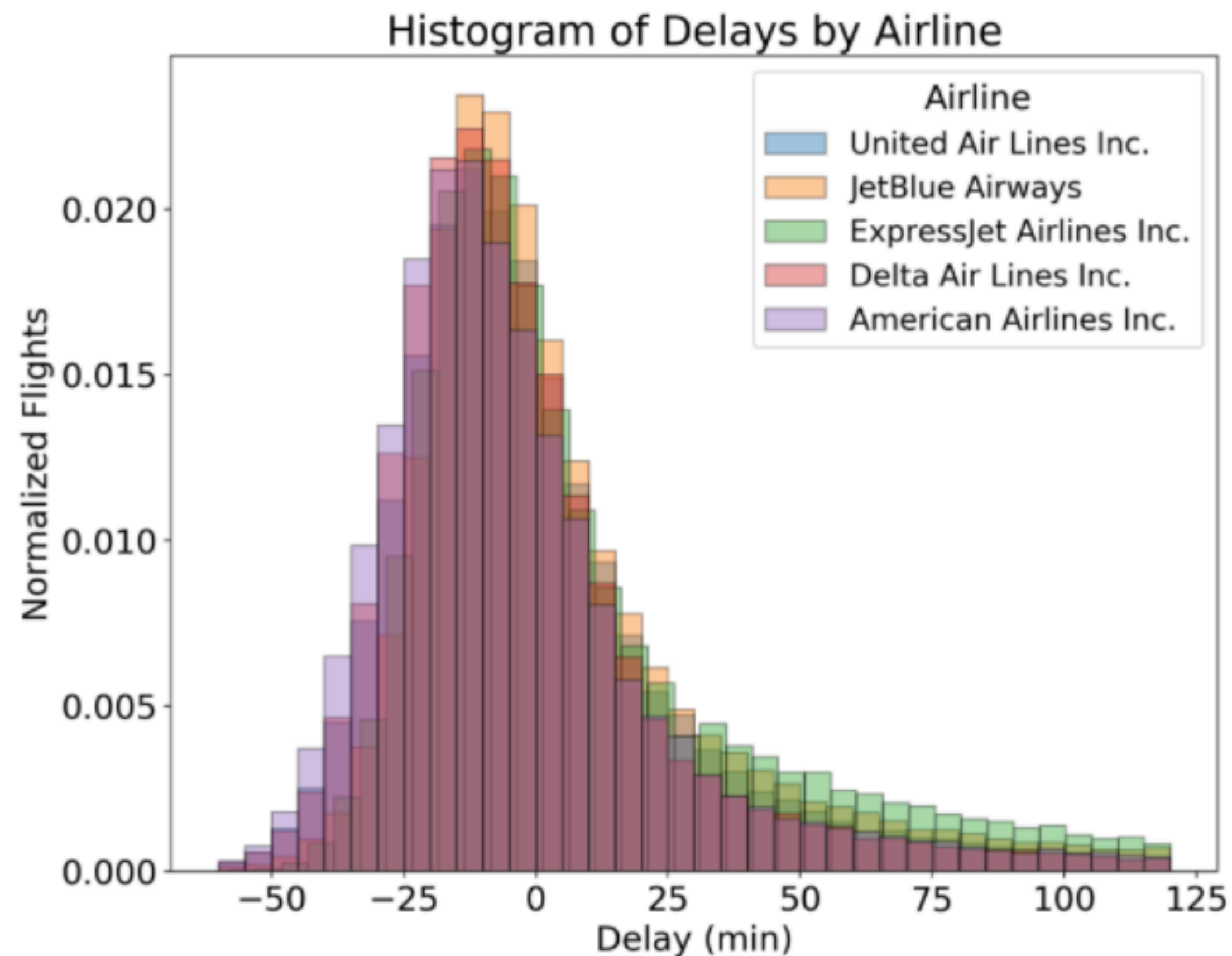
**KDE** - most common form of estimation

a continuous curve (the kernel) is drawn at individual data point and all of these curves are then added together to make a single smooth density estimation.

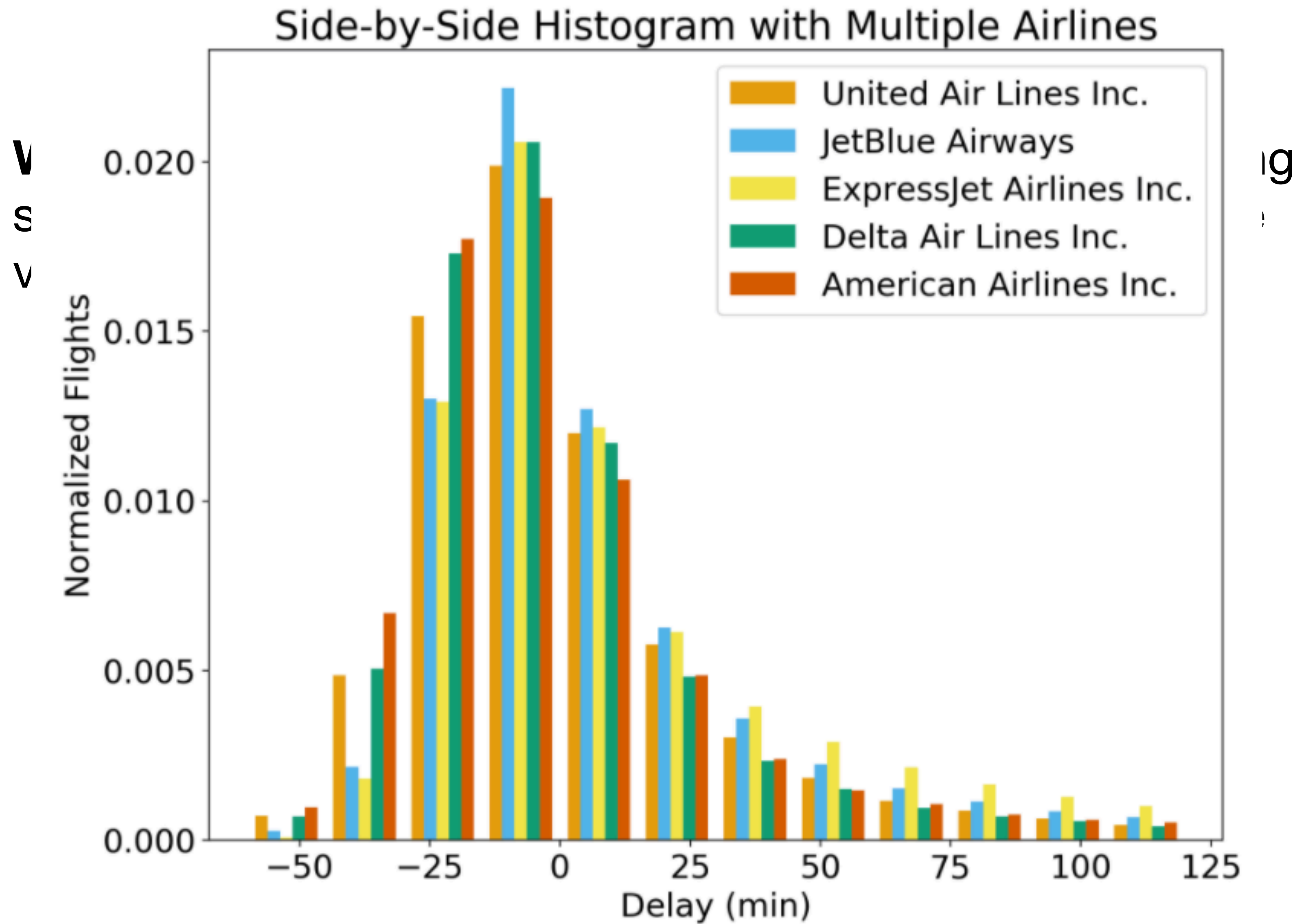


# Histograms & KDE

**When histograms fail** — Histograms are great for exploring single variable from one category, but not great for a single variable across multiple categories.

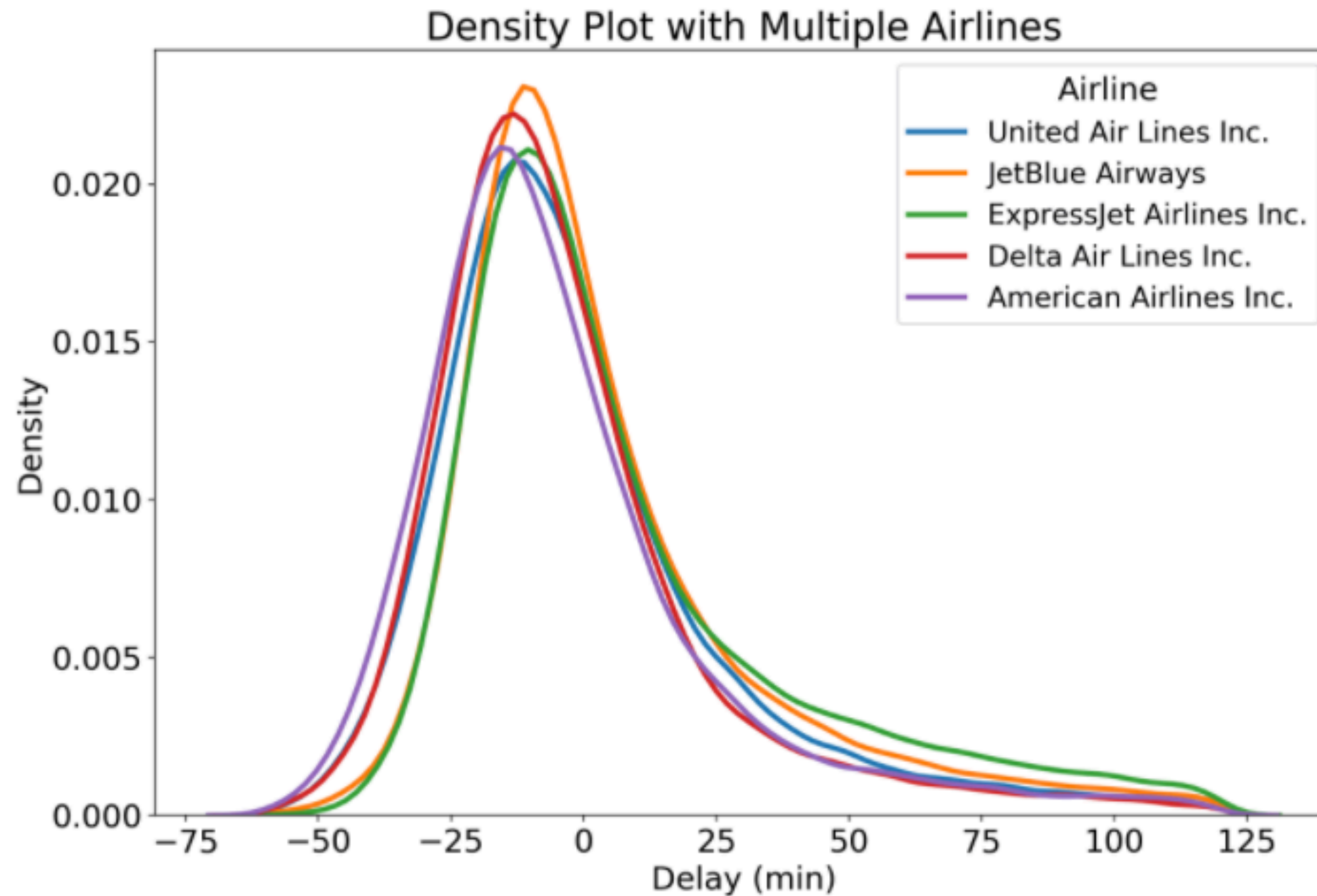






# Density plot

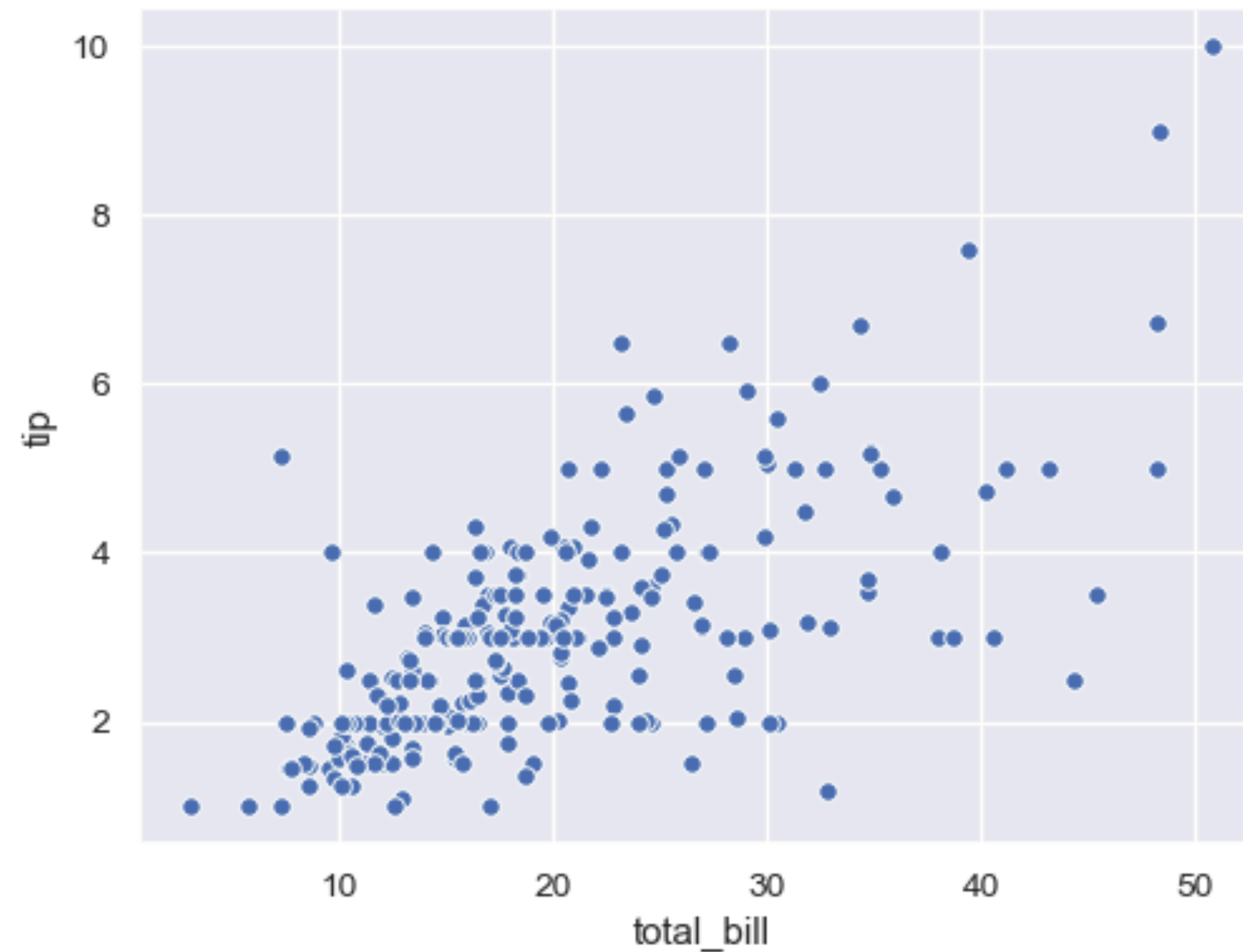
A smoothed, continuous version of a histogram estimated from the data.



