Data Scientifique

GradProSkills

Introduction to R: Day 3
Data Visualization

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Workshop Schedule

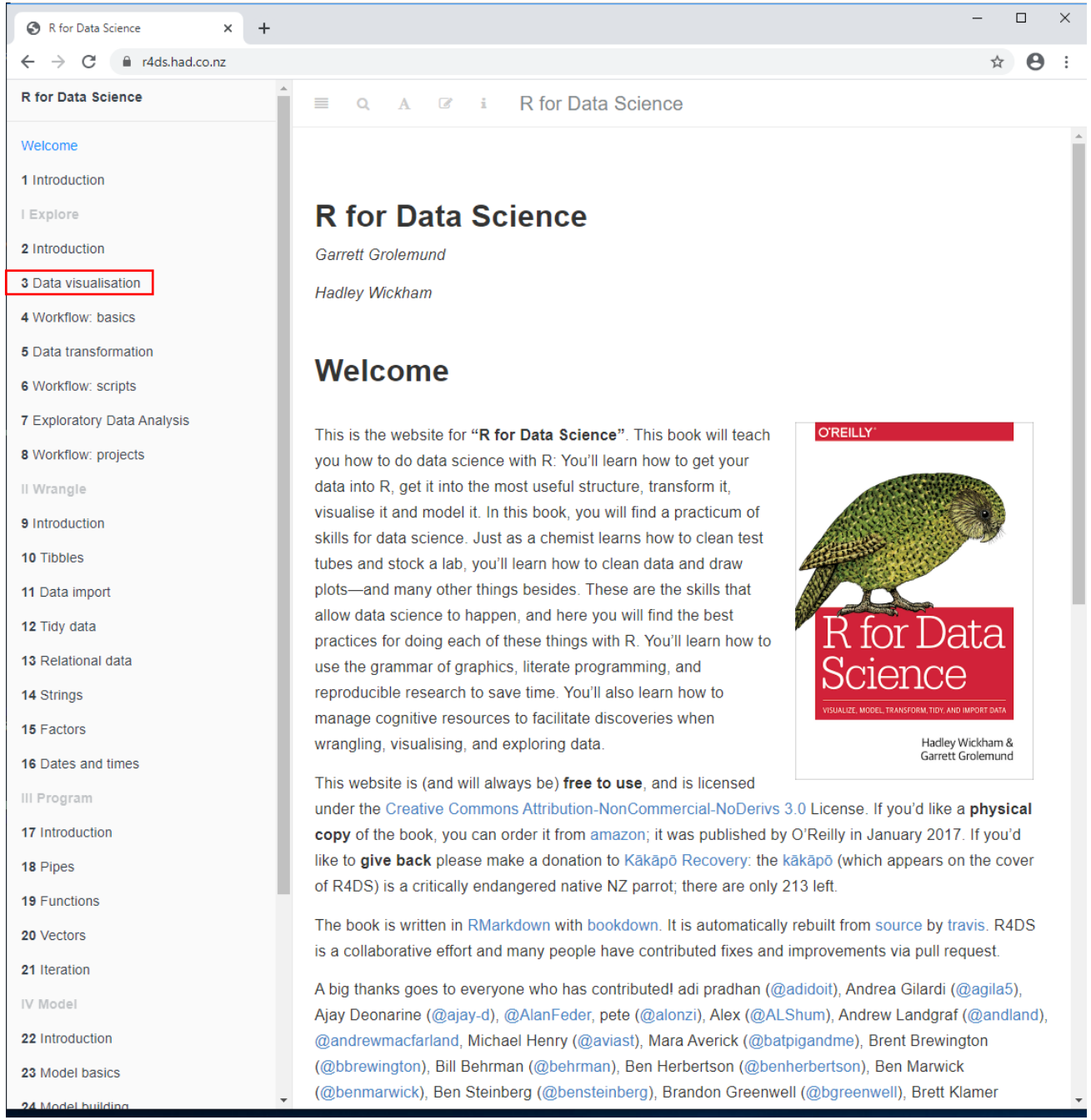
- In this workshop, the following topics will be covered:
  - Mastering R basics: will include an introduction to the R environment, packages and data types
  - Describing data: will demonstrate how to generate descriptive statistics, table outputs and simple statistical tests (i.e. t tests)
  - Visualizing data: will show participants how to generate multiple types of plots and charts

Plan for today

- First 15 minutes will consist of a quick review
- Go to https://github.com/YaraRAA/DataSci_IntrotoR
- Scroll down to: **README.md**
- You will see some poll links with numbers
- Let’s begin!

Plan for today ctd...

- Then….
- I will talk for an hour (if you're lucky, maybe less) about how to draw plots
  - using base R
  - using ggplot2
- You will get a chance to practice afterwards with the exercises provided. Again, I recommend that you work in pairs.



Textbook for ggplot2

- https://r4ds.had.co.nz/
- Chapter 3 of R for Data Science
- Why ggplot2?
  - Very popular package
  - A lot of examples online
  - Part of the tidyverse collection of packages https://www.tidyverse.org/

me →

Some words about the tidyverse

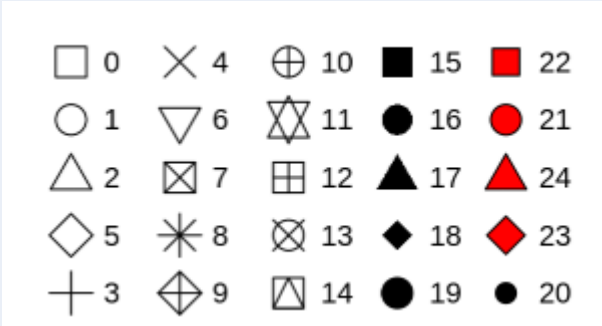
- I personally do not use tidyverse packages, so bear with me
- I also disagree with some of the suggestions in this textbook but feel free to follow it if it works for you!
- FYI, I like the following packages:
  - base R for basic plotting (if I need to quickly check a distribution)
  - mgcv plot of a gam model for the non-linear shape of a dose-response (gam stands for generalized additive model)
  - plotly for fancy plots for publication (they have a lot of easy tutorials online)
  - data.table for data management - data.table is excellent for very large (100 million + rows) datasets

Image source: https://www.amazon.co.uk/3dRose-Inches-Grumposaurus-Dinosaur-mp_104180_1/dp/B00BMCQSQU

Plots using base R

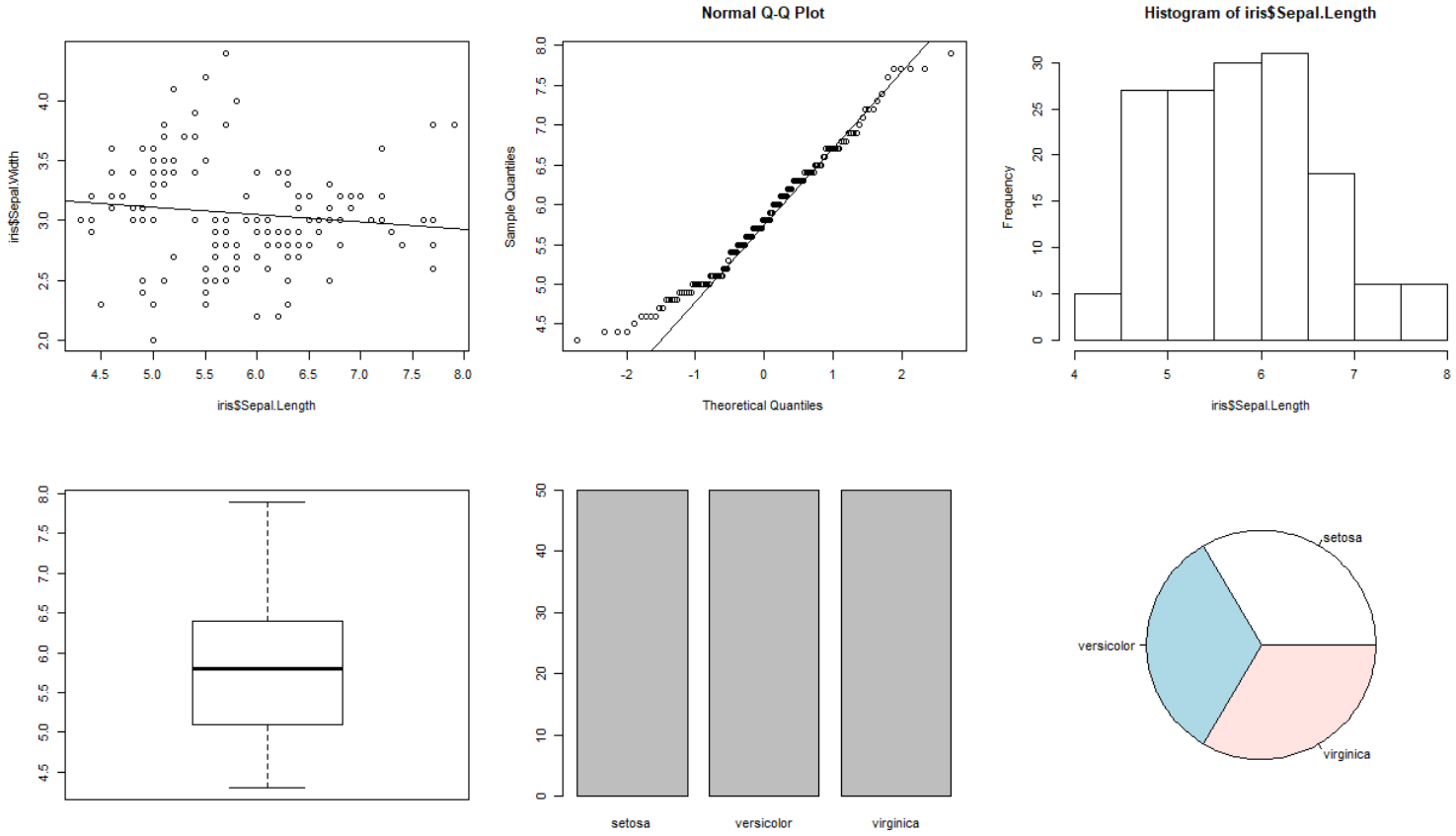
Function	Input	Output
Plot abline	(x , y) 2 numeric vectors i.e. 2 columns (a = constant, b = slope)	Scatter plot Adds one or more straight lines through the current plot
hist	One numeric vector i.e. one column	Histogram of a continuous variable
qqnorm qqline	One numeric vector i.e. one column	Normal qqplot of a continuous variable Adds a line to a “theoretical”, by default normal, quantile-quantile plot
boxplot	One numeric vector i.e. one column	Box plot
pie	A table object	Pie chart
barplot	A table object	Bar plot

