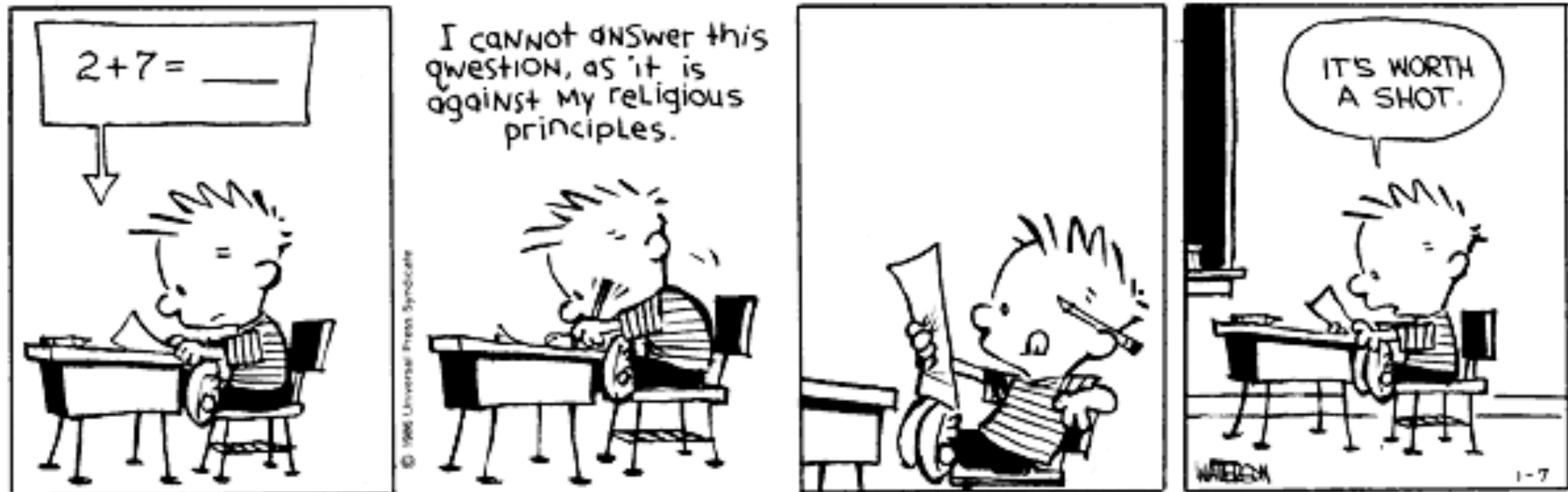




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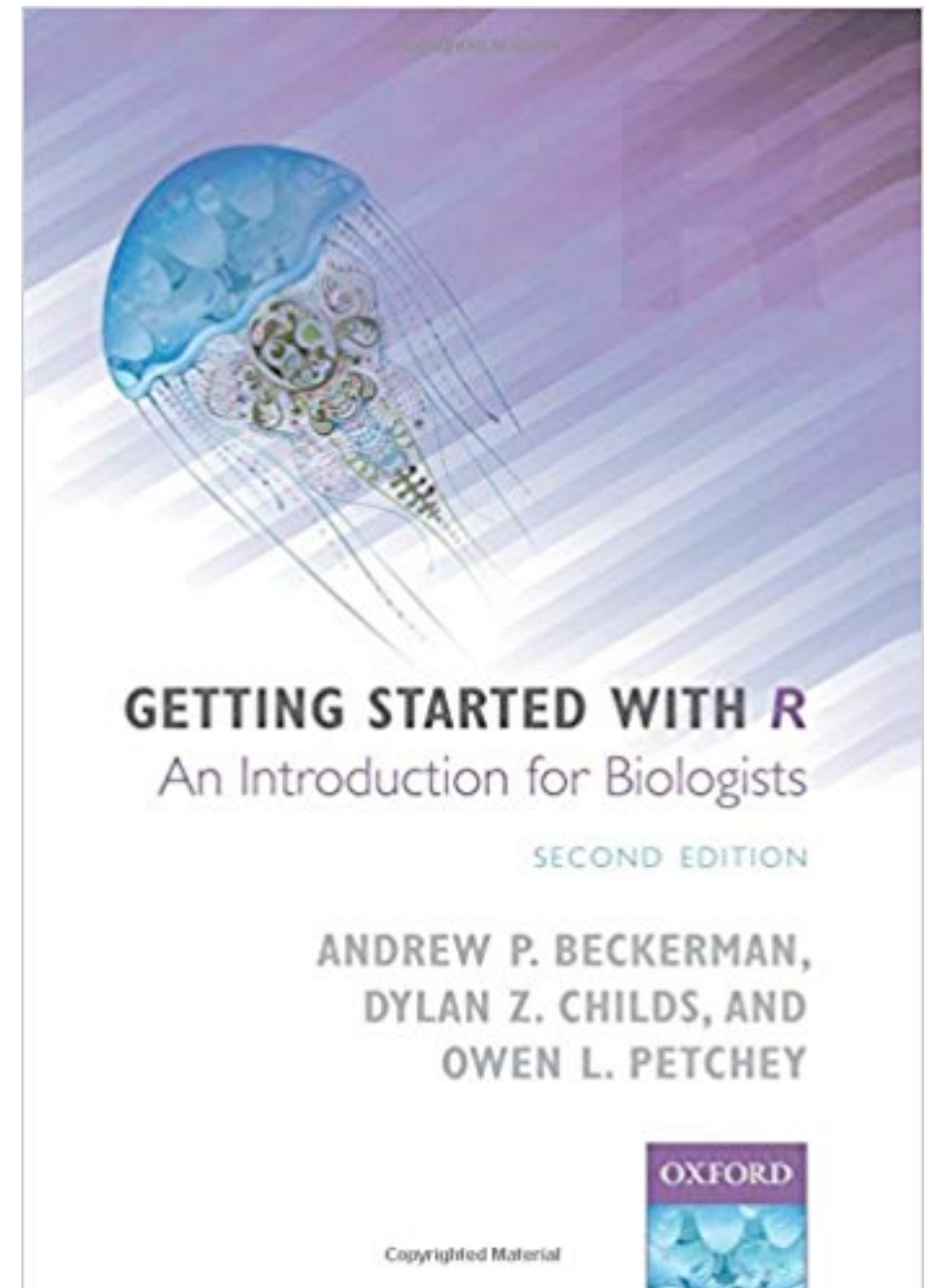
Introductory statistics with R