# Common visualizations for low dimensional data (1D/2D)

**Bivariate** - 2 variables.

**Scatterplot** - Qualitative (continuous) x Quant (continuous), correlation

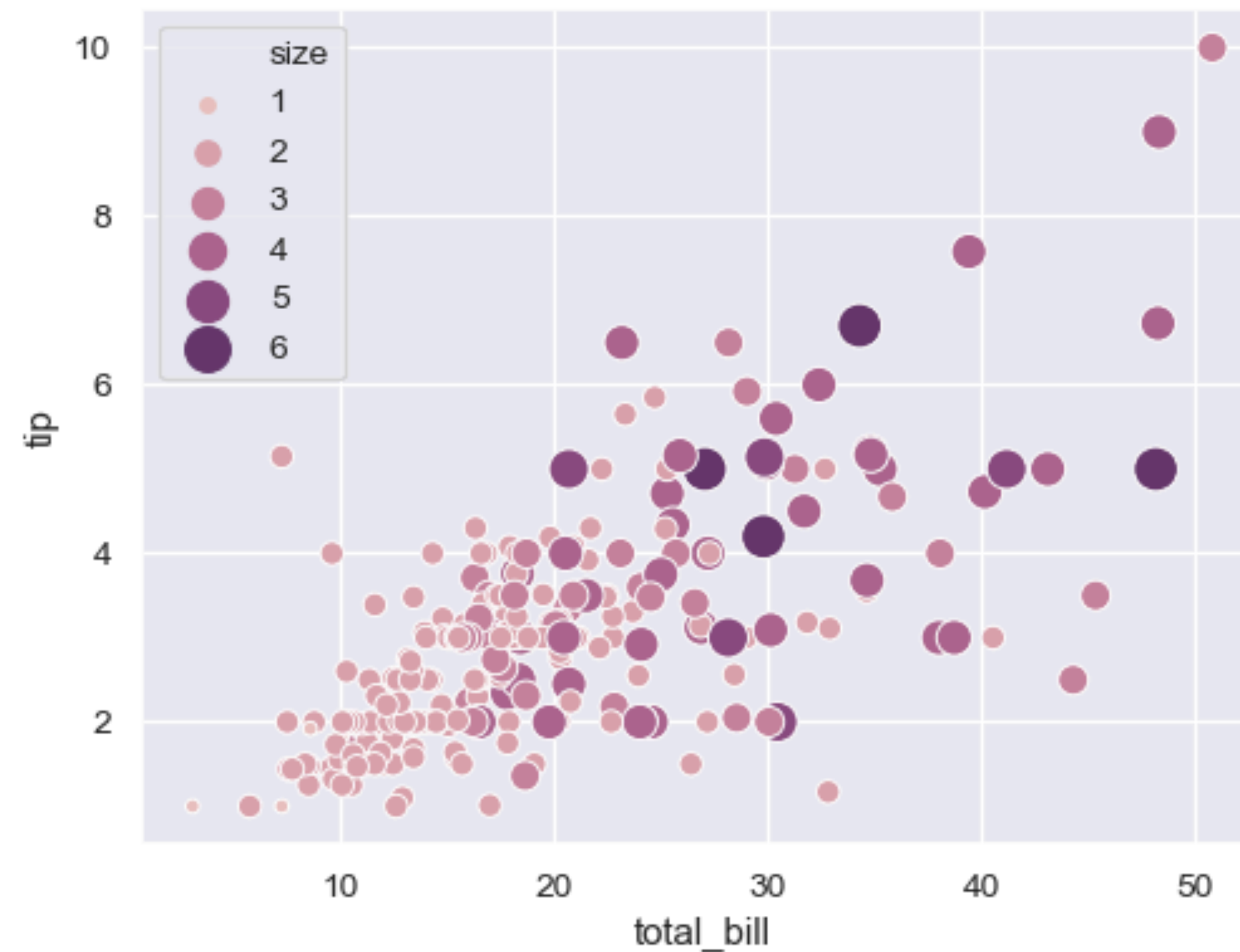




# Common visualizations for low dimensional data (1D/2D)

**Bivariate** - 2 variables.

**Scatterplot** - Qualitative (continuous) x Quant (continuous), correlation

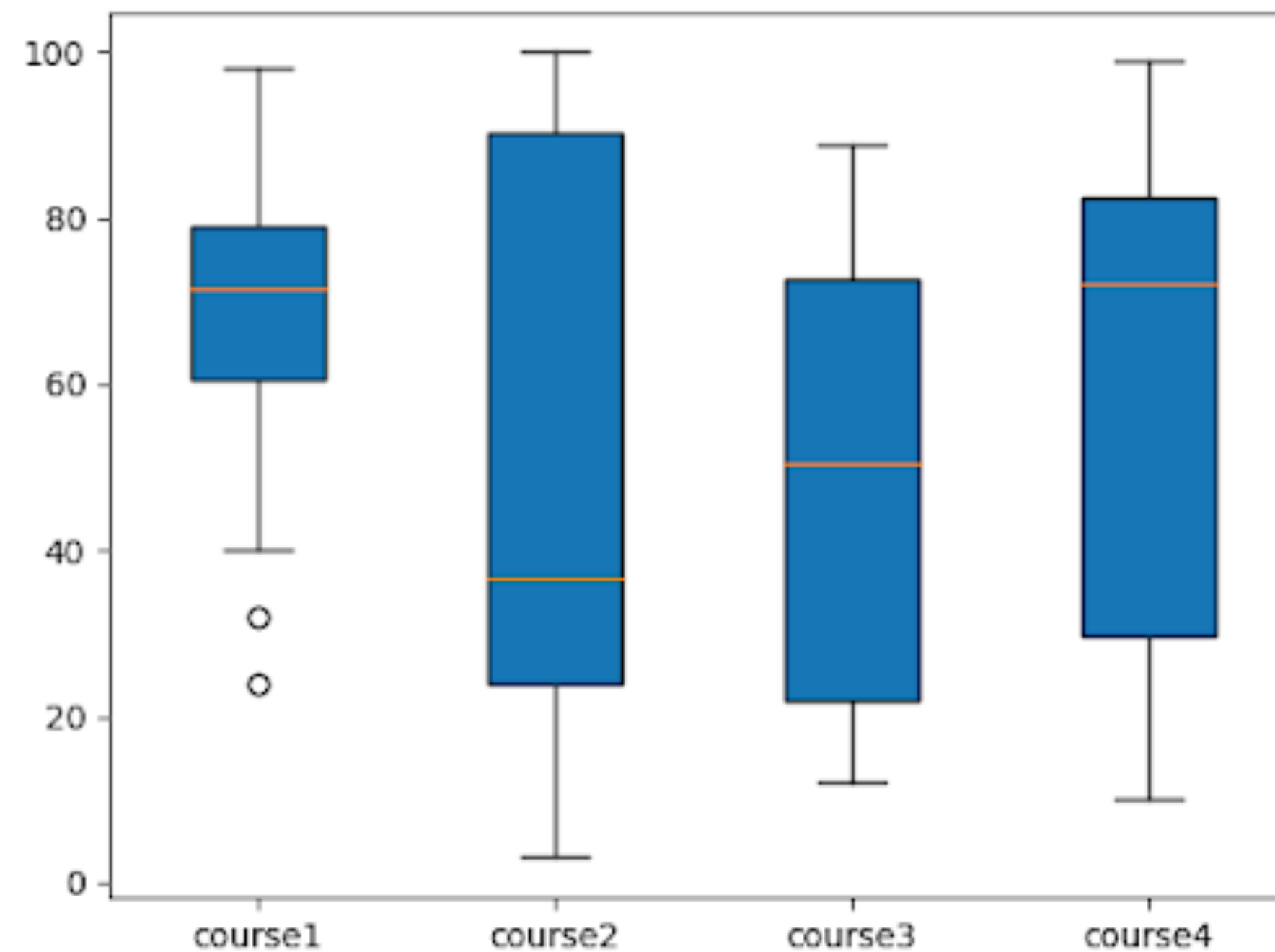


# Common visualizations for low dimensional data (1D/2D)

**Bivariate** - 2 variables.

**Boxplot** - Continuous x Categorical

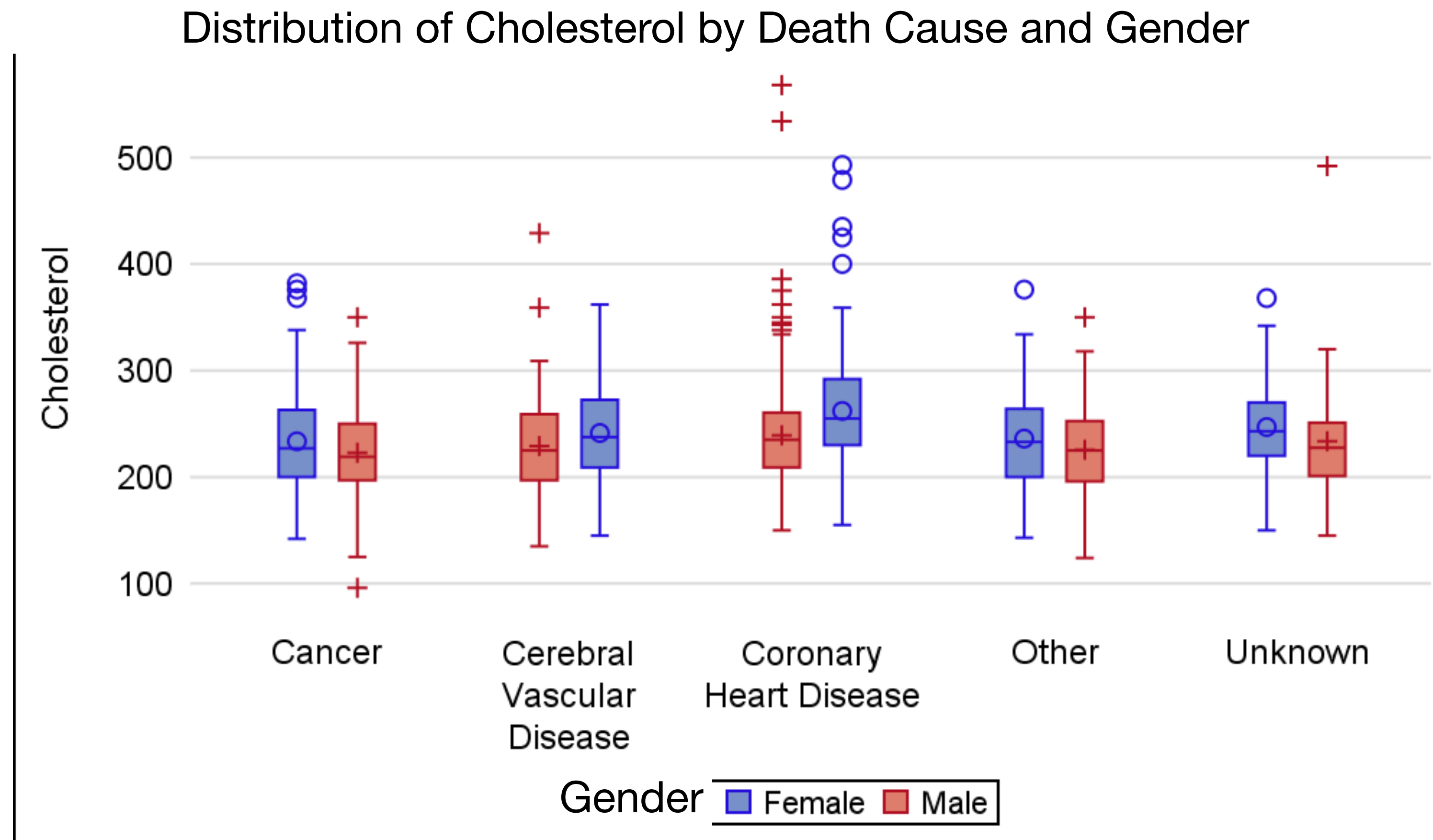
**Grade** Distribution by Course



# Common visualizations for low dimensional data (1D/2D)

**Bivariate** - 2 variables.

**Boxplot** - Continuous x Categorical

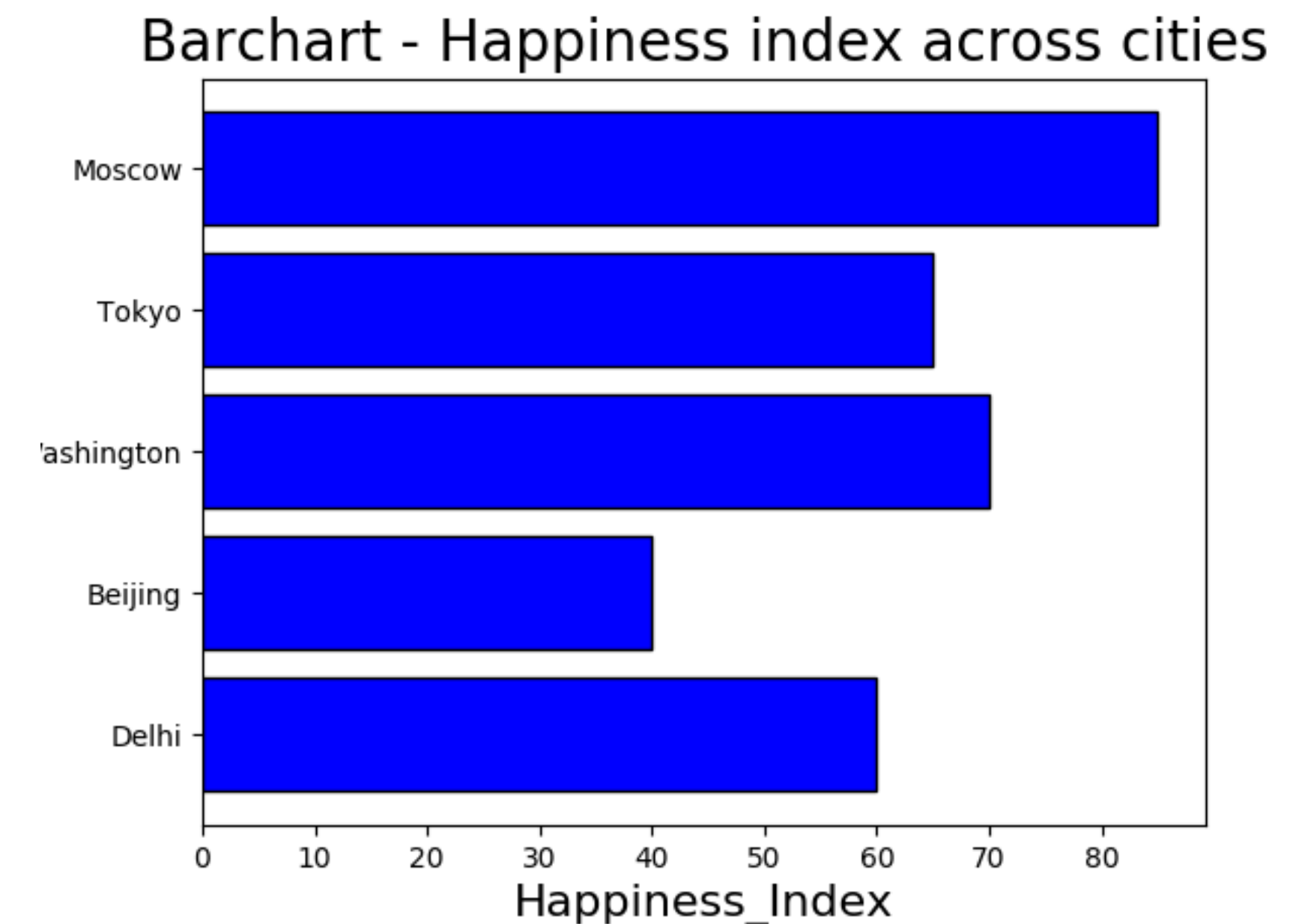
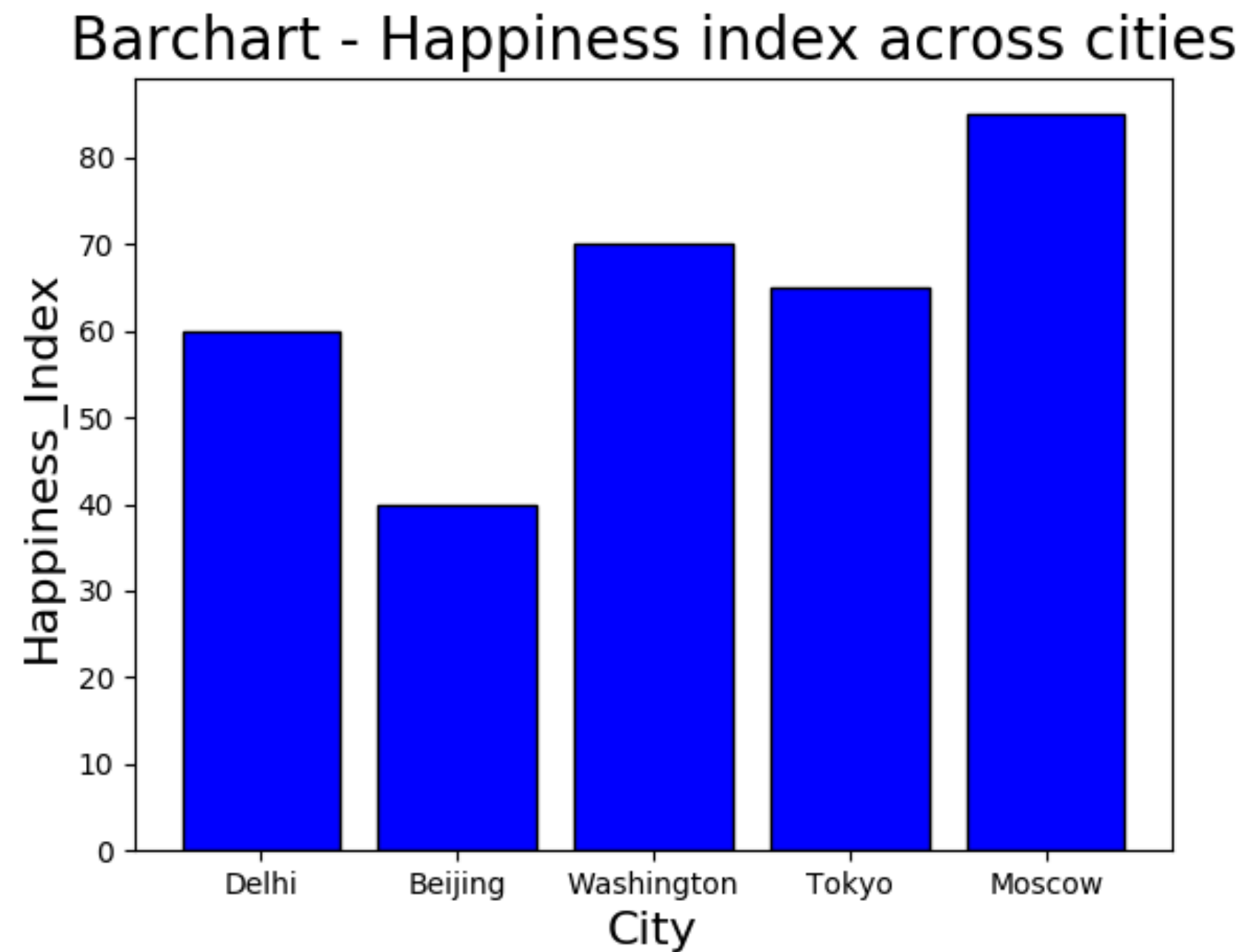