Useful arguments in base R plot functions

Argument	Input	What it does
main = , xlab = , ylab =	'title in quotes'	Adds main title, x axis and y axis title to plot respectively
add =	T	Add this new plot to the previous
xlim =, ylim =	c(min, max)	Sets beginning and end of x axis & y axis respectively
col =	a color in quotations i.e. 'red' OR rgb(red,green,blue,transparency)	Determines the colour of the plot
cex =	number	Relative size of text
pch =	number 	Determines shape of points Same numbers are also used in ggplot2 syntax but instead of pch = , the syntax is shape =

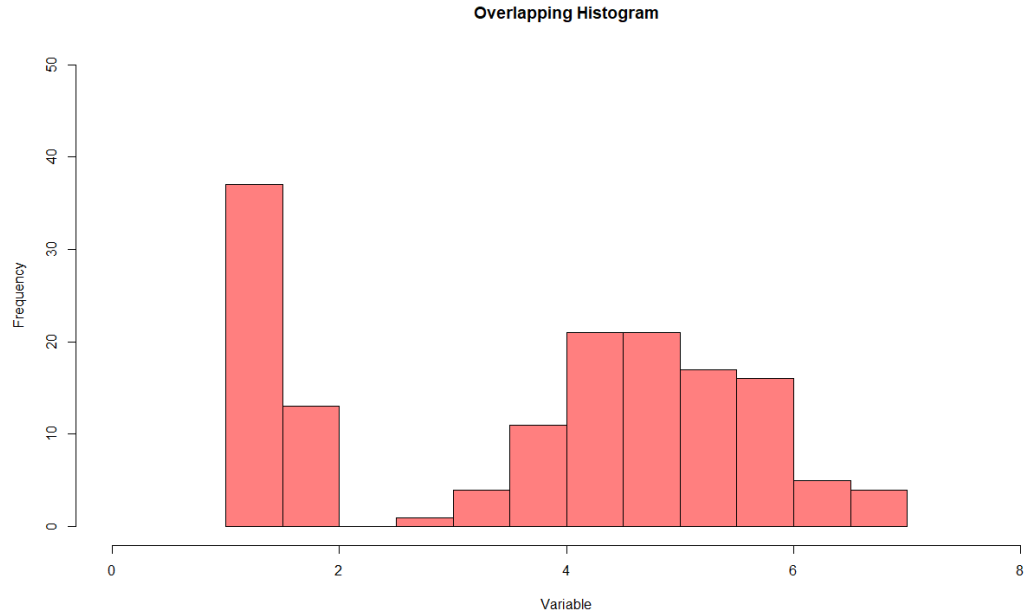
Example: multiple plots in one

```
par(mfrow =c(2,3))
plot(iris$Sepal.Length, iris$Sepal.Width)
abline(lm(Sepal.Width ~ Sepal.Length, data = iris ))
qqnorm(iris$Sepal.Length)
qqline(iris$Sepal.Length)
hist(iris$Sepal.Length)
boxplot(iris$Sepal.Length)
barplot(table(iris$Species))
pie(table(iris$Species))
```

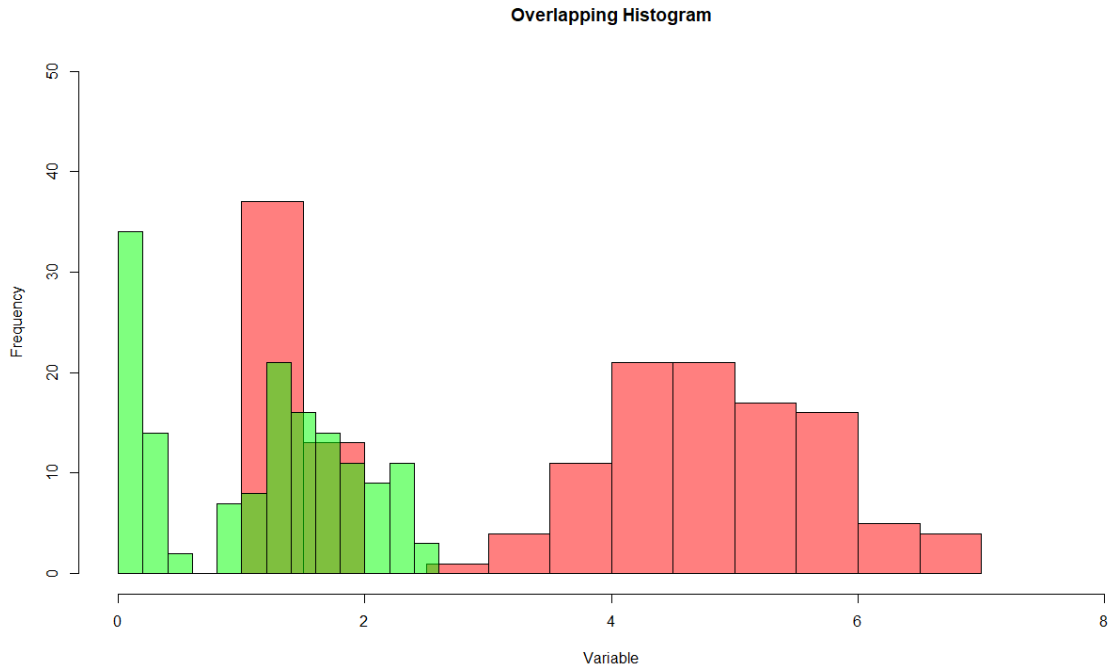


Example: overlaying plots

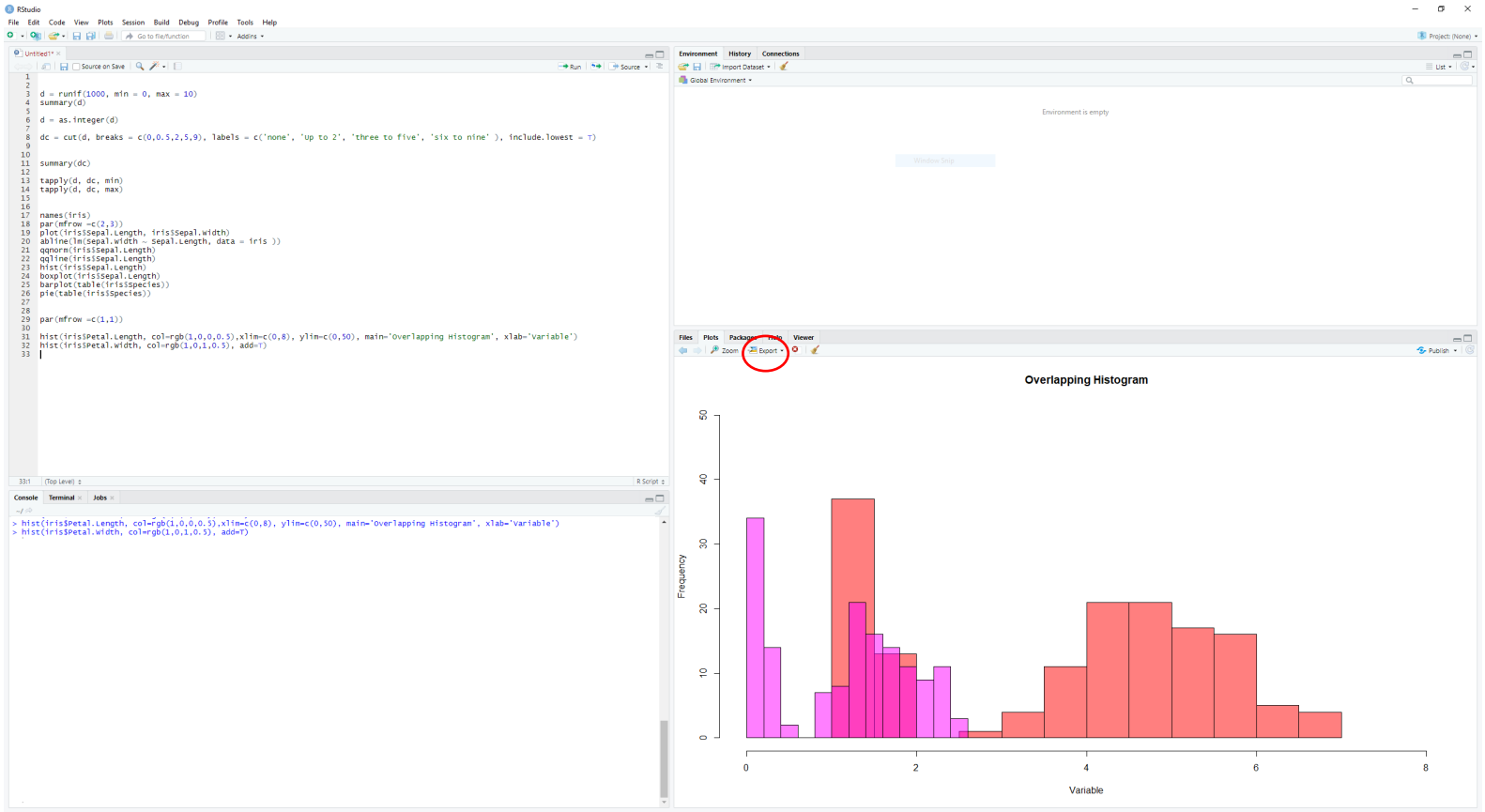
```
hist(iris$Petal.Length, col=rgb(1,0,0,0.5),xlim=c(0,8),
ylim=c(0,50), main='Overlapping Histogram', xlab='Variable')
```




```
hist(iris$Petal.Width, col=rgb(0,1,0,0.5), add=T)
```