Solutions to exercises

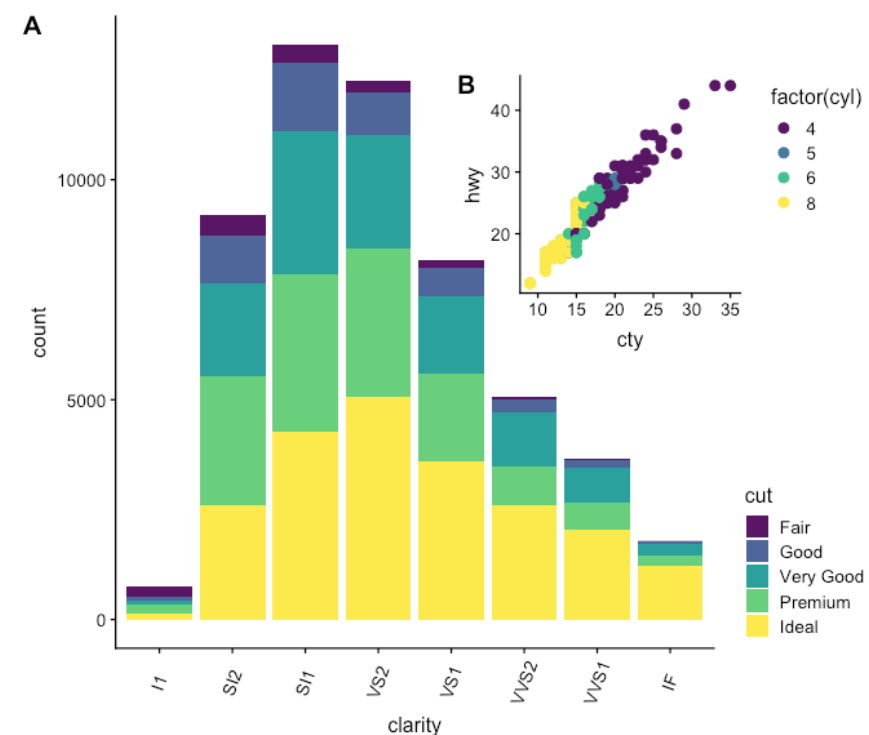
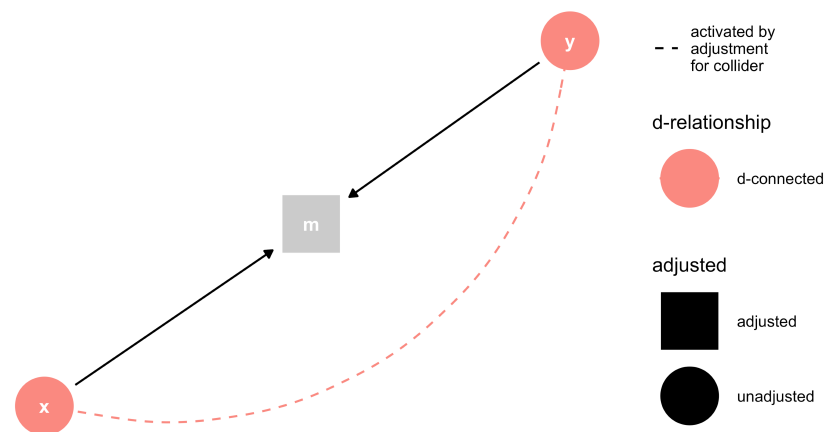
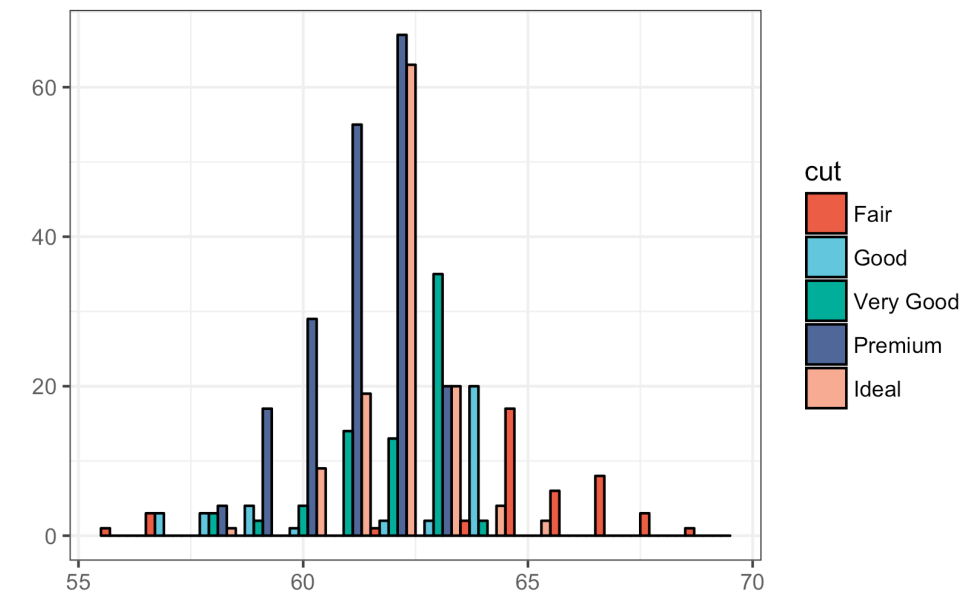
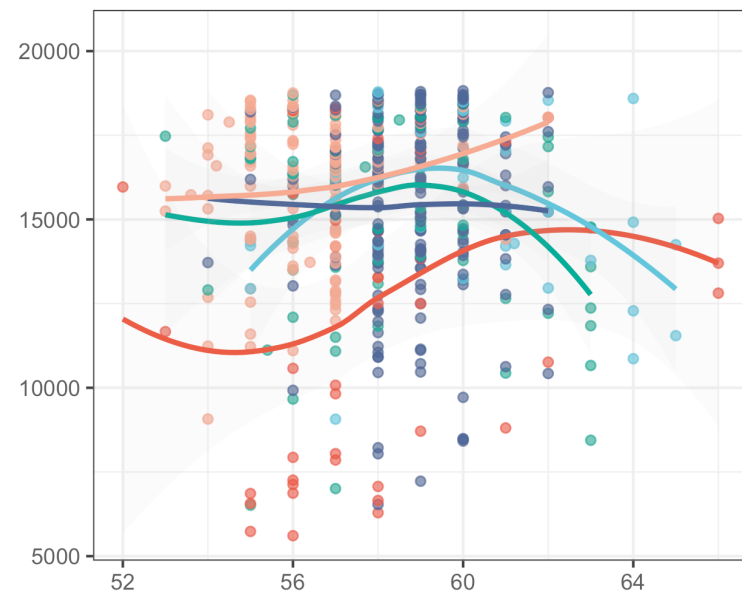


For next week

- We will read chapter 5 (only from 5.1 to 5.3).
Optional reading: Chapter 8 (As I will not go through R basics anymore, it is up to you if you want to pimp you figures).



How to visualise your data: ggplot



Why ggplot?



First plot with ggplot2: qplot