# Common visualizations for low dimensional data (1D/2D)

**Bivariate** - 2 variables.

**bar chart** - Continuous x Categorical

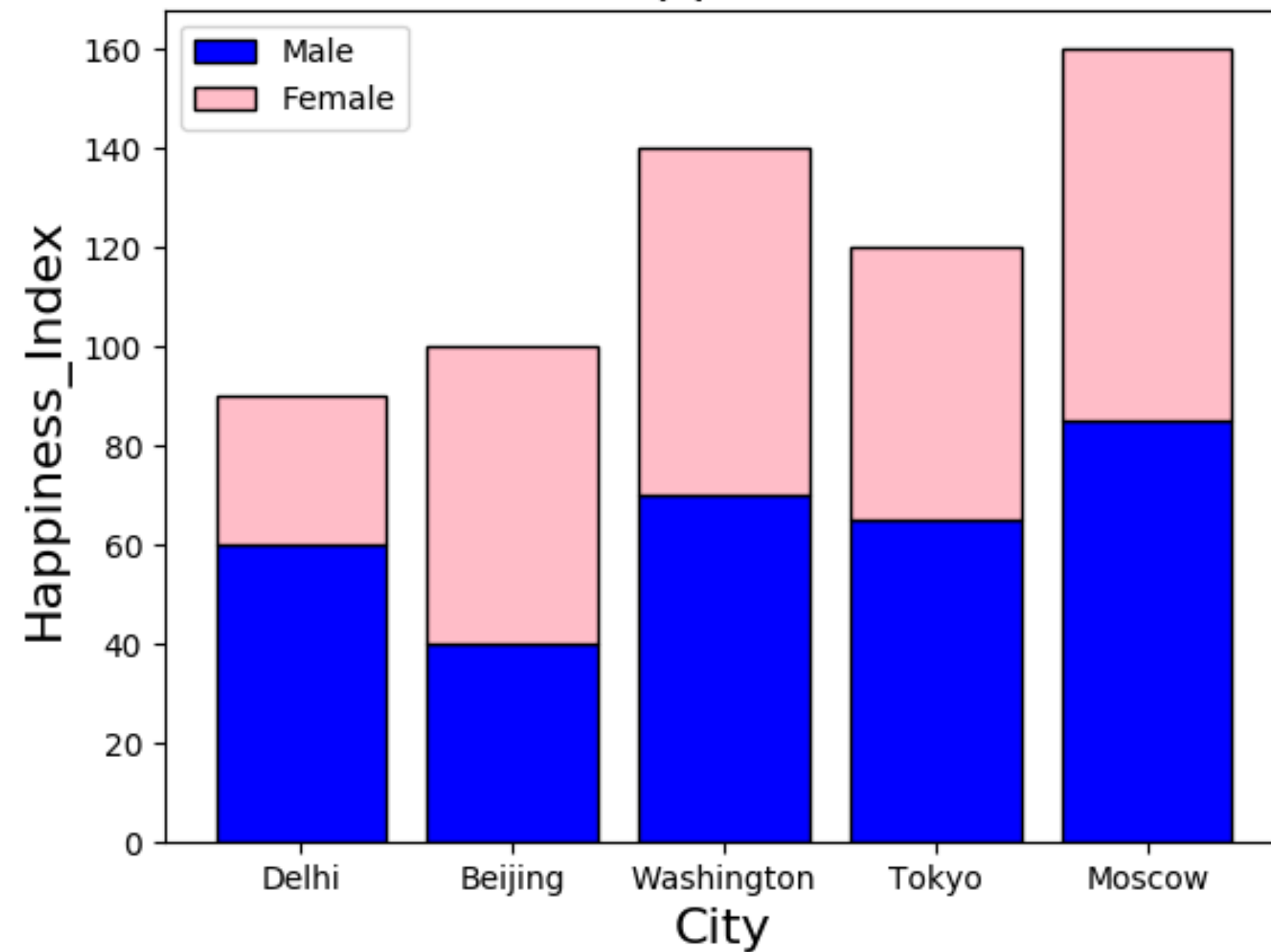


# Common visualizations for low dimensional data (1D/2D)

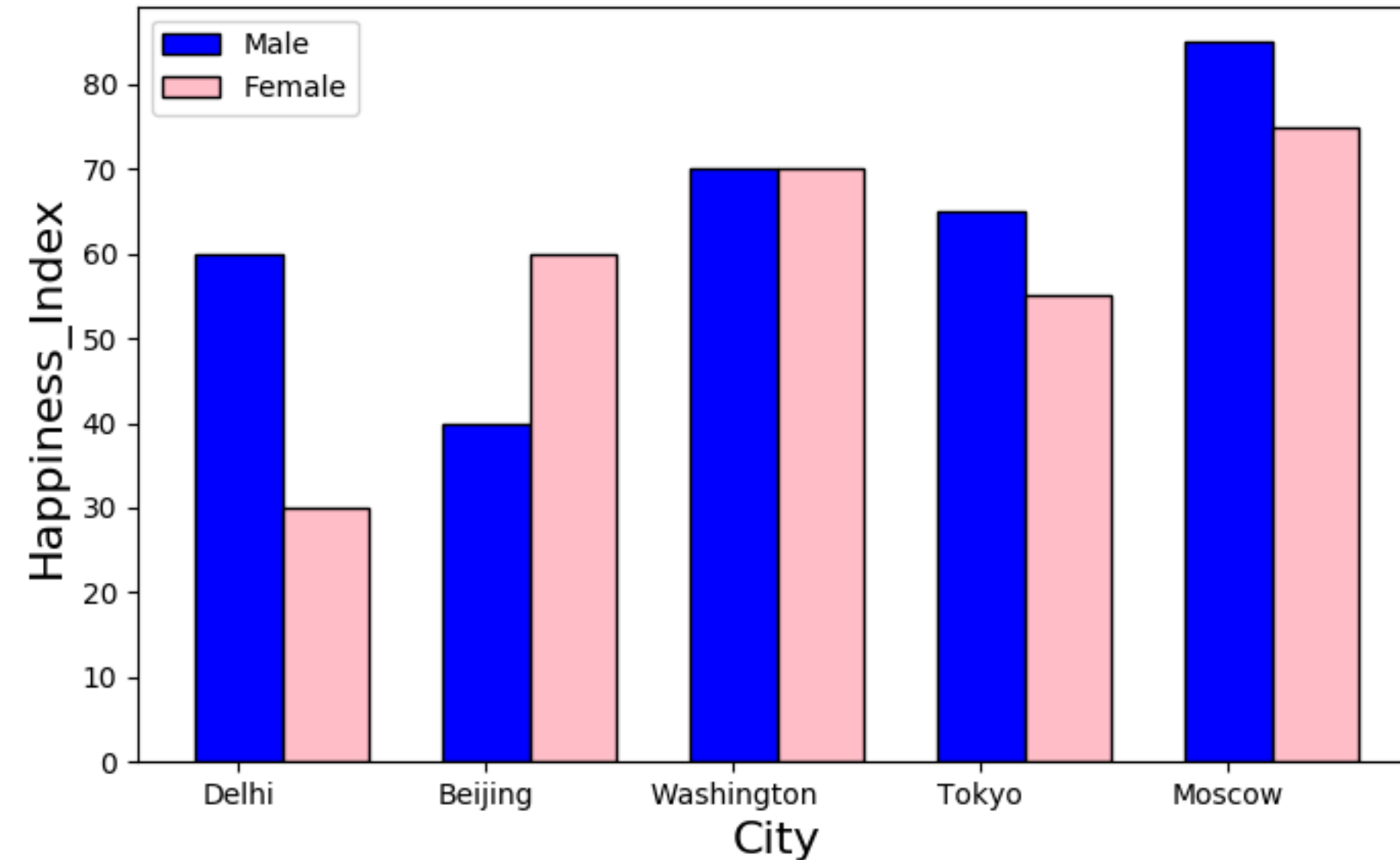
**Bivariate** - 2 variables.

**bar chart** - Continuous x Categorical

Stacked Barchart - Happiness index across cities



Group Barchart - Happiness index across cities By Gender



# Data visualization basics (needed for exploring data)

Human brain is exceptionally good at visual pattern

- Time-series: A single variable is captured over a period of time - **line chart**
- Ranking: Categorical subdivisions are ranked in ascending or descending order - **bar chart**
- Part-to-whole: Categorical subdivisions are measured as a ratio to the whole (i.e., a percentage out of 100%) - **pie chart**
- Deviation: Categorical subdivisions are compared against a reference - **bar chart**
- Frequency distribution: Shows the number of observations of a particular variable for given interval - **histogram**
- Correlation: Comparison between observations represented by two variables (X,Y) to determine if they tend to move in the same or opposite directions - **scatter plot**
- Nominal comparison: Comparing categorical subdivisions in no particular order - **bar chart**
- Geographic or geospatial: Comparison of a variable across a map - **cartogram**



Code to visual - ggplot

# Basic plotting

ggplot basics

ggplot2 is now over 10 years old and is used by hundreds of thousands of people to make millions of plots!

It is a system for creating graphics, based on The Grammar of Graphics.

## Installation

```
The easiest way to get ggplot2 is to install the whole tidyverse:
install.packages("tidyverse")

Alternatively, install just ggplot2:
install.packages("ggplot2")
```

<http://r-statistics.co/ggplot2-Tutorial-With-R.html#1.%20The%20Setup>

# Basic plotting

ggplot basics

ggplot2 is now over 10 years old and is used by hundreds of thousands of people to make millions of plots!

**Setup:** First tell ggplot what data to use.

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```

<http://r-statistics.co/ggplot2-Tutorial-With-R.html#1.%20The%20Setup>

# Basic plotting ggplot basics

## Setup:

- First tell ggplot what data to use.
- Add whatever aesthetics you want to apply to your ggplot (inside **aes()** argument) - such as X and Y axis by specifying the respective variables from the dataset.
- The variable based on which the color, size, shape and stroke should change can also be specified

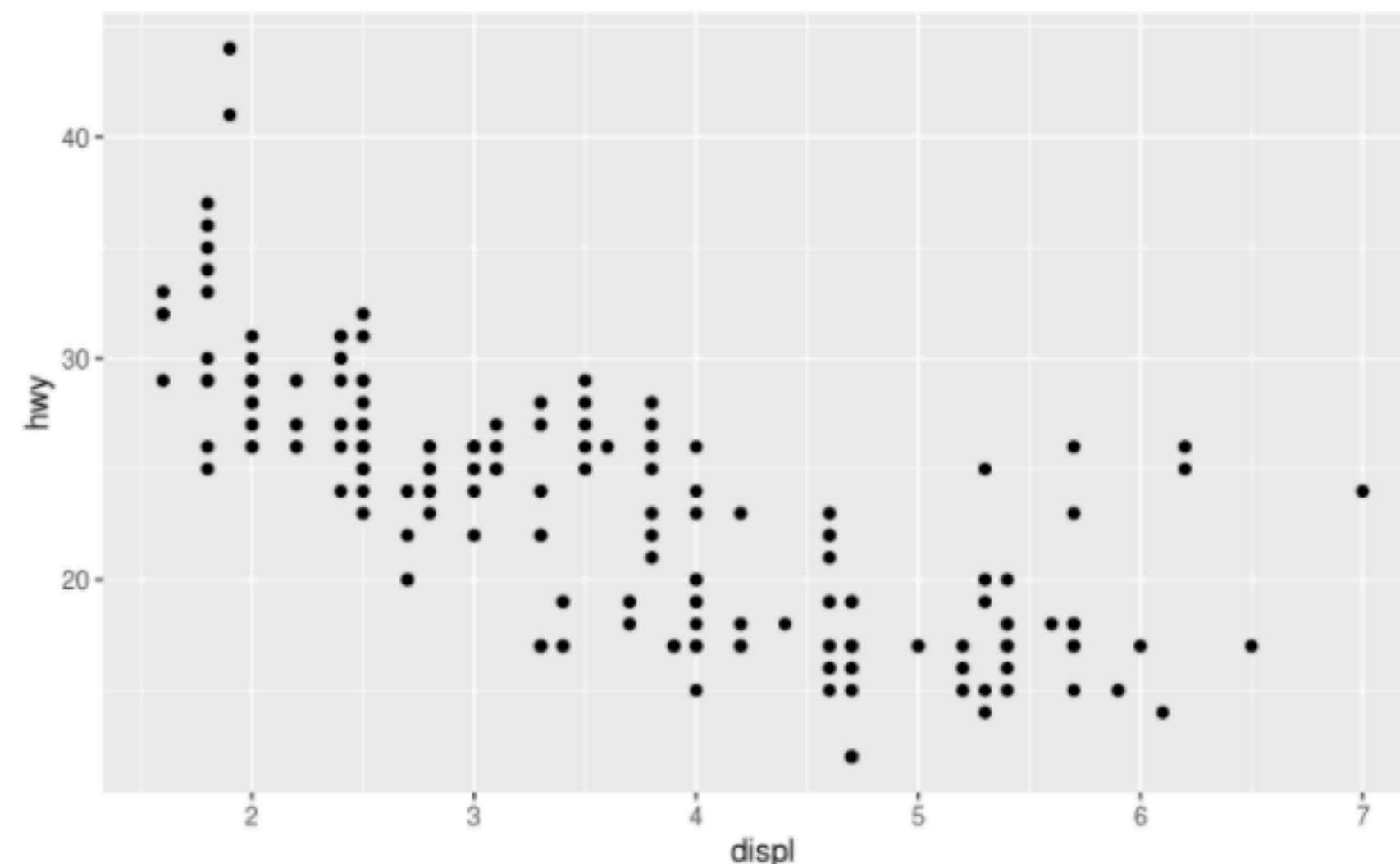
```
ggplot(data = mpg) +
 geom_point(mapping = aes(x = displ, y = hwy))
```

displ: displacement — represents engine size

hwy: highway miles per gallon — represents Fuel efficiency

What does the plot tell you?

Can you figure out how X and Y are related?





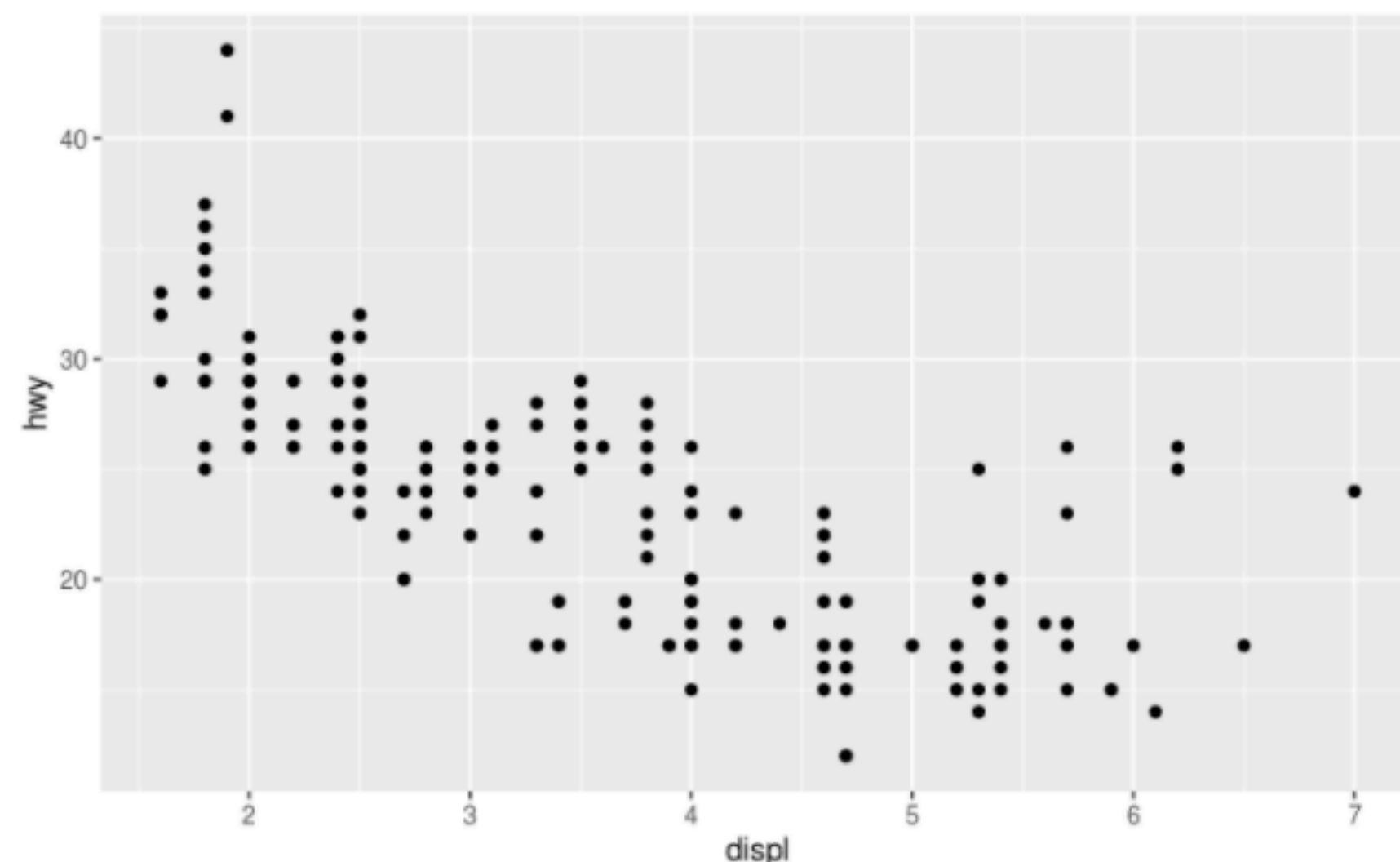
# A graphing template

R for Data Science: <https://r4ds.had.co.nz/data-visualisation.html>

```
ggplot(data = <DATA>) +
 <GEOM_FUNCTION>(mapping = aes(<MAPPINGS>))
```

Understanding the **<MAPPINGS>** component.

```
ggplot(data = mpg) +
 geom_point(mapping = aes(x = displ, y = hwy))
```



A **geom** is the geometrical object that a plot uses to represent data. Here we are using **point** geom. (bar charts use **bar** geoms, line charts use **line** geoms, **boxplots** use boxplot geoms)

Each geom function takes a **mapping** argument. This defines how variables in your dataset are mapped to visual properties.

The **mapping** argument is always paired with aes ( ), and the **x** and **y** arguments of aes ( ) specify which variables to map to the x and y axes.