Saving your plot



Alternatively, before plotting run code:

pdf('nameoffile.pdf')

plot(x, y)

dev.off()

Saves your plot to a pdf file named 'nameoffile.pdf' in your working directory.

Now to ggplot2

- First install the tidyverse packages and then load them:

```
install.packages("tidyverse")
library(tidyverse)
```

- Alternatively:

```
install.packages(“ggplot2")
library(ggplot2)
```

We will be using the mpg data frame which is found in ggplot2

```
mpg
#> # A tibble: 234 x 11
#>   manufacturer model displ  year   cyl trans      drv     cty   hwy fl    class
#>   <chr>        <chr> <dbl> <int> <int> <chr>      <chr> <int> <int> <chr> <chr>
#> 1 audi         a4      1.8  1999     4 auto(l5)   f        18    29 p     compa…
#> 2 audi         a4      1.8  1999     4 manual(m5) f        21    29 p     compa…
#> 3 audi         a4      2    2008     4 manual(m6) f        20    31 p     compa…
#> 4 audi         a4      2    2008     4 auto(av)   f        21    30 p     compa…
#> 5 audi         a4      2.8  1999     6 auto(l5)   f        16    26 p     compa…
#> 6 audi         a4      2.8  1999     6 manual(m5) f        18    26 p     compa…
#> # … with 228 more rows
```

Among the variables in mpg are:

1. displ, a car's engine size, in litres.

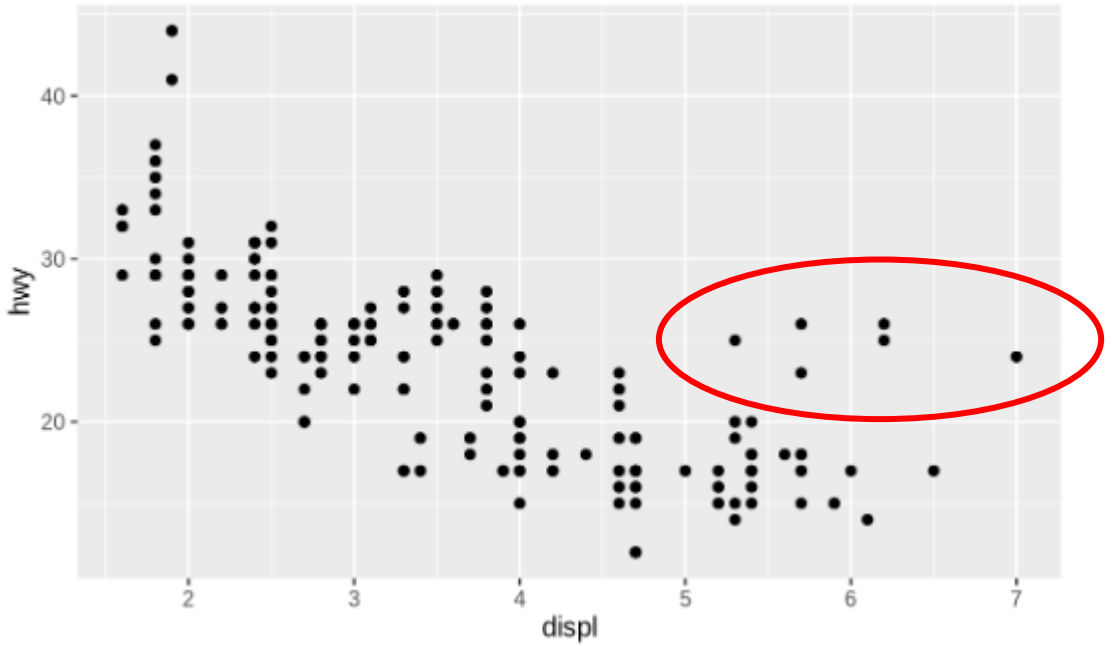
2. hwy, a car's fuel efficiency on the highway, in miles per gallon (mpg). A car with a low fuel efficiency consumes more fuel than a car with a high fuel efficiency when they travel the same distance.

Creating a ggplot

- To plot mpg, run this code to put displ on the x-axis and hwy on the y-axis:

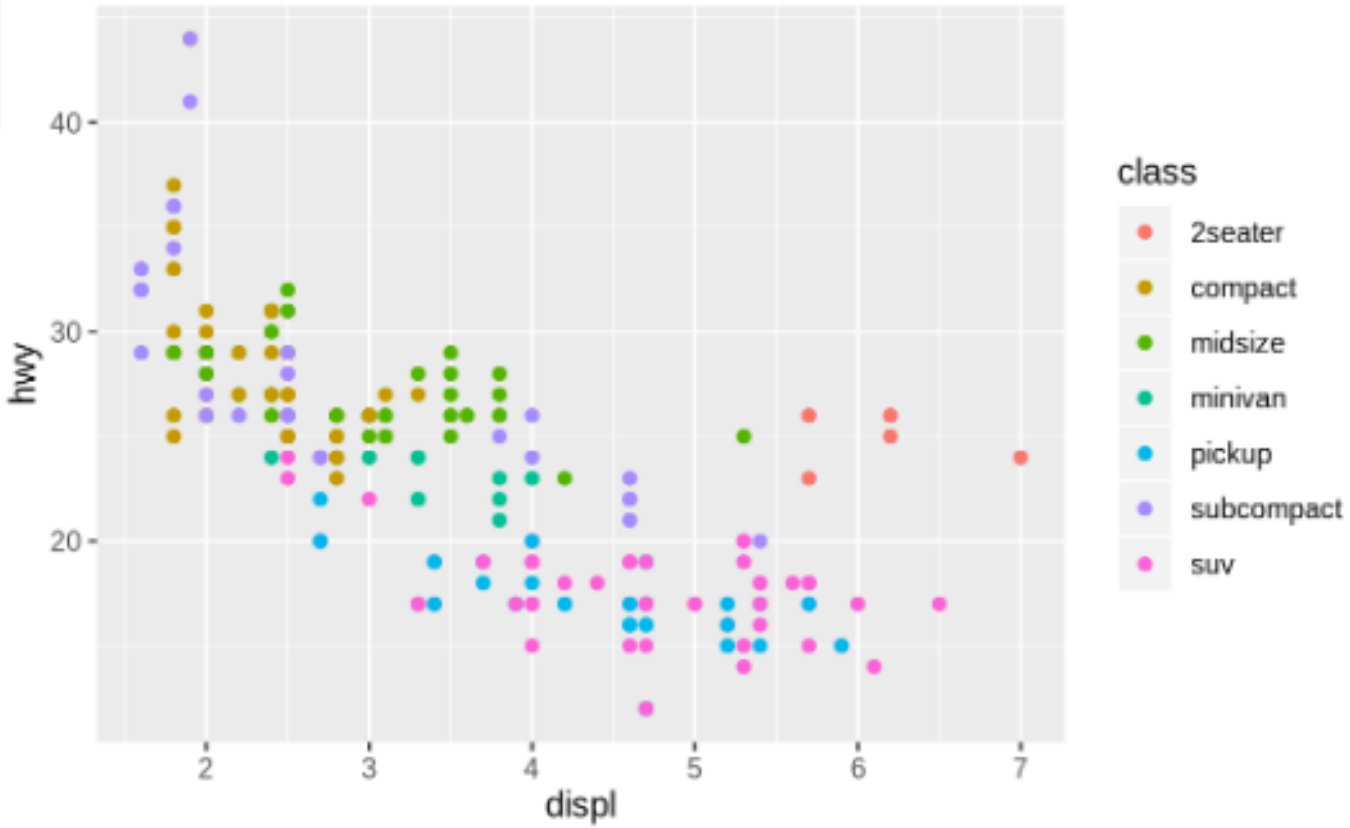
```
ggplot(data = <DATA>) +
  <GEOM_FUNCTION>(mapping = aes(<MAPPINGS>))
```