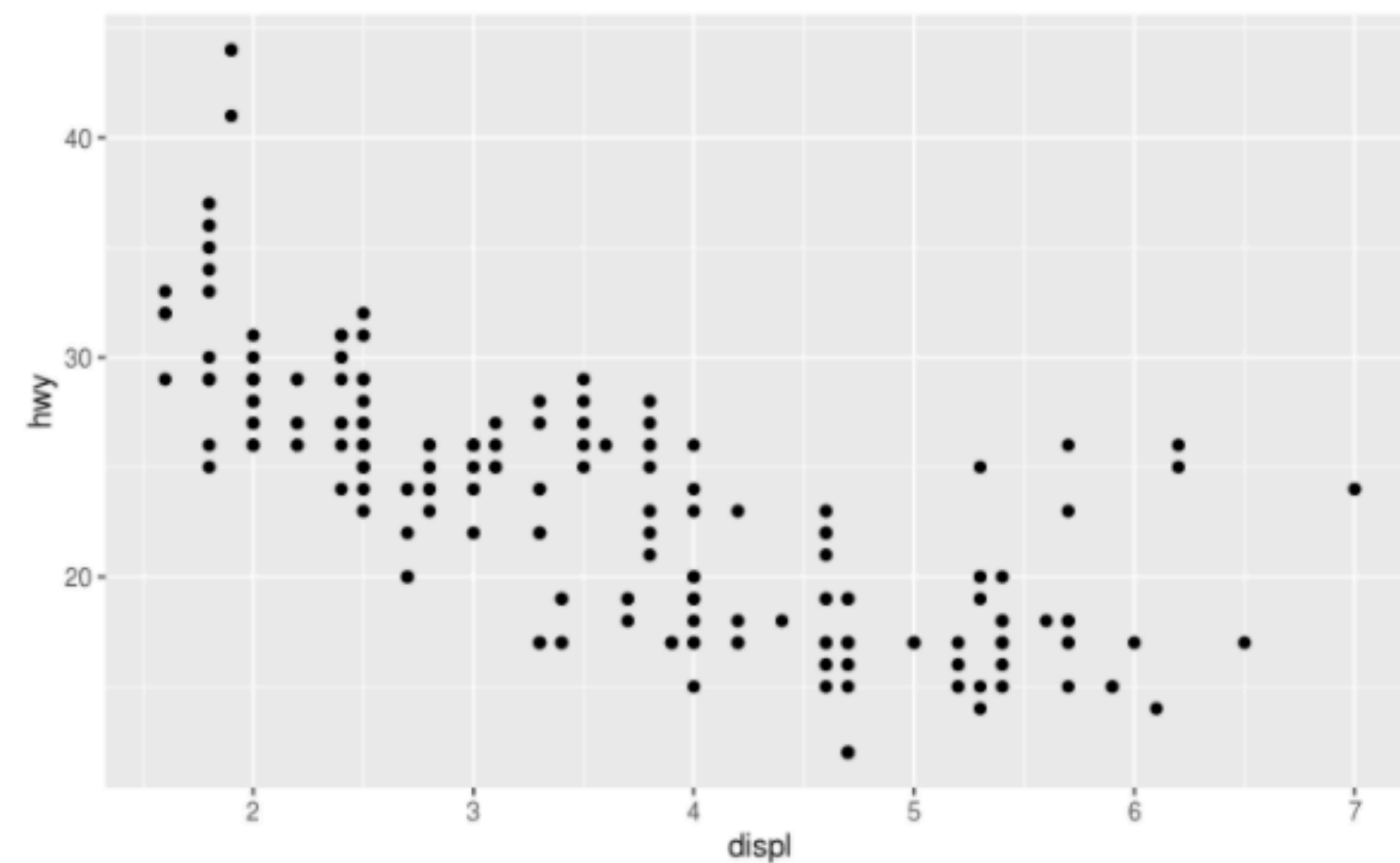
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R for Data Science: <https://r4ds.had.co.nz/data-visualisation.html>

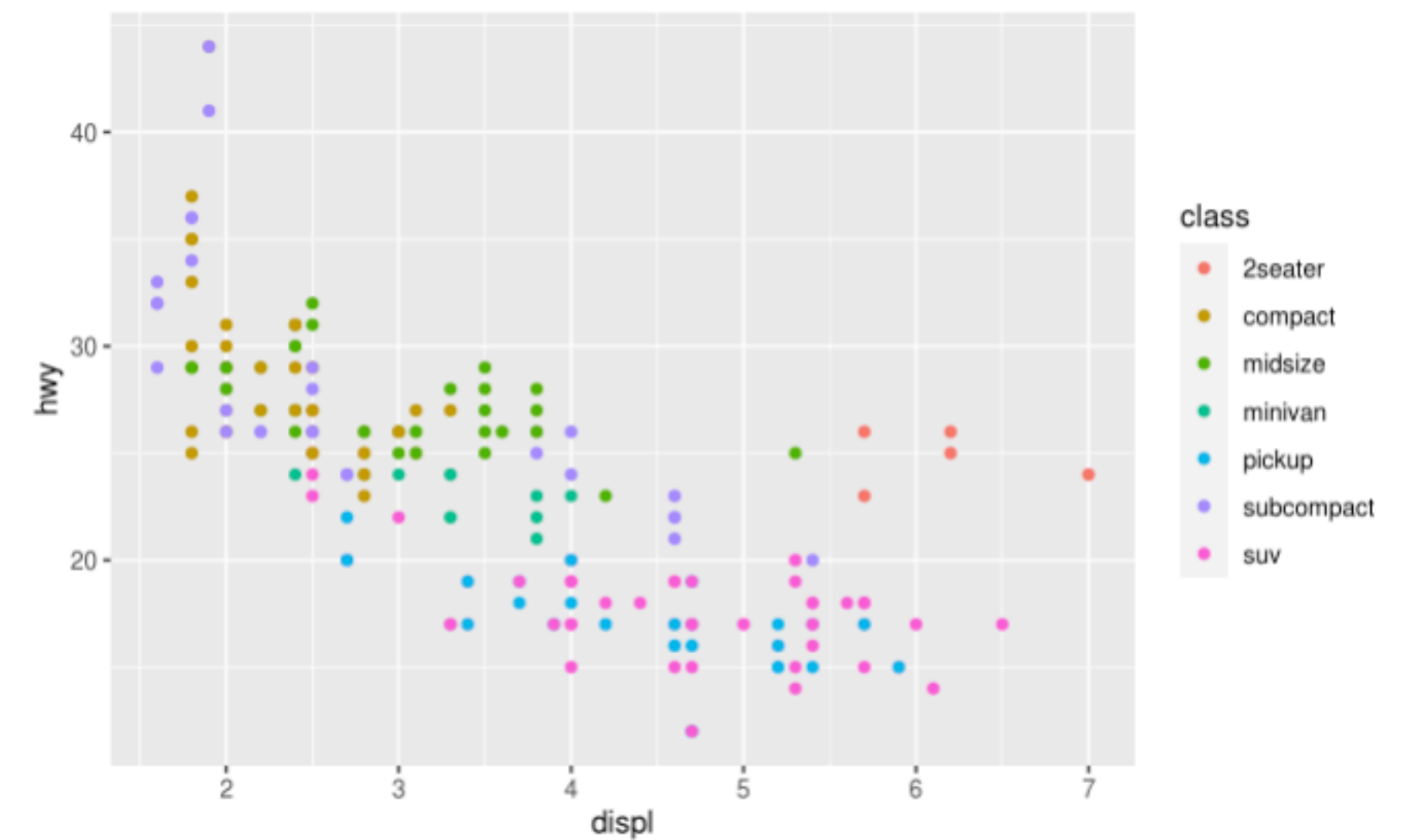
```
ggplot(data = <DATA>) +
 <GEOM_FUNCTION>(mapping = aes(<MAPPINGS>))
```

Understanding the **<MAPPINGS>** component.

```
ggplot(data = mpg) +
 geom_point(mapping = aes(x = displ, y = hwy))
```



```
ggplot(data = mpg) +
 geom_point(mapping = aes(x = displ, y = hwy, color = class))
```



# A graphing template

```
ggplot(data = <DATA>) +
 <GEOM_FUNCTION>(mapping = aes(<MAPPINGS>))
```

Understanding the **<MAPPINGS>** component.

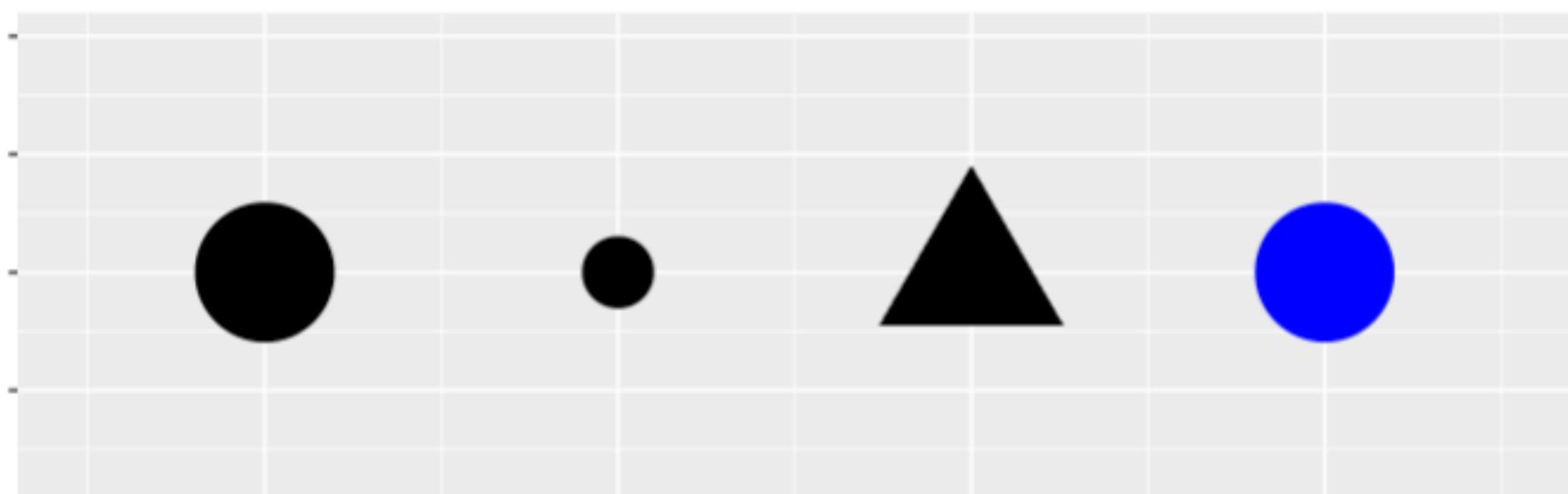
## Aesthetic mappings

A visual property of the objects in your plot.

Aesthetics include:

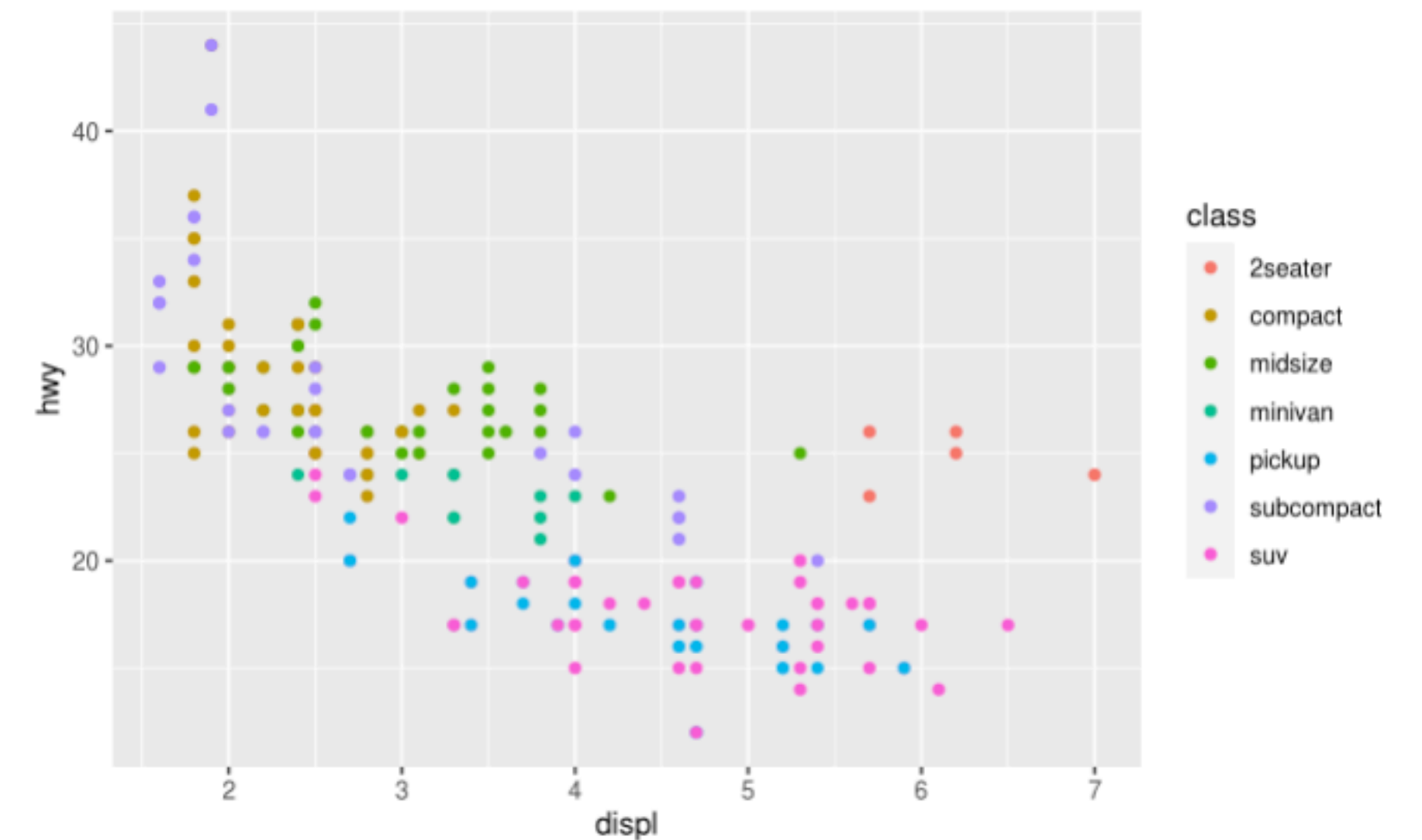
- size,
- shape, or
- color of your points.

Display a point in different ways by changing the values of its aesthetic properties.



## Mapping a variable to the **color** aesthetic

```
ggplot(data = mpg) +
 geom_point(mapping = aes(x = displ, y = hwy, color = class))
```



# A graphing template

```
ggplot(data = <DATA>) +
 <GEOM_FUNCTION>(mapping = aes(<MAPPINGS>))
```

Understanding the **<MAPPINGS>** component.

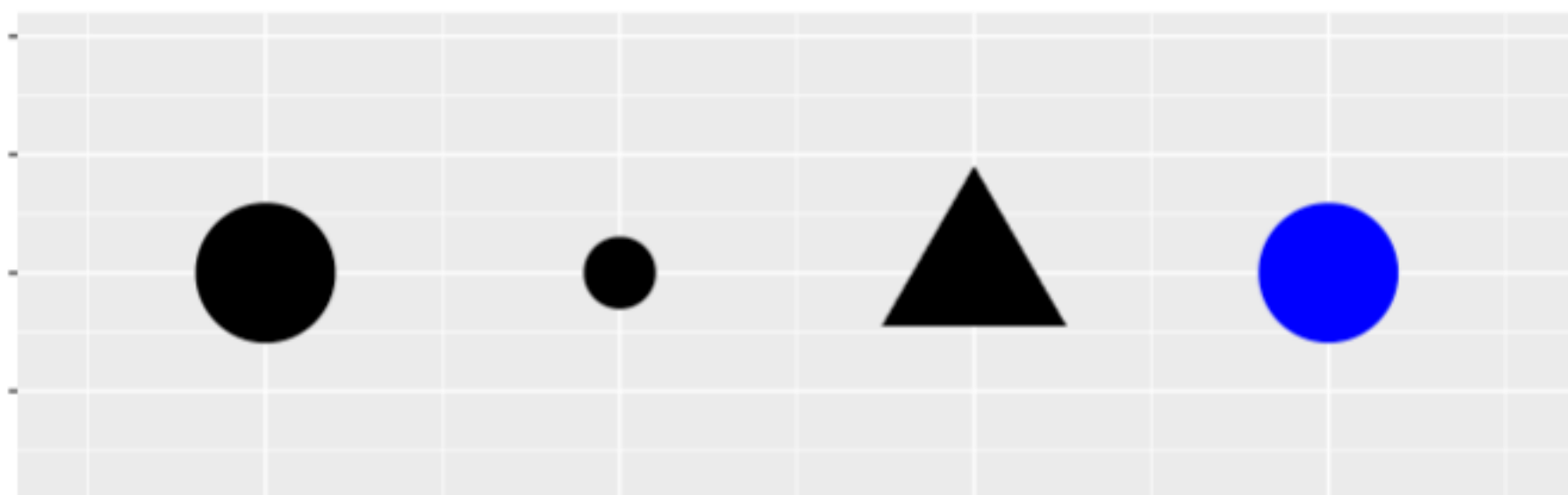
## Aesthetic mappings

A visual property of the objects in your plot.

Aesthetics include:

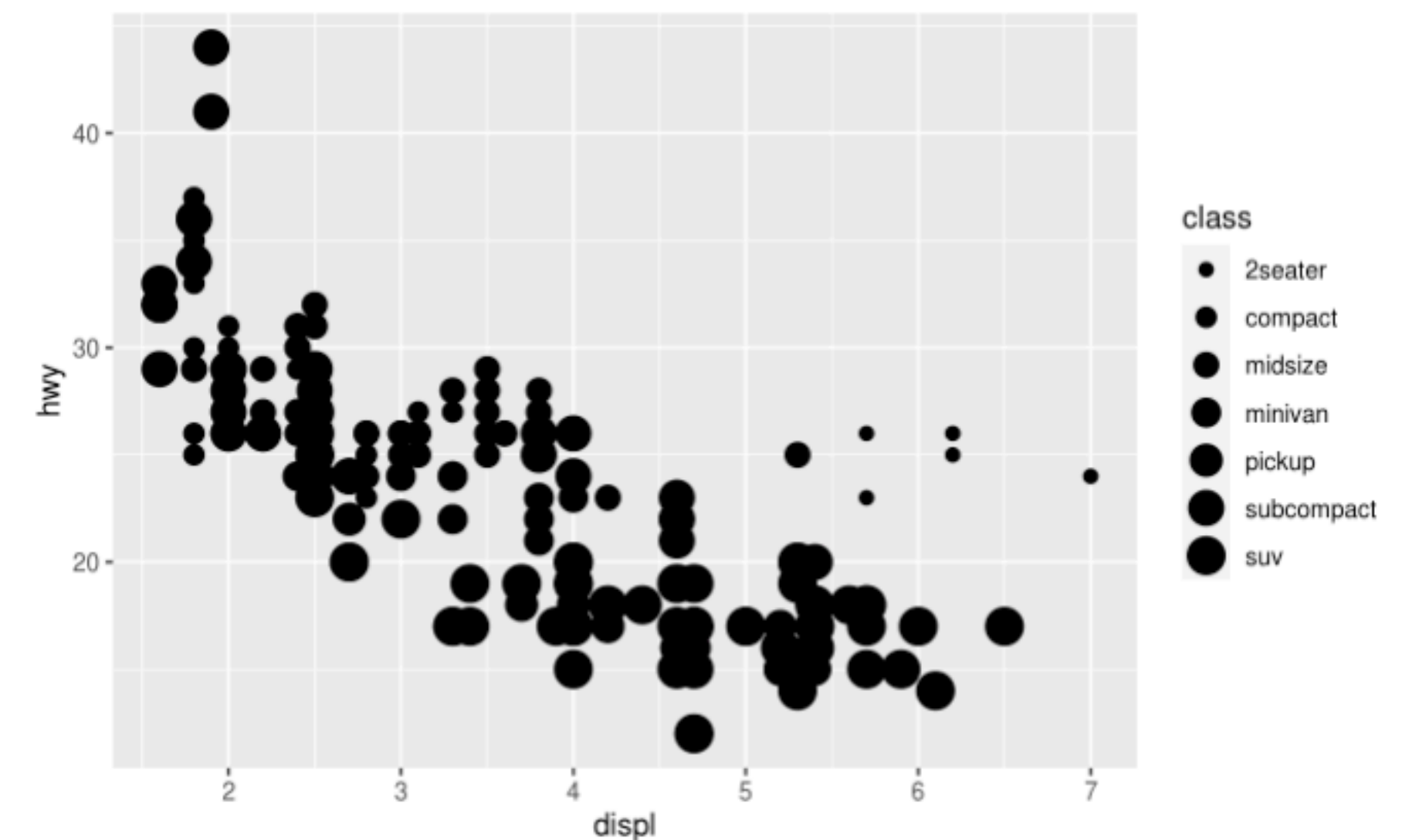
- size,
- shape, or
- color of your points.

Display a point in different ways by changing the values of its aesthetic properties.



## Mapping a variable to the **size** aesthetic

```
ggplot(data = mpg) +
 geom_point(mapping = aes(x = displ, y = hwy, size = class))
#> Warning: Using size for a discrete variable is not advised.
```





# A graphing template

```
ggplot(data = <DATA>) +
 <GEOM_FUNCTION>(mapping = aes(<MAPPINGS>))
```

Understanding the **<MAPPINGS>** component.

## Aesthetic mappings

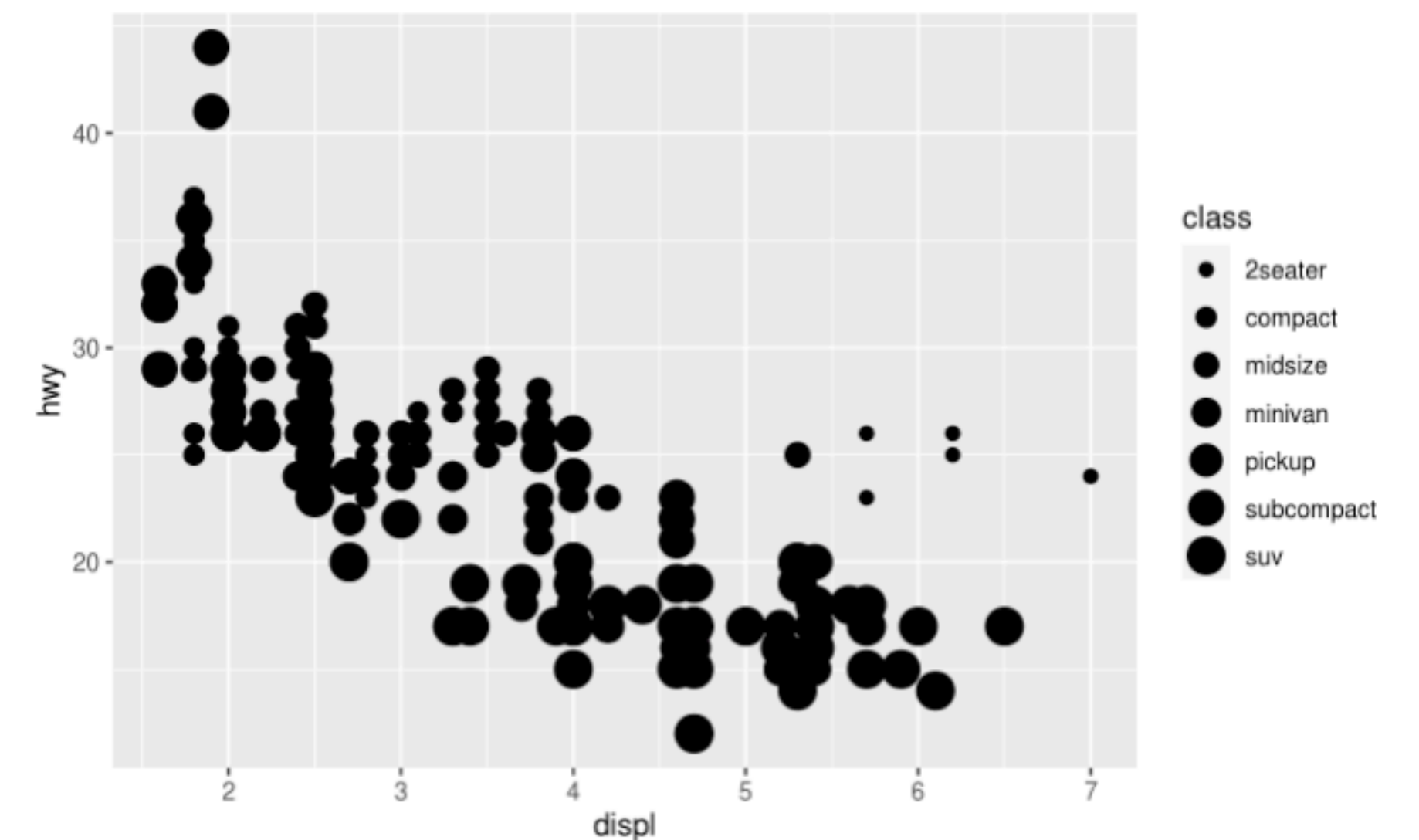
All three give the same result:

```
ggplot(mpg) +
 geom_point(aes(x=displ, y=hwy))
```

```
ggplot(mpg, aes(x=displ, y=hwy)) +
 geom_point()
```

## Mapping a variable to the **size** aesthetic

```
ggplot(data = mpg) +
 geom_point(mapping = aes(x = displ, y = hwy, size = class))
#> Warning: Using size for a discrete variable is not advised.
```



# A graphing template

```
ggplot(data = <DATA>) +
 <GEOM_FUNCTION>(mapping = aes(<MAPPINGS>))
```

Understanding the **<MAPPINGS>** component.

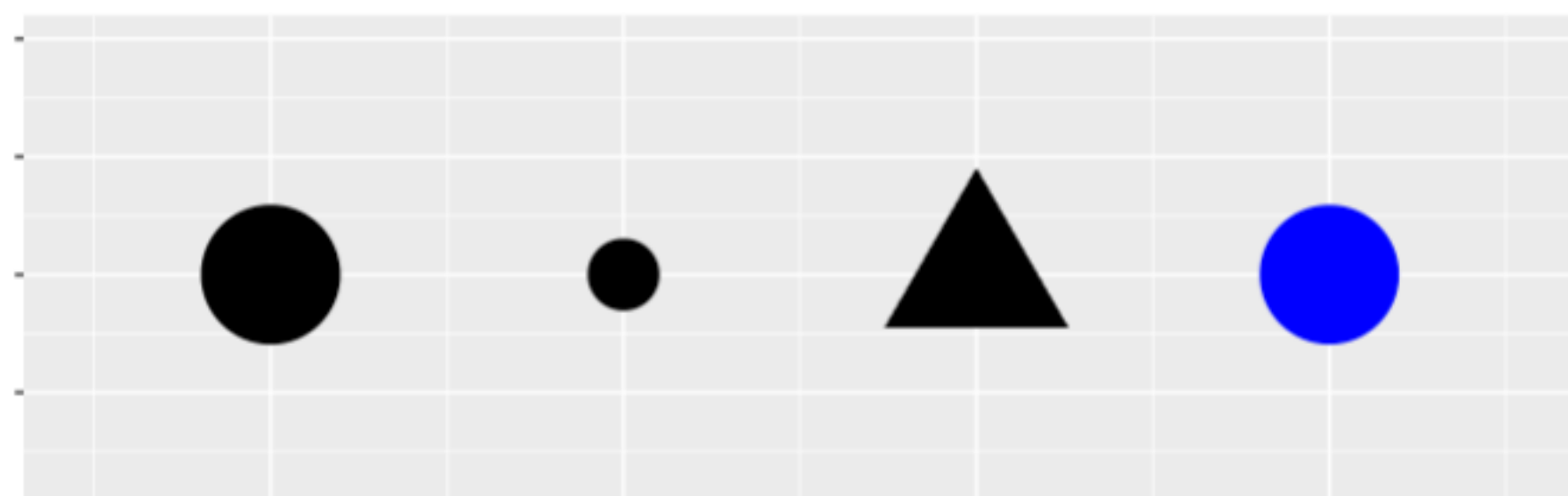
## Aesthetic mappings

A visual property of the objects in your plot.

Aesthetics include:

- size,
- shape, or
- color of your points.

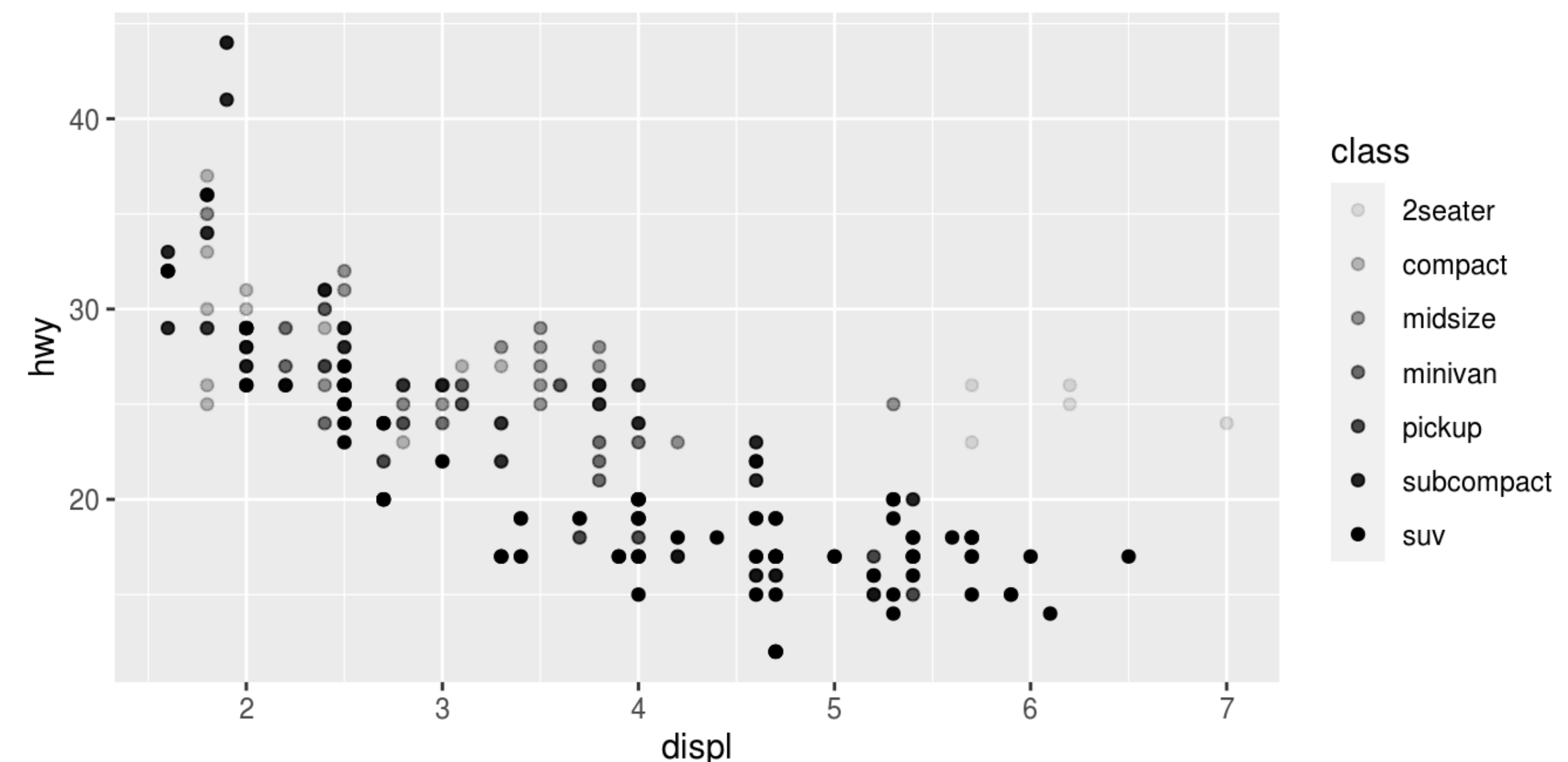
Display a point in different ways by changing the values of its aesthetic properties.



## Mapping a variable to the **alpha** aesthetic

*Controls the transparency of the points*

```
ggplot(data = mpg) +
 geom_point(mapping = aes(x = displ, y = hwy, alpha = class))
```



# A graphing template

```
ggplot(data = <DATA>) +
 <GEOM_FUNCTION>(mapping = aes(<MAPPINGS>))
```

Understanding the **<MAPPINGS>** component.

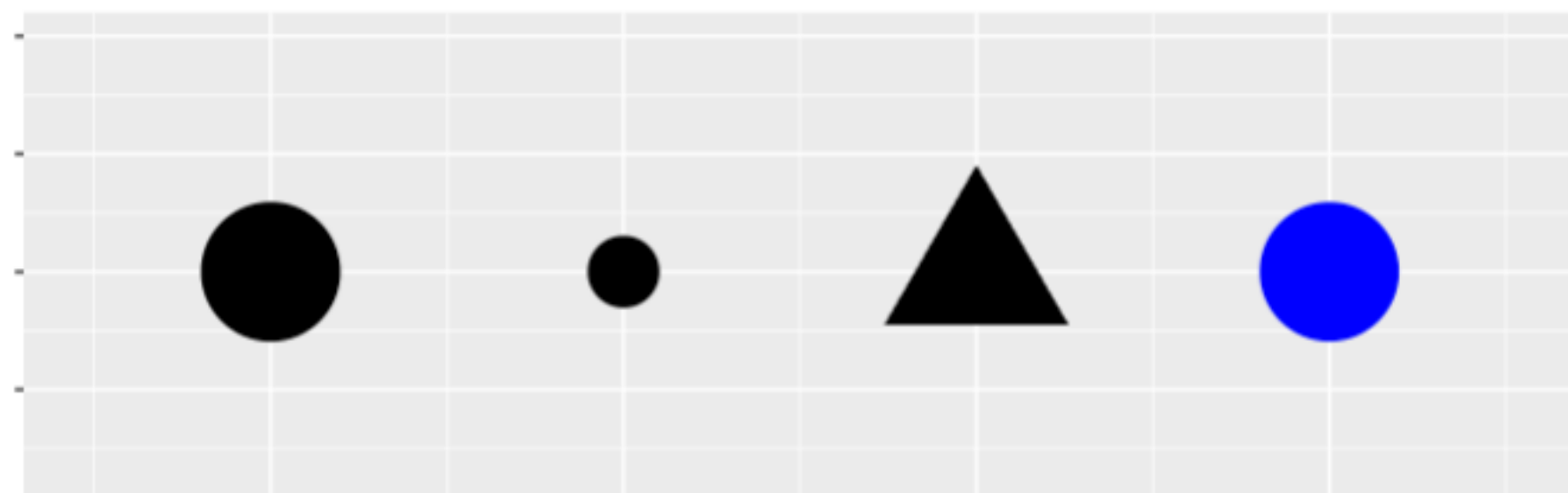
## Aesthetic mappings

A visual property of the objects in your plot.

Aesthetics include:

- size,
- shape, or
- color of your points.