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



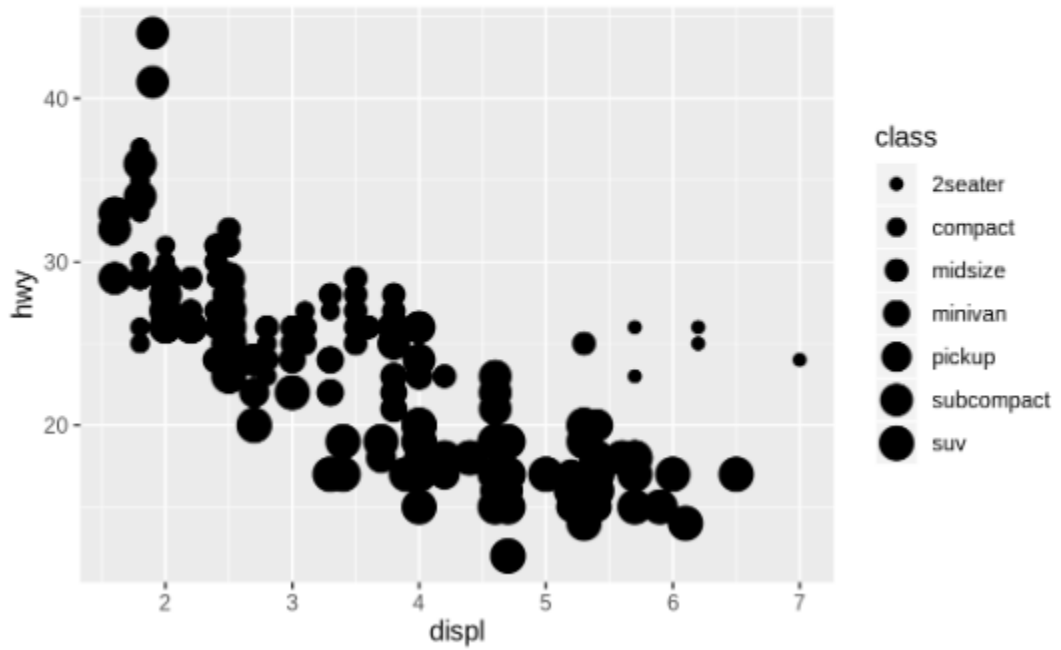
Colour coded point data

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



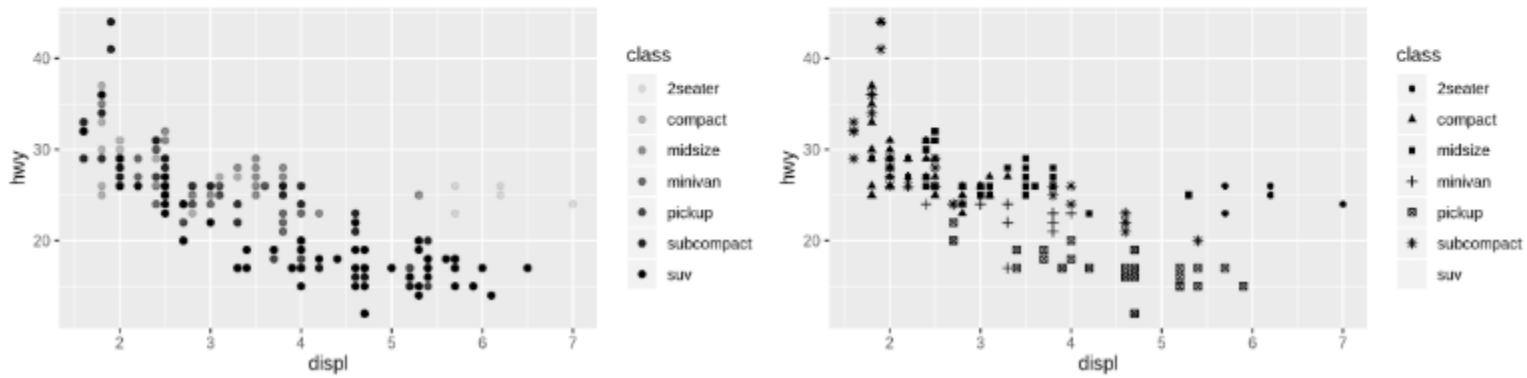
Indicating class by point size

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



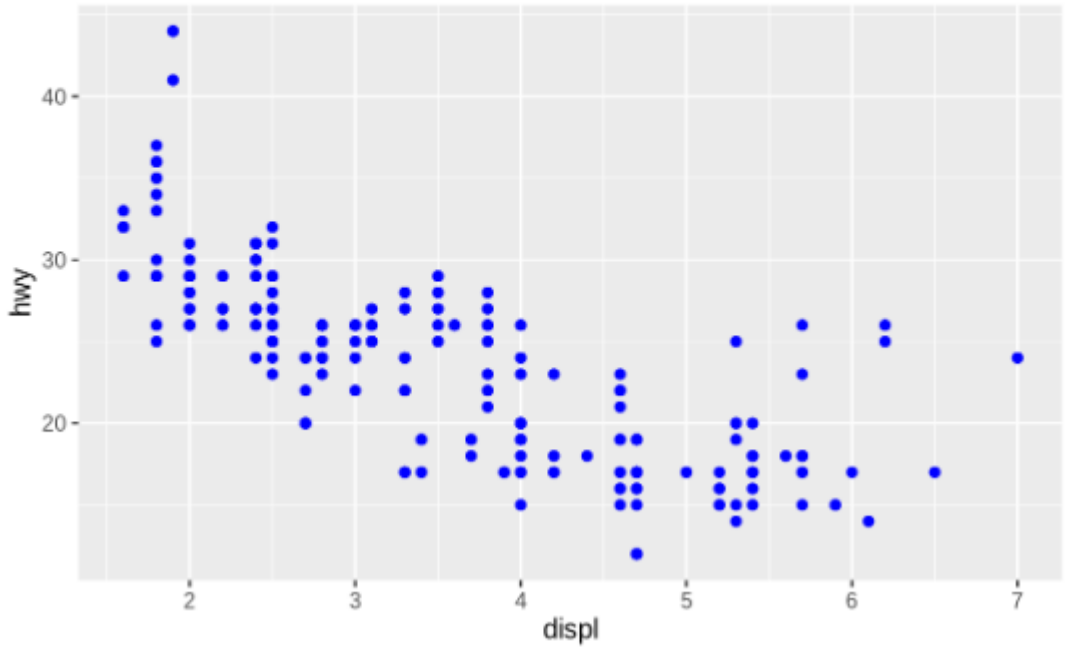
Indicating class by point shape / transparency

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

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



Selecting the color of your points

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



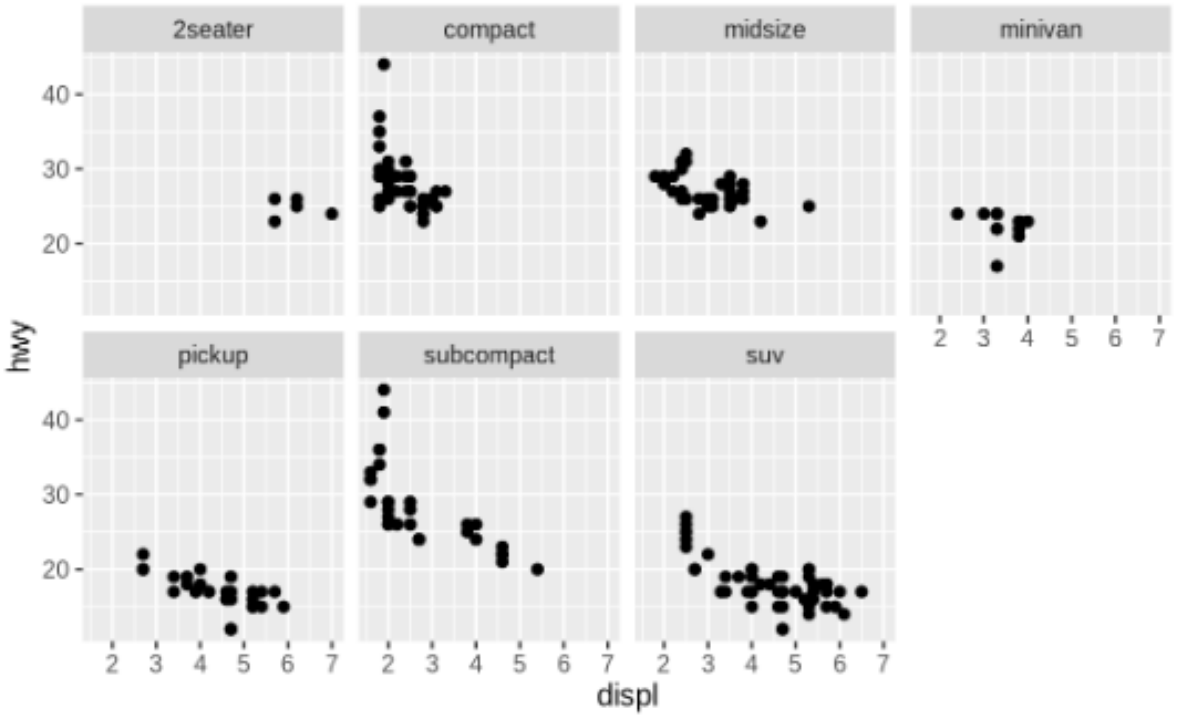
Avoiding errors in your code

- Make sure:
  - you have entered the code exactly as it is presented
  - commas are in the right place
  - every opening quote " has a closing quote "
  - every open bracket (has a closing bracket)
  - specifically in ggplot2, the + has to come at the end of the line