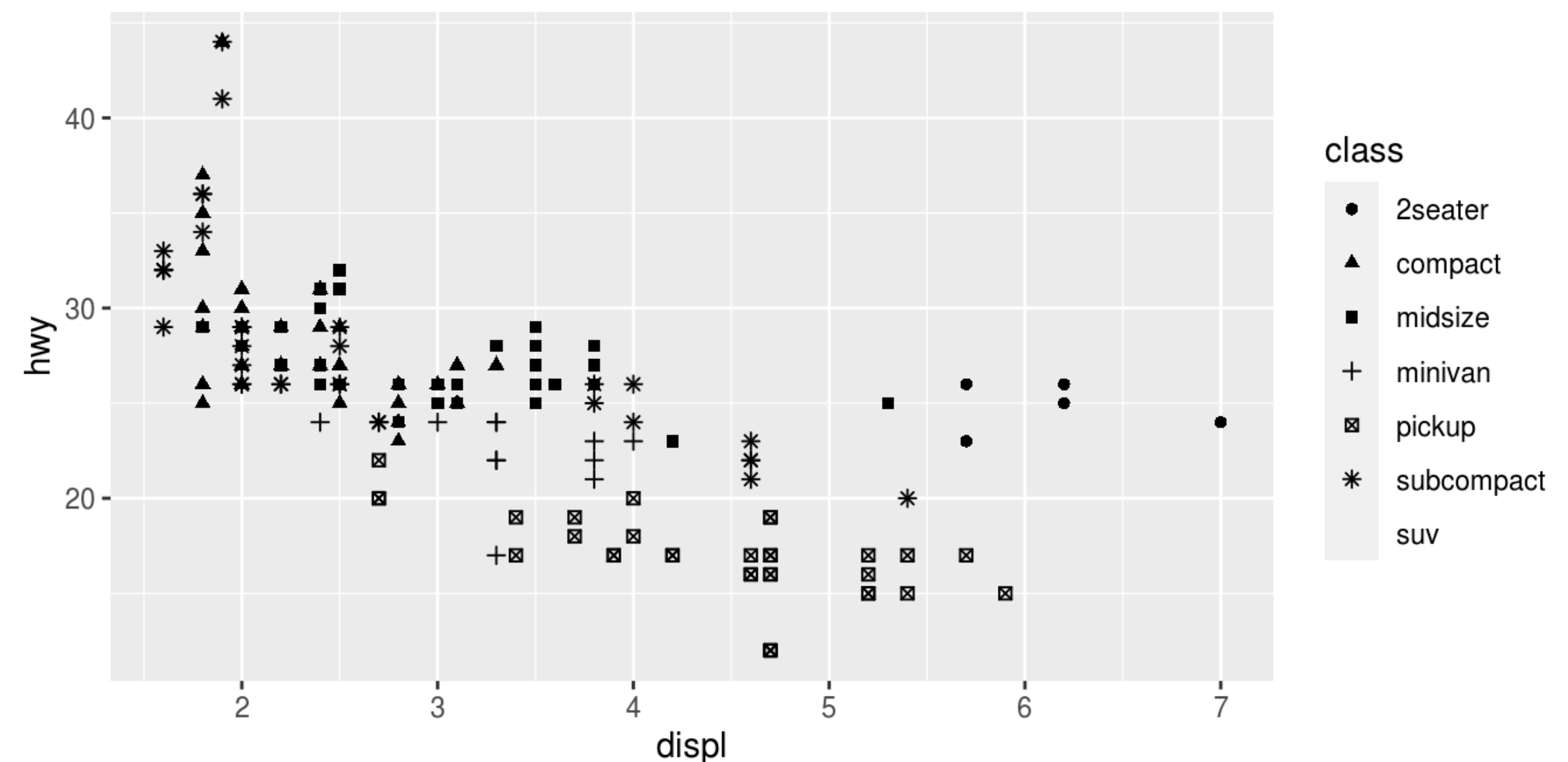
Display a point in different ways by changing the values of its aesthetic properties.



## Mapping a variable to the **shape** aesthetic

*Controls the transparency of the points*

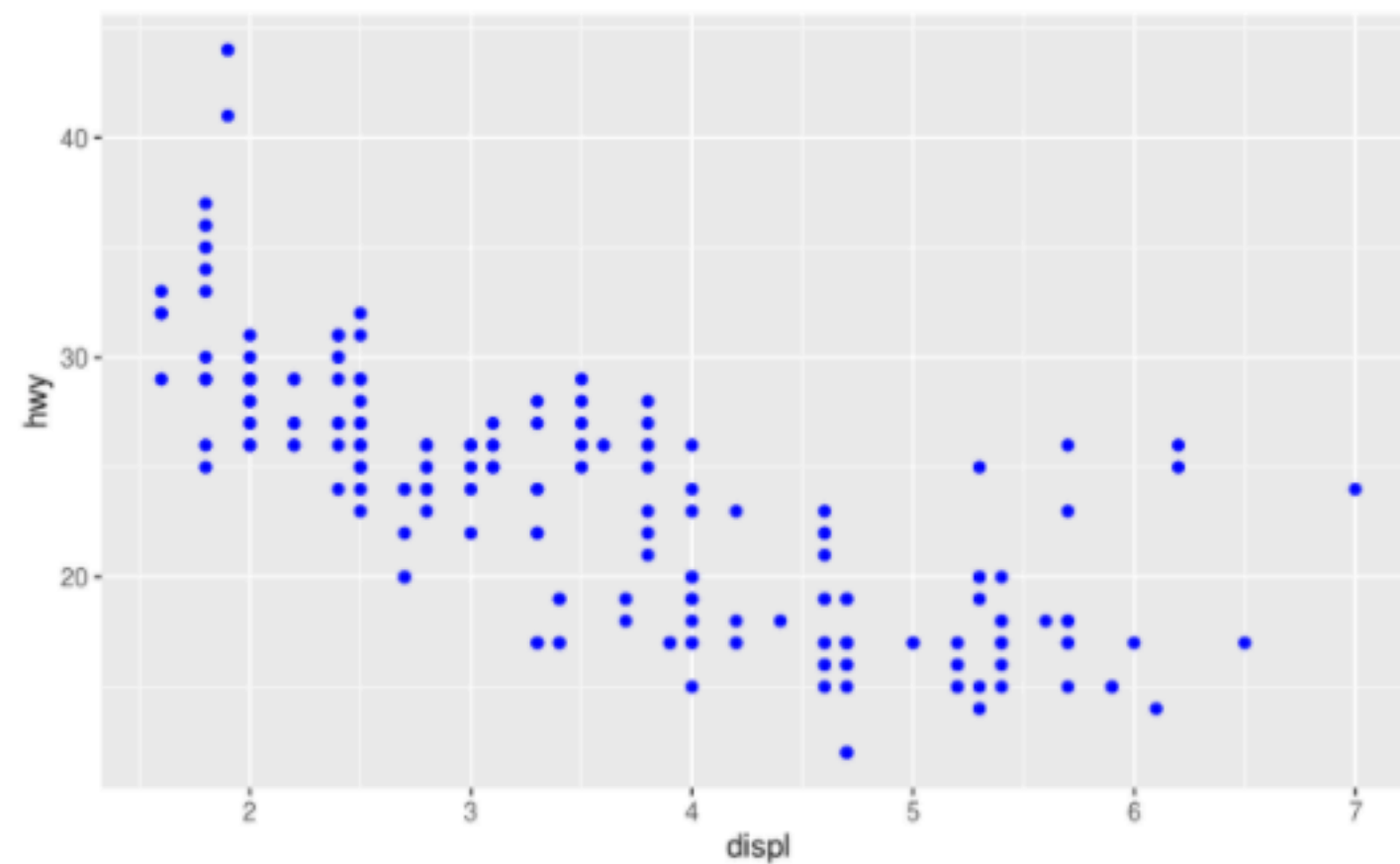
```
ggplot(data = mpg) +
 geom_point(mapping = aes(x = displ, y = hwy, shape = class))
```



# Setting aesthetic manually

To set an aesthetic manually, set the aesthetic by name as an argument of your geom function; i.e. it goes *outside* of `aes()`.

```
ggplot(data = mpg) +
 geom_point(mapping = aes(x = displ, y = hwy), color = "blue")
```





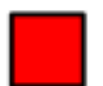







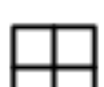














Here color doesn't convey information about a variable, but only changes the appearance of the plot.



# Setting aesthetic manually

To set an aesthetic manually, set the aesthetic by name as an argument of your geom function; i.e. it goes *outside* of `aes()`.

|                                                                                       |                                                                                       |                                                                                        |                                                                                         |                                                                                          |
|---------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------|
|  0   |  4   |  10   |  15   |  22   |
|  1   |  6   |  11   |  16   |  21   |
|  2 |  7 |  12 |  17 |  24 |
|  5 |  8 |  13 |  18 |  23 |
|  3 |  9 |  14 |  19 |  20 |

R has 25 built in shapes that are identified by numbers.

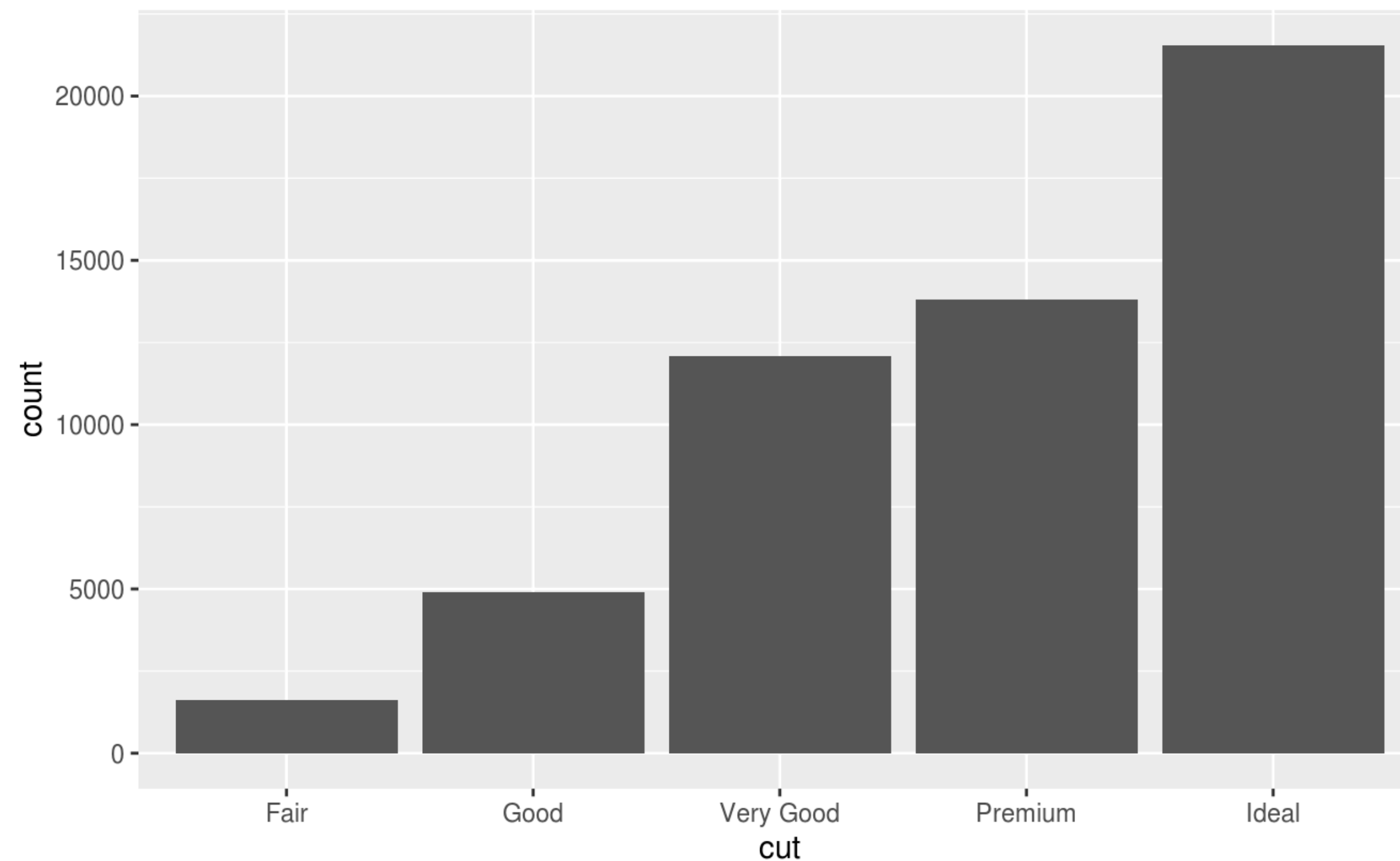
You'll need to pick a level that makes sense for that aesthetic:

- The name of a color as a character string.
- The size of a point in mm.
- The shape of a point as a number, as shown

# Changing the geom in your plot (statistical transformations of the data)

Change the geom function

```
ggplot(data = diamonds) +
 geom_bar(mapping = aes(x = cut))
```



# Statistical transformations

1. **geom_bar()** begins with the **diamonds** data set

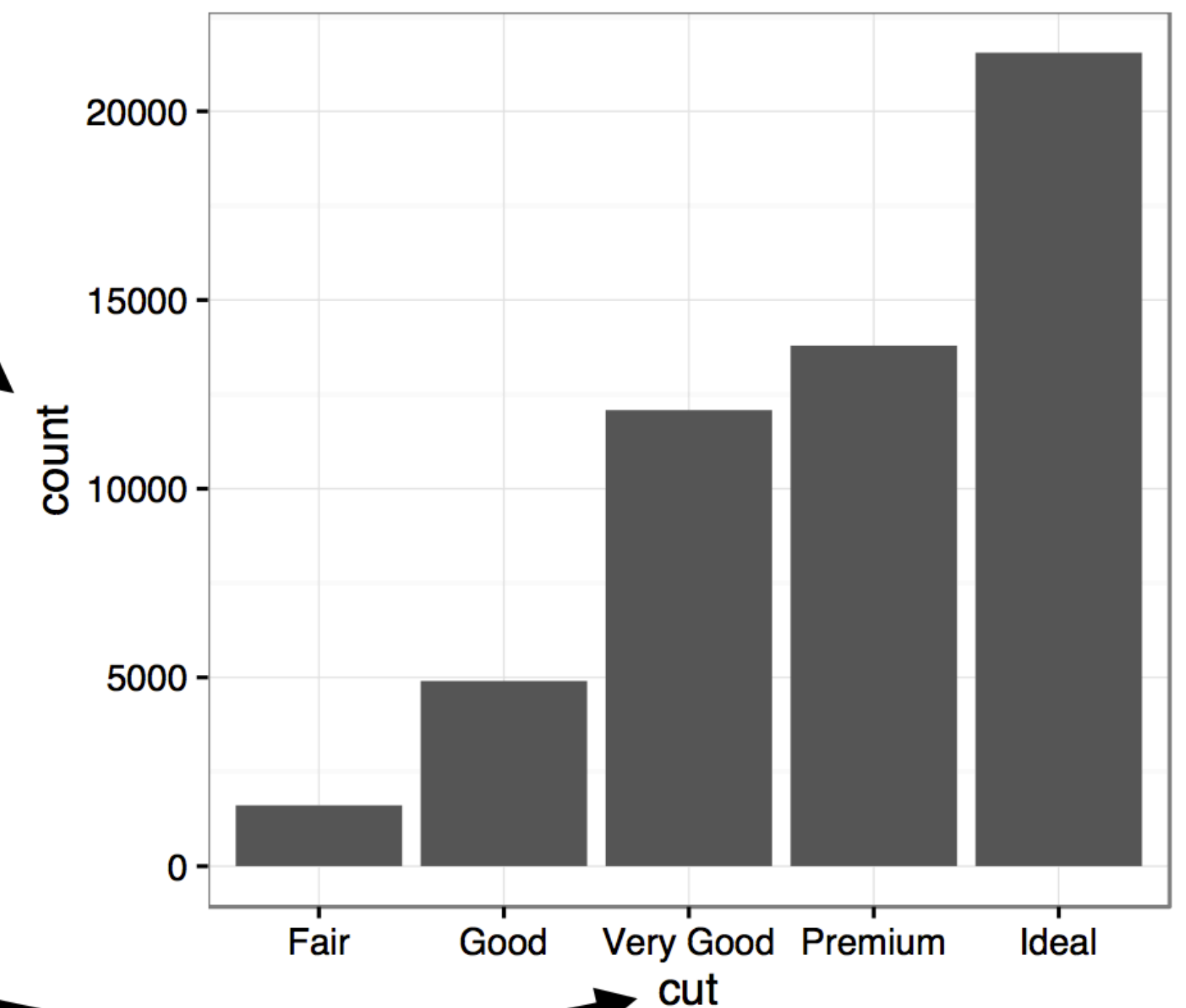
| carat | cut     | color | clarity | depth | table | price | x    | y    | z    |
|-------|---------|-------|---------|-------|-------|-------|------|------|------|
| 0.23  | Ideal   | E     | SI2     | 61.5  | 55    | 326   | 3.95 | 3.98 | 2.43 |
| 0.21  | Premium | E     | SI1     | 59.8  | 61    | 326   | 3.89 | 3.84 | 2.31 |
| 0.23  | Good    | E     | VS1     | 56.9  | 65    | 327   | 4.05 | 4.07 | 2.31 |
| 0.29  | Premium | I     | VS2     | 62.4  | 58    | 334   | 4.20 | 4.23 | 2.63 |
| 0.31  | Good    | J     | SI2     | 63.3  | 58    | 335   | 4.34 | 4.35 | 2.75 |
| ...   | ...     | ...   | ...     | ...   | ...   | ...   | ...  | ...  | ...  |

**stat_count()**

2. **geom_bar()** transforms the data with the "count" stat, which returns a data set of cut values and counts.

| cut       | count | prop |
|-----------|-------|------|
| Fair      | 1610  | 1    |
| Good      | 4906  | 1    |
| Very Good | 12082 | 1    |
| Premium   | 13791 | 1    |
| Ideal     | 21551 | 1    |

3. **geom_bar()** uses the transformed data to build the plot. cut is mapped to the x axis, count is mapped to the y axis.