- If you run code and nothing happens, you will see a + sign in your console. This means that R is waiting for you to close a function that you started (conversely, > means that R is ready for a new command):
  - when this happens, hit the escape button to abort the current command and check your code

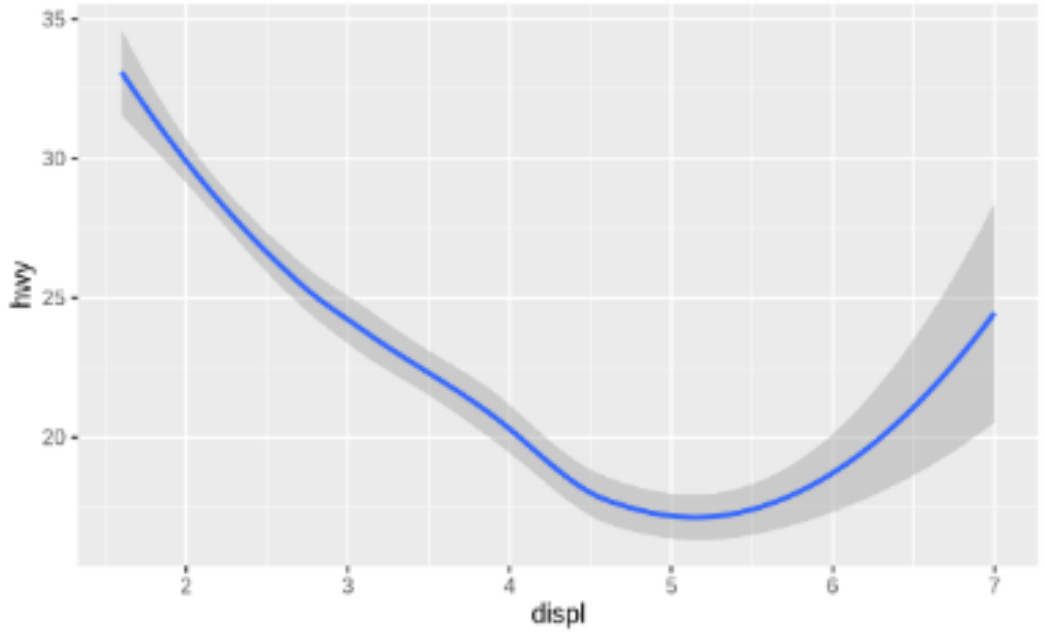
Plotting subsets of data

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

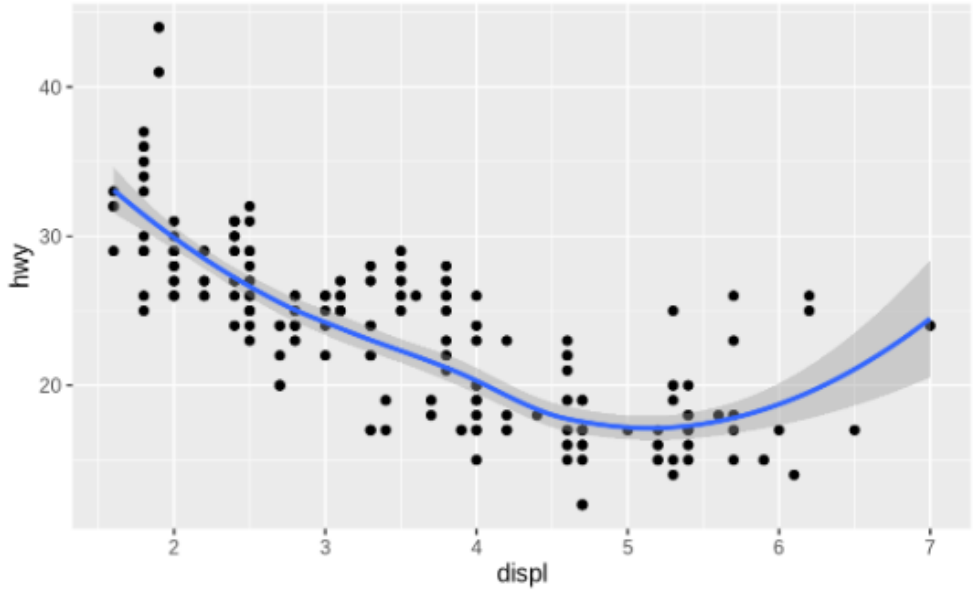


Applying a smooth function to your scatter plot

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

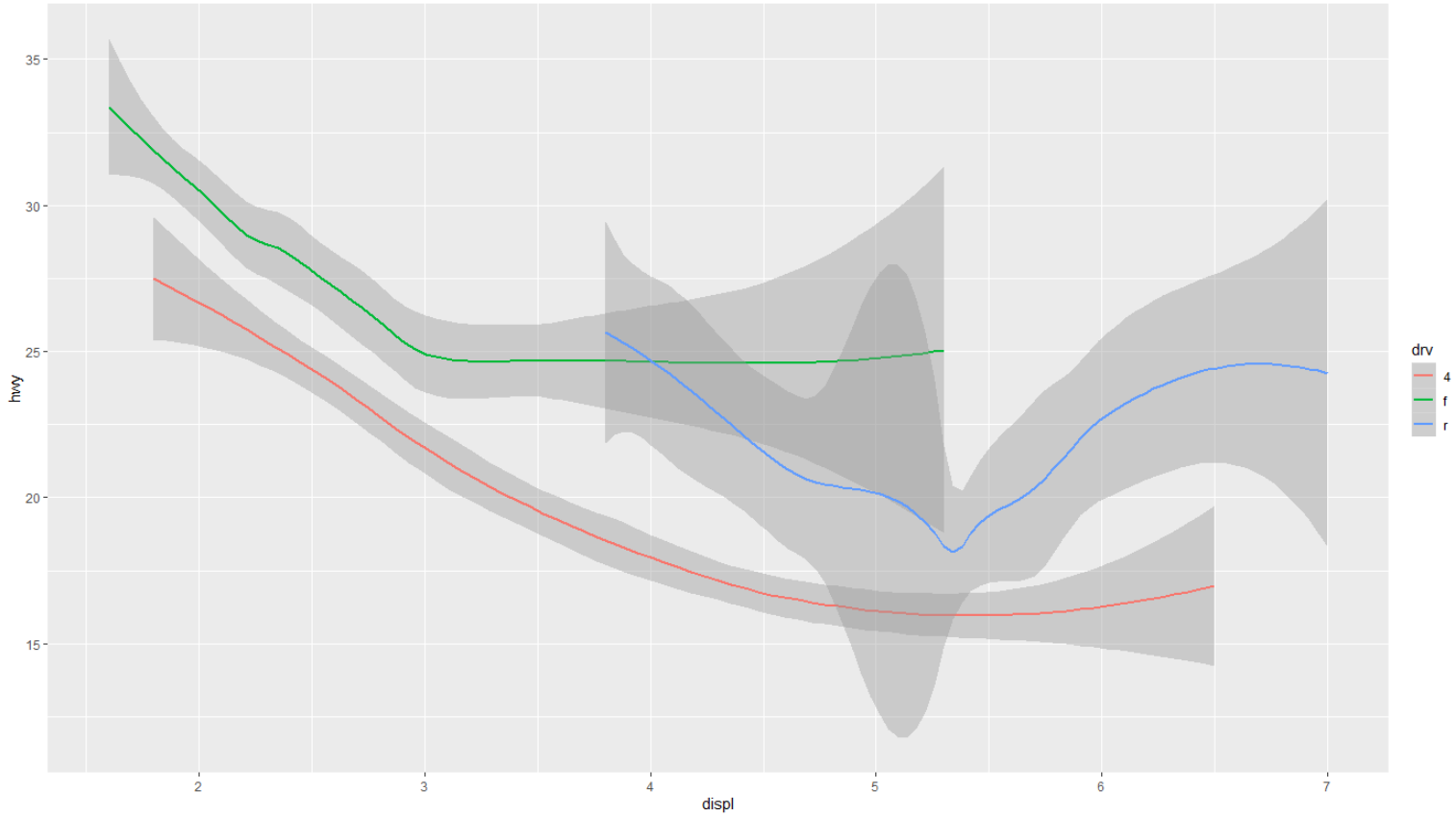



```
ggplot(data = mpg) +
  geom_point(mapping = aes(x = displ, y = hwy)) +
  geom_smooth(mapping = aes(x = displ, y = hwy))
```