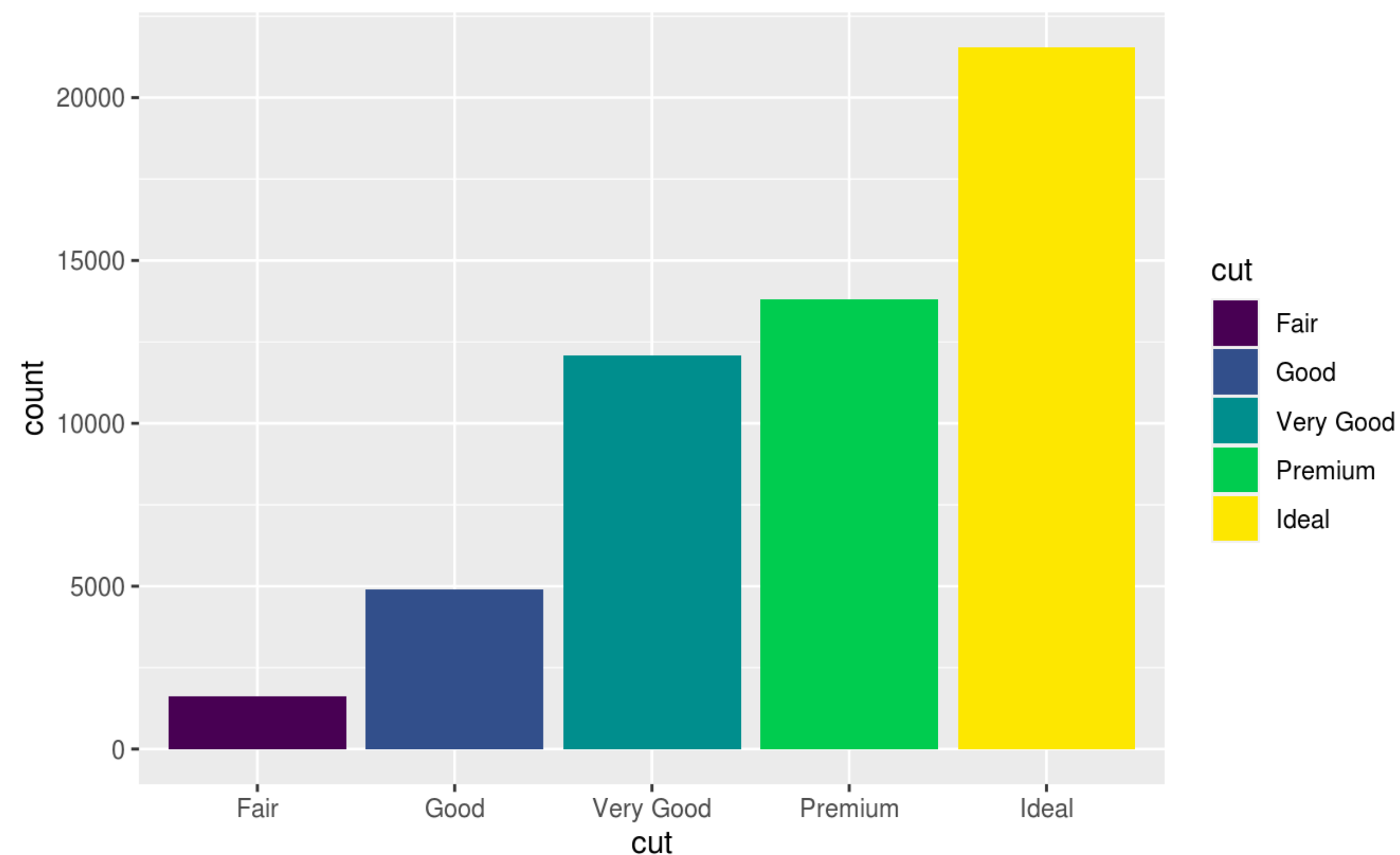


# geom_bar

<https://r4ds.had.co.nz/data-visualisation.html>

You can color a bar chart using either the **color** aesthetic, or, more usefully, **fill**:

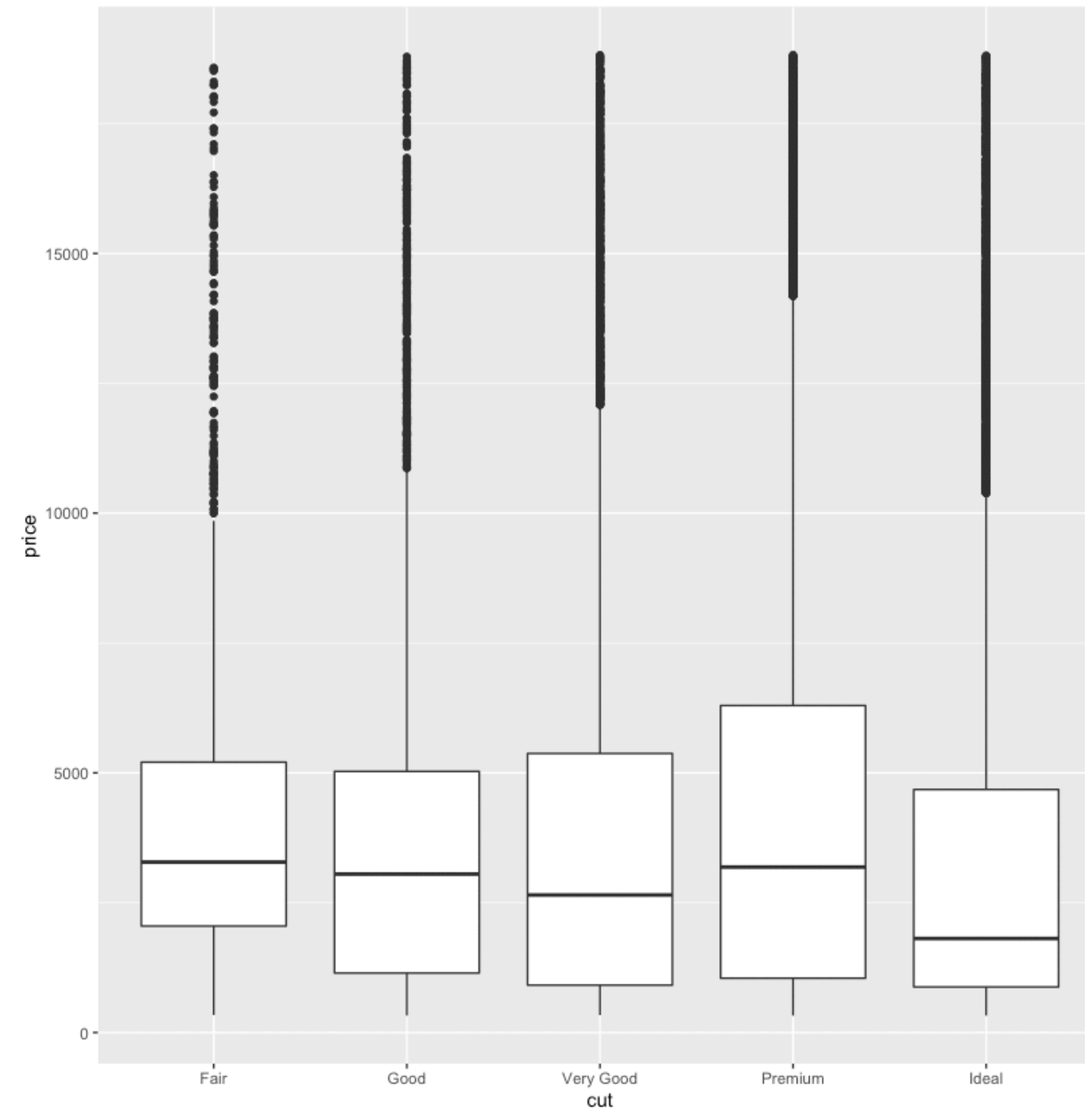
```
ggplot(data = diamonds) +
 geom_bar(mapping = aes(x = cut, fill = cut))
```





# geom_boxplot

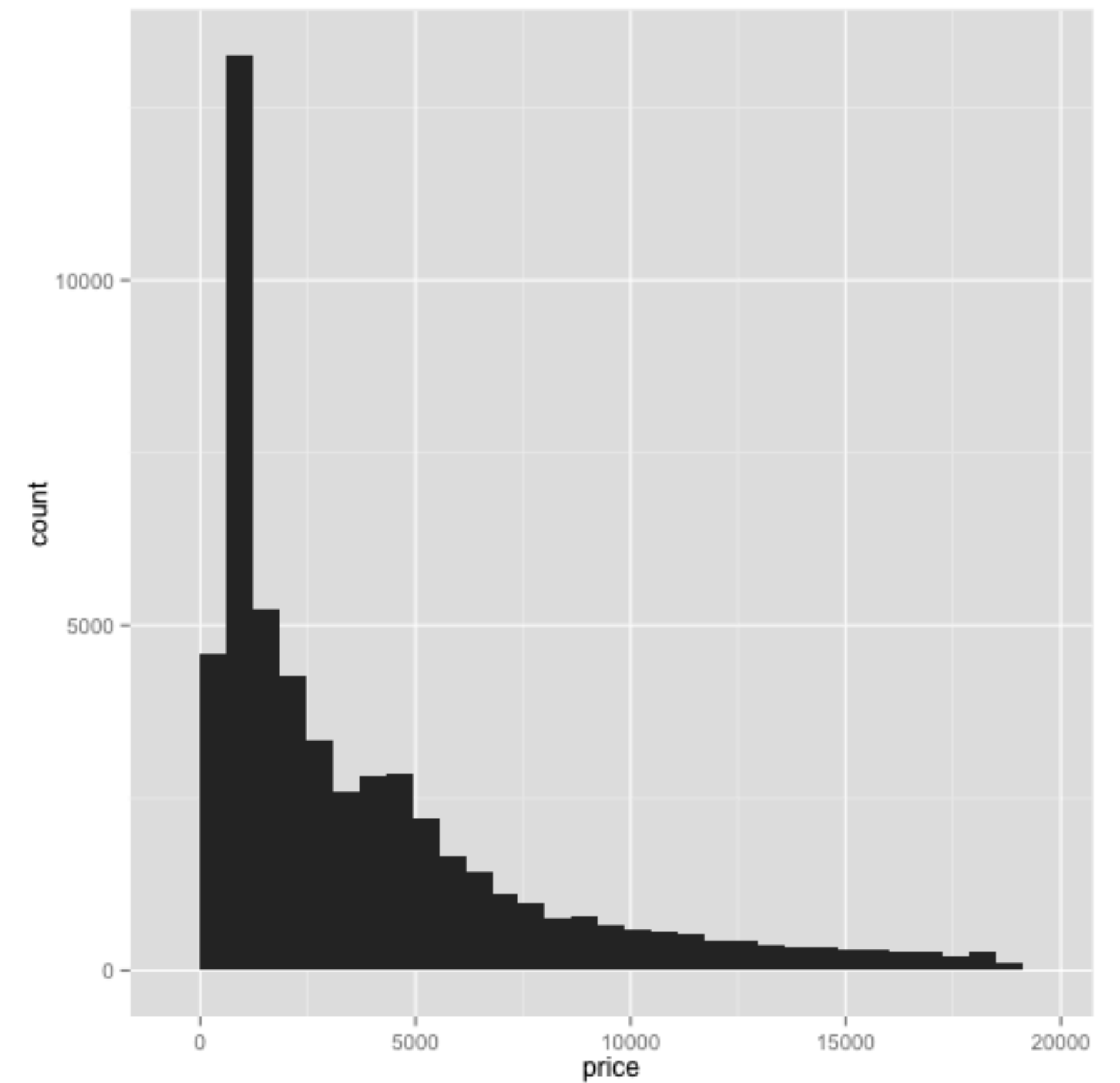
```
ggplot(data = diamonds) +
 geom_boxplot(aes(x=cut, y=price))
```



# geom_histogram

```
ggplot(diamonds, aes(x=price)) + geom_histogram()
```

visualization of the density of the  
distribution of price.

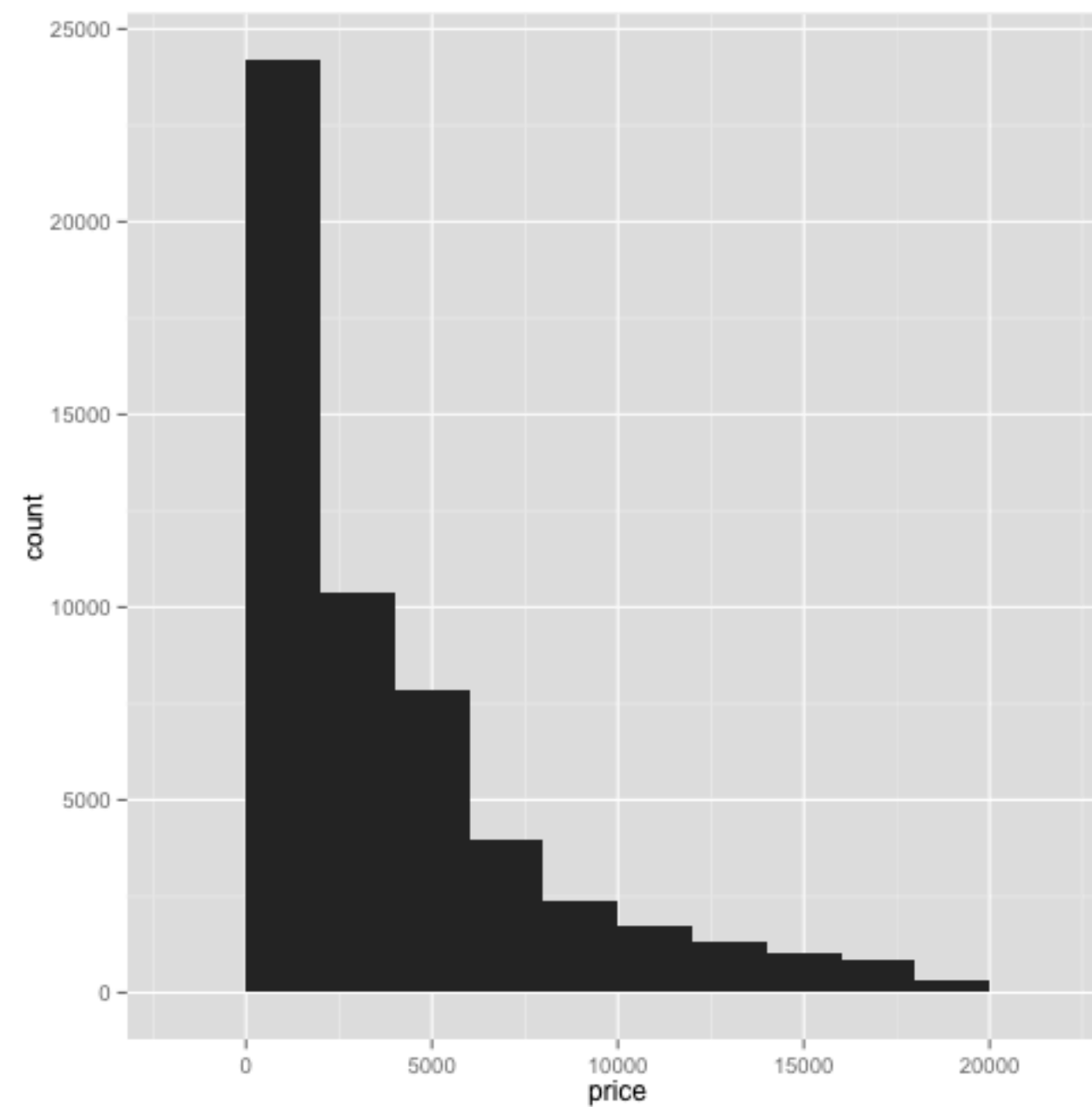


# geom_histogram

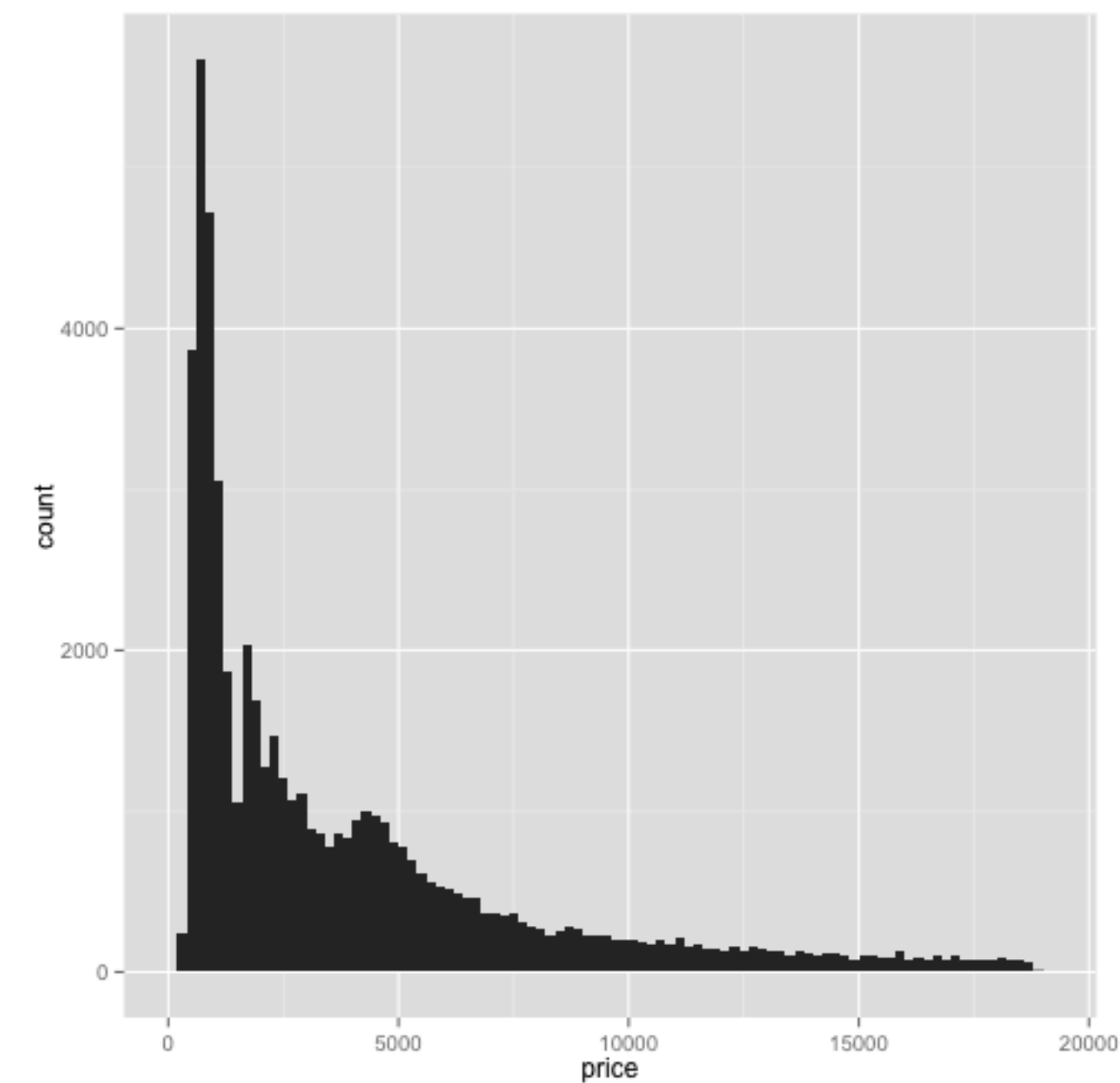
visualization of the density of the distribution of price.