Apply smooth functions to different subsets

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



drv
f = front-wheel drive,
r = rear wheel drive,
4 = 4wd

Note: by default, geom_smooth
Uses mgcv gam when there are
> 1000 observations

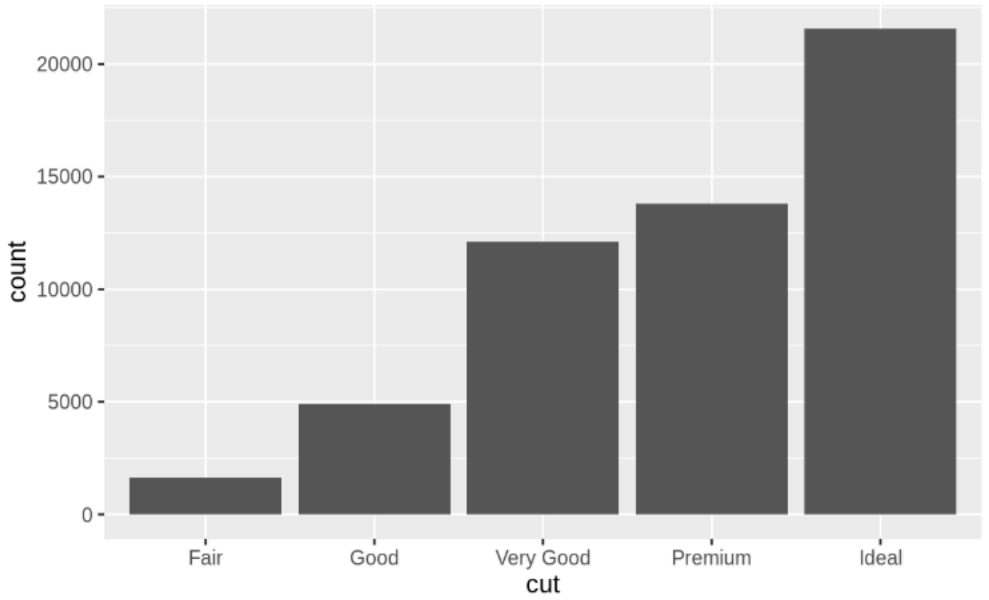
What about categorical data?

- Data frame diamonds in ggplot2

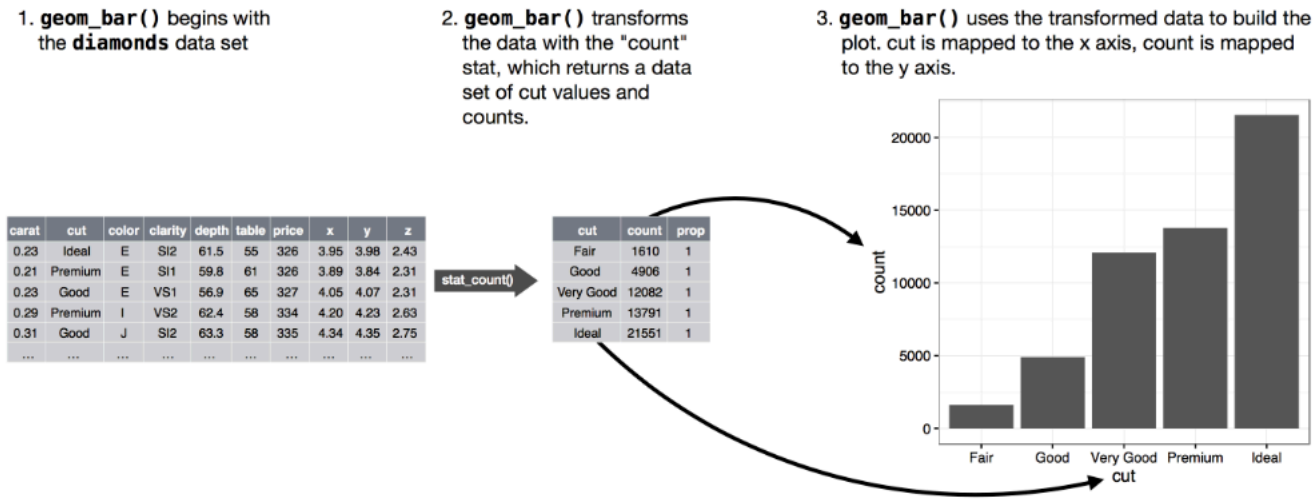
```
> head(diamonds)
# A tibble: 6 x 10
  carat cut       color clarity depth table price     x     y     z
  <dbl> <ord>     <ord> <ord>   <dbl> <dbl> <int> <dbl> <dbl> <dbl>
1 0.23  Ideal     E     SI2      61.5    55   326  3.95  3.98  2.43
2 0.21  Premium   E     SI1      59.8    61   326  3.89  3.84  2.31
3 0.23  Good      E     VS1      56.9    65   327  4.05  4.07  2.31
4 0.290 Premium   I     VS2      62.4    58   334  4.2   4.23  2.63
5 0.31  Good      J     SI2      63.3    58   335  4.34  4.35  2.75
6 0.24  Very Good J     VVS2     62.8    57   336  3.94  3.96  2.48
```

Bar plot: Frequency of diamonds by quality of cut

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

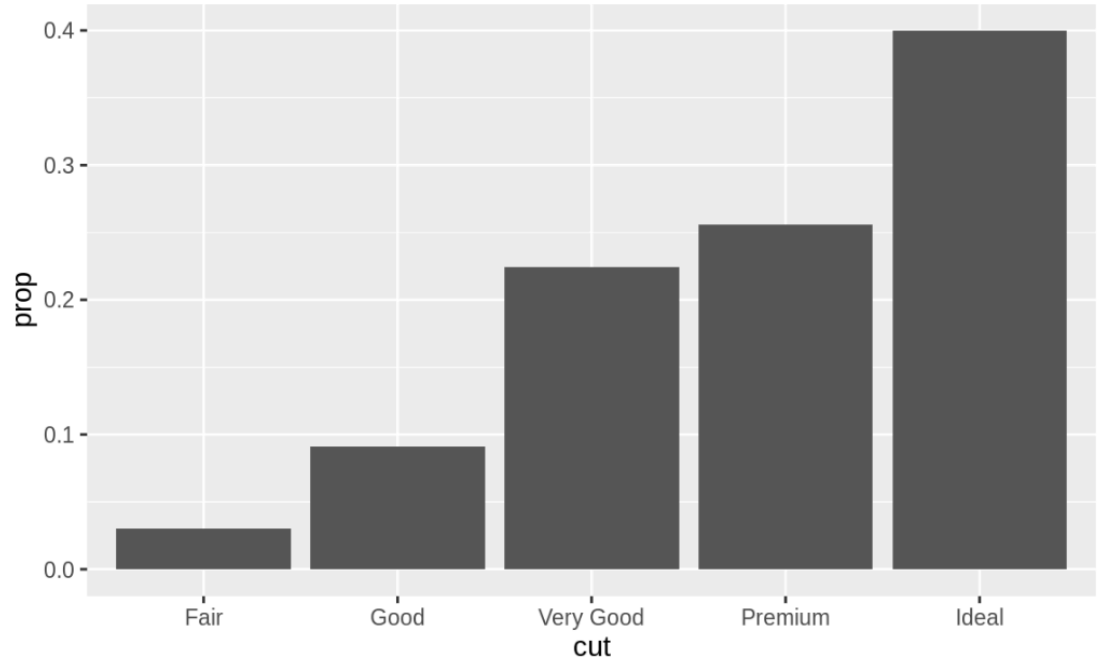


Bar plot: A count variable is created!



Bar plot: what if I want to plot proportion instead?

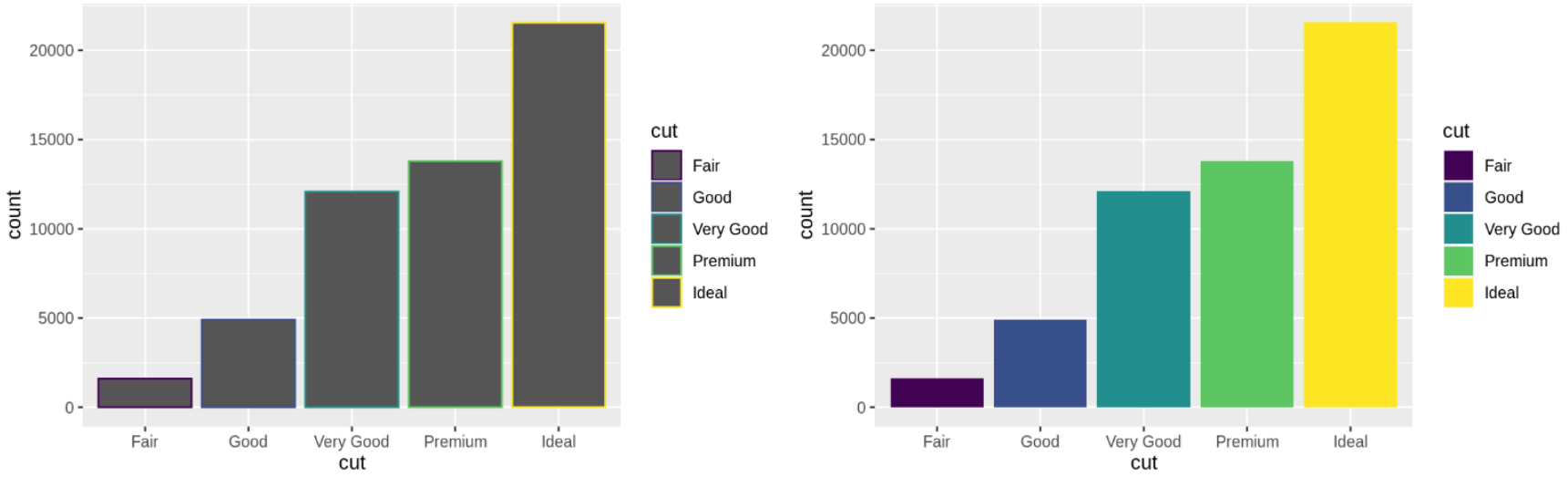
```
ggplot(data = diamonds) +
  geom_bar(mapping = aes(x = cut, y = stat(prop), group = 1))
```