Changing width of each bin to make them wider or thinner

```
ggplot(diamonds, aes(x=price)) +
geom_histogram(binwidth=2000)
```



```
ggplot(diamonds, aes(x=price)) +
geom_histogram(binwidth=200)
```

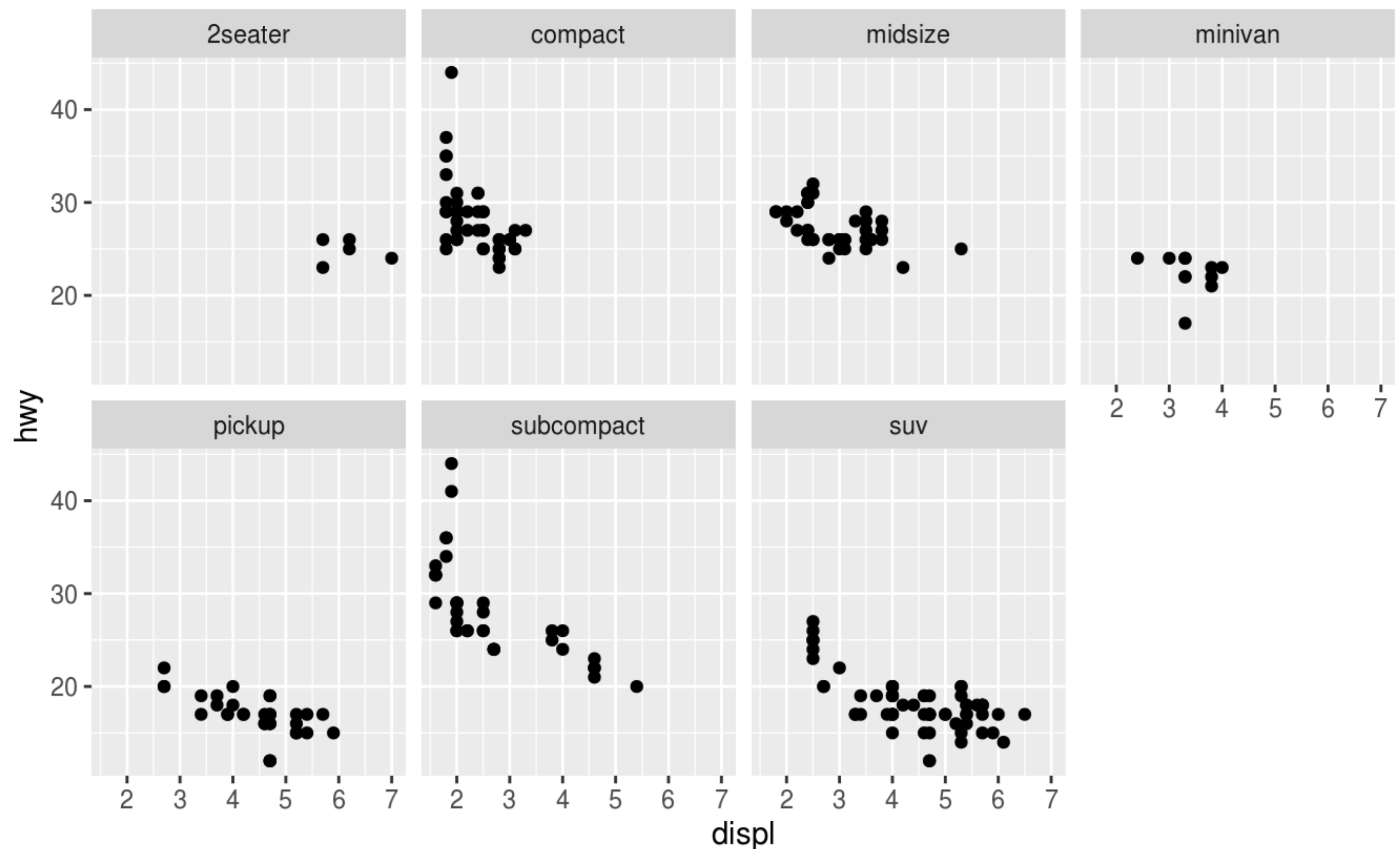


# Facets

subplots that each display one subset of the data

To facet your plot by a single variable, use `facet_wrap()`. The first argument of `facet_wrap()` should be a formula, which you create with `~` followed by a variable name

```
ggplot(data = mpg) +
 geom_point(mapping = aes(x = displ, y = hwy)) +
 facet_wrap(~ class, nrow = 2)
```

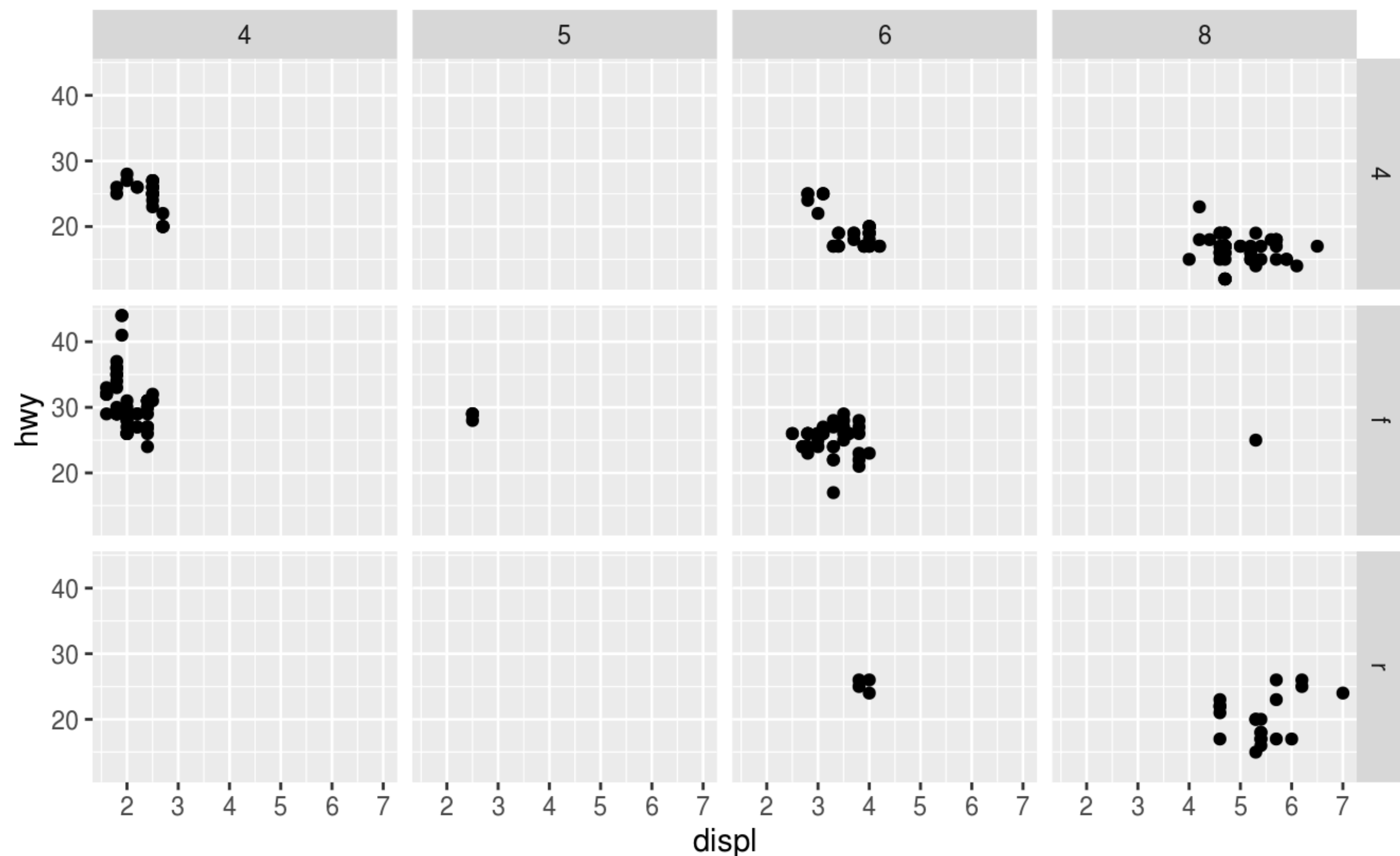


# Facets

subplots that each display one subset of the data

To facet your plot on a combination of 2 variables, use `facet_grid()`. The first argument of `facet_wrap()` is also a formula, which should be 2 variable names separated with `~`

```
ggplot(data = mpg) +
 geom_point(mapping = aes(x = displ, y = hwy)) +
 facet_grid(drv ~ cyl)
```



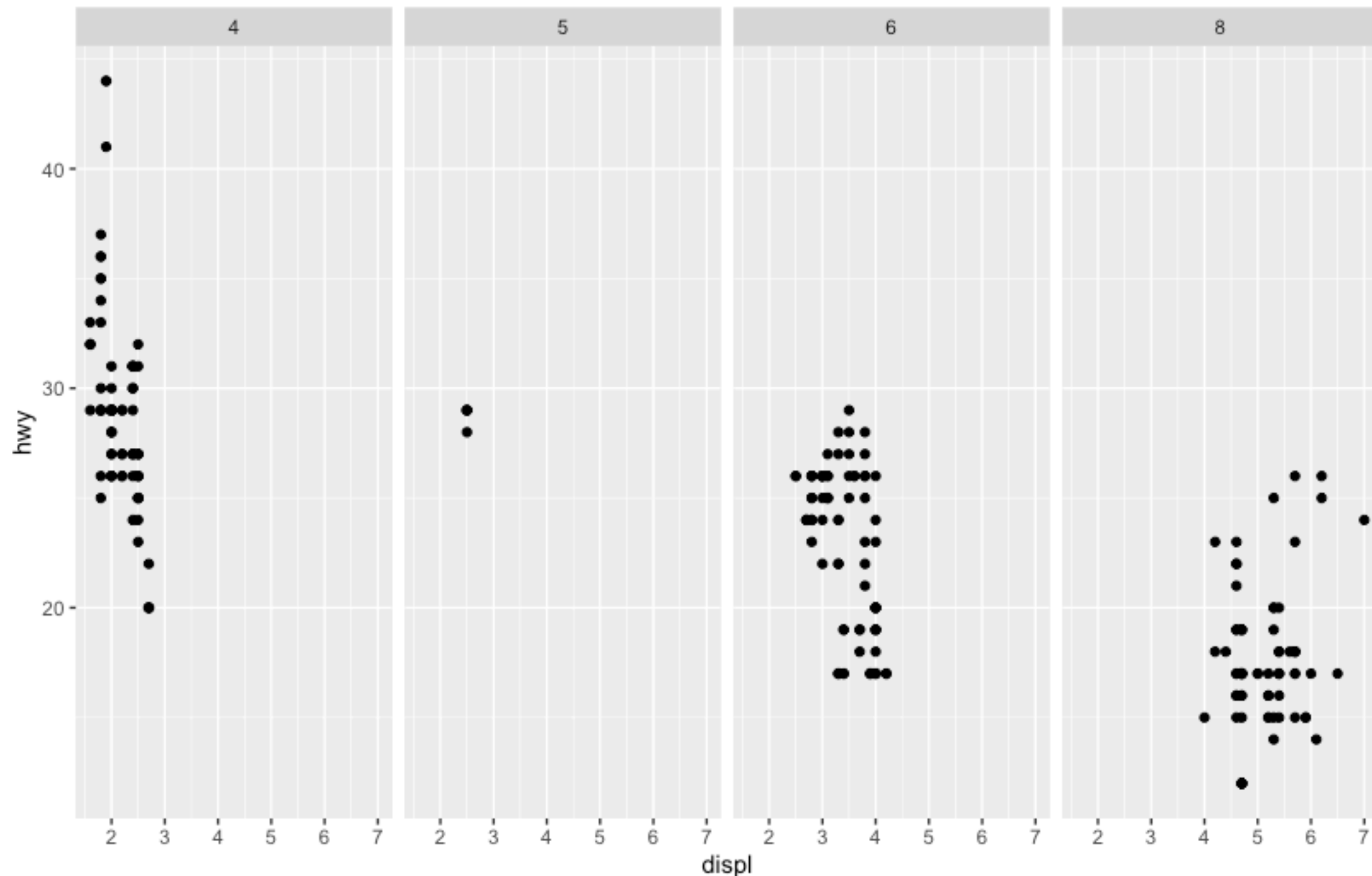


# Facets

subplots that each display one subset of the data

If you prefer to not facet in the rows or columns dimension, use a `.` instead of a variable name, e.g. `facet_grid(. ~ cyl)`.

```
ggplot(data = mpg) +
 geom_point(mapping = aes(x = displ, y = hwy)) +
 facet_grid(. ~ cyl)
```

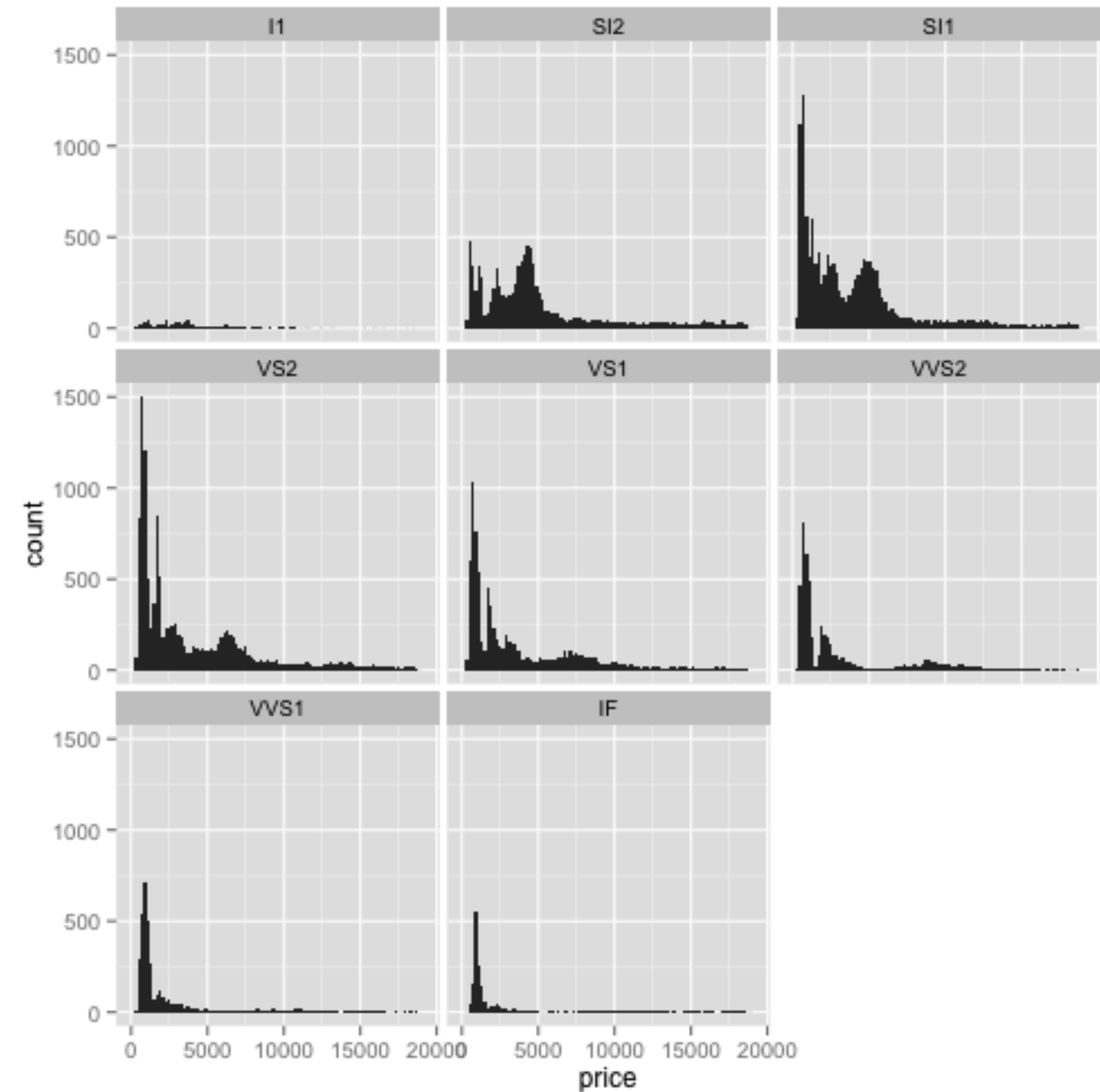


# Facets

subplots that each display one subset of the data

Another example

```
ggplot(diamonds, aes(x=price)) +
 geom_histogram(binwidth=200) +
 facet_wrap(~ clarity)
```



# Common Problems

As you start to run R code, you're likely to run into problems. Don't worry — it happens to everyone.

- Start by carefully comparing the code that you're running to the code in the book.
- R is extremely picky, and a misplaced character can make all the difference.
- Make sure that every ( is matched with a ) and every " is paired with another ".
- Sometimes you'll run the code and nothing happens. Check the left-hand of your console: if it's a +, it means that R doesn't think you've typed a complete expression and it's waiting for you to finish it. In this case, it's usually easy to start from scratch again by pressing ESCAPE to abort processing the current command.
- One common problem when creating ggplot2 graphics is to put the + in the wrong place: it has to come at the end of the line, not the start. In other words, make sure you haven't accidentally written code like this:

```
ggplot(data = mpg)
+ geom_point(mapping = aes(x = displ, y = hwy))
```

# LAB: Exploring the titanic data

# LAB: Exploring the titanic data

Sample solution: <https://bio304-class.github.io/bio304-fall2017/data-story-titanic.html>

Other useful references to learn plotting:

[Data Visualization with ggplot2](#)

[Lesson 2: Visualizing Data Using ggplot2](#)

<https://bookdown.org/mikemahoney218/IDEAR/visualizing-your-data.html>

<https://r4ds.had.co.nz/data-visualisation.html>



# Next class

dplyr

TUE, OCT 13

Data Manipulation

*DUE: Problem Set 1 - Exploring Data*

Required Reading:

★ Wickham and Grolemund (2017). R for Data Science.  
Chapters 9-14. OReilly Media, Inc.

*Lab: Working with Data Part I*

DUE

- Lab2: upload by tomorrow (Oct 9) 9am PT
- PS1: deadline next Tuesday (Oct 13) 9am PT