Note: 'ggplot2 provides over 20 stats for you to use. Each stat is a function, so you can get help in the usual way, e.g. ?stat_bin'

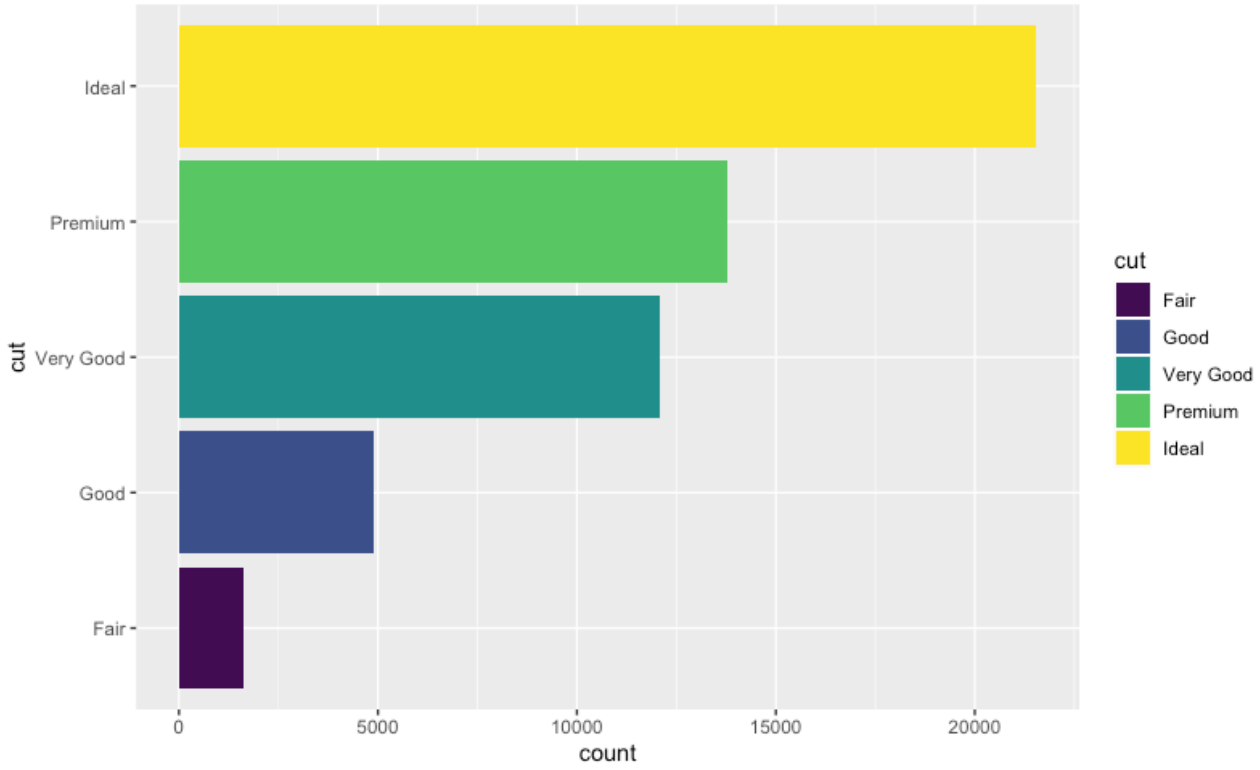
Bar plot: adding color

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



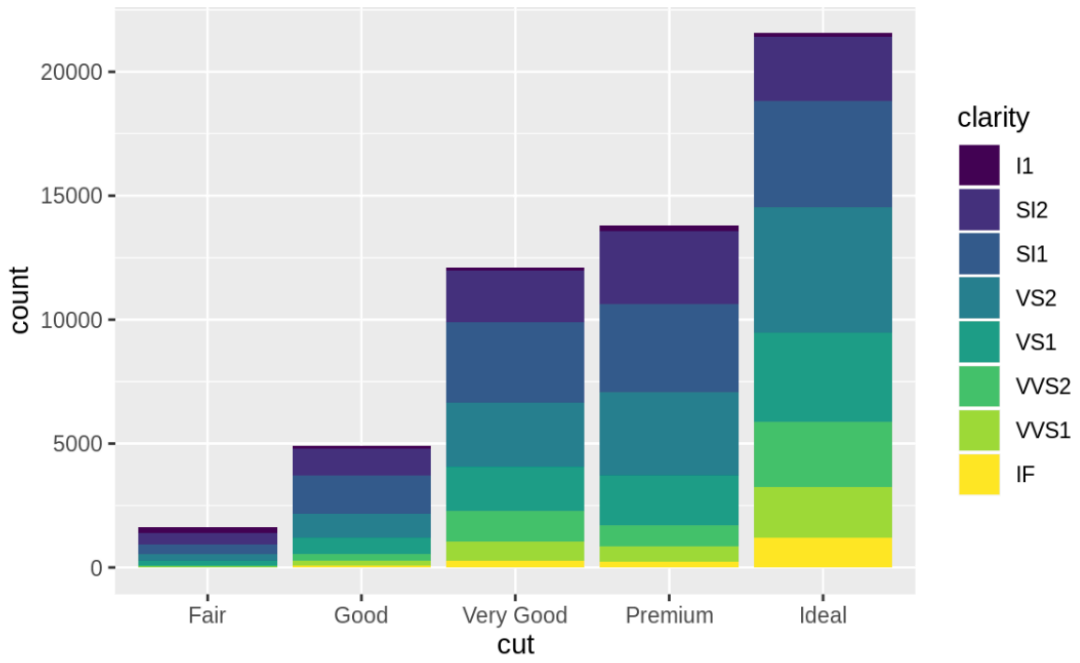
Bar plot: flipped to the side

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

bar + coord_flip()
```



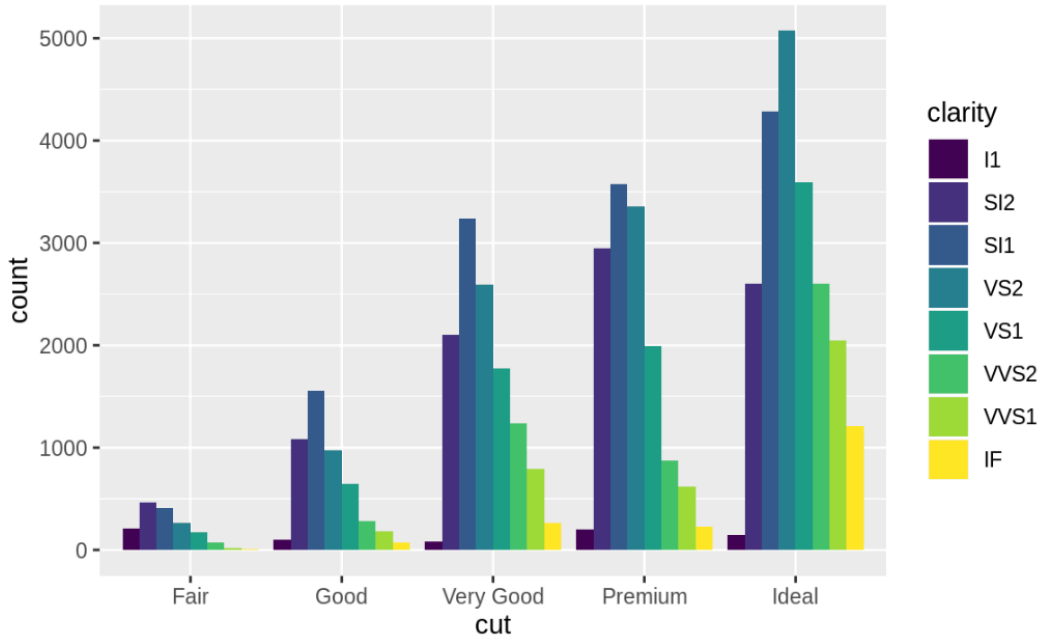
Bar plot: stacked bar plots

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



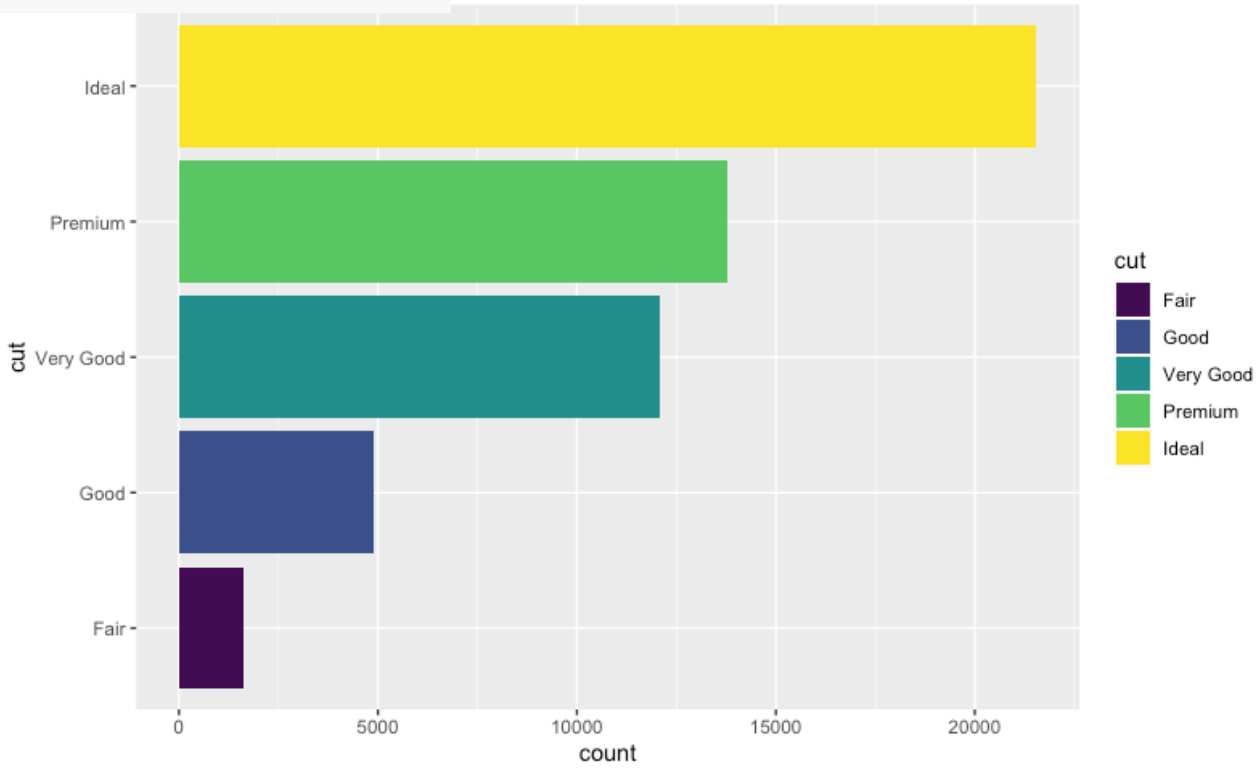
Bar plots: grouped bar plots

```
ggplot(data = diamonds) +
  geom_bar(mapping = aes(x = cut, fill = clarity), position = "dodge")
```



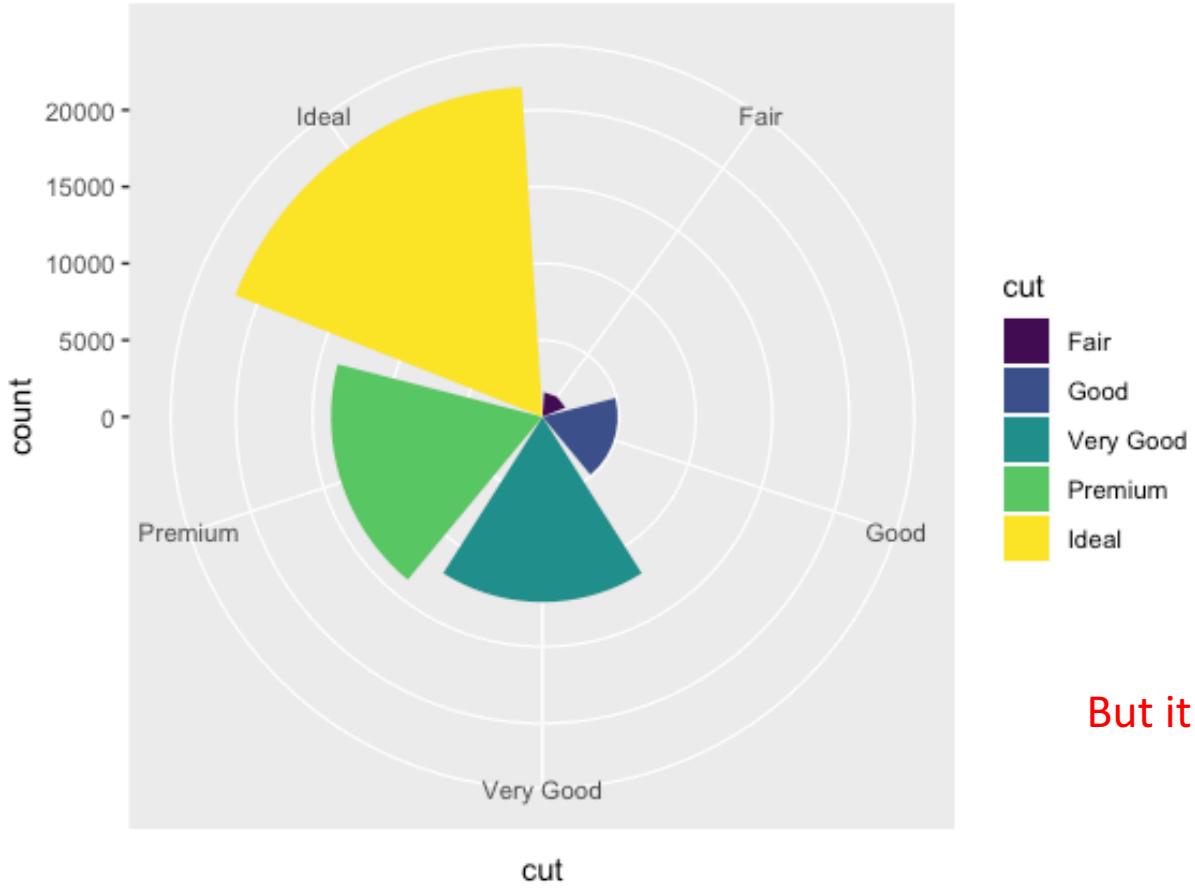
Pie Chart: Remember how we flipped the bar plot?

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

bar + coord_flip()
```



Pie chart: similarly, we can create a pie chart

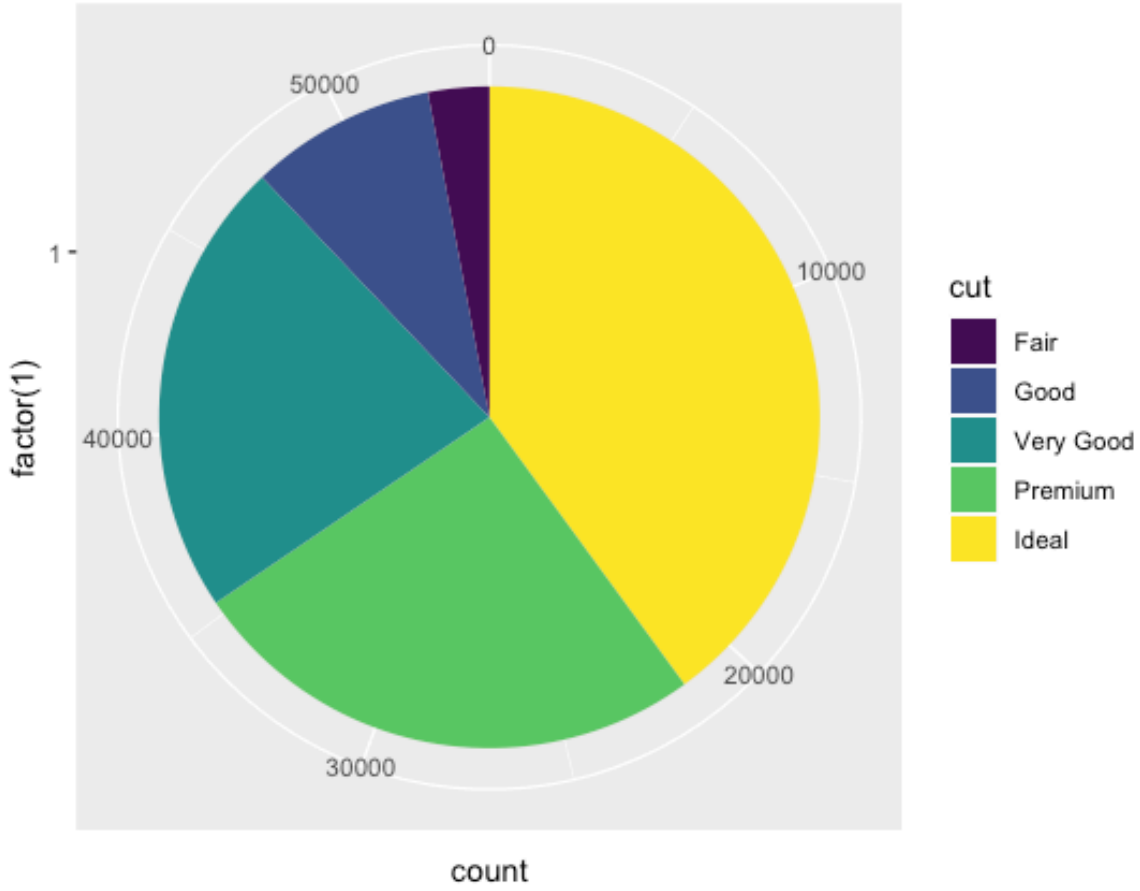
```
bar <- ggplot(data = diamonds) +
 geom_bar( mapping = aes(x = cut, fill = cut))

bar + coord_polar()
```



But it looks funky

Pie Chart

```
ggplot(diamonds, aes(x=factor(1), fill=cut))+
geom_bar(width = 1)+  coord_polar("y")
```



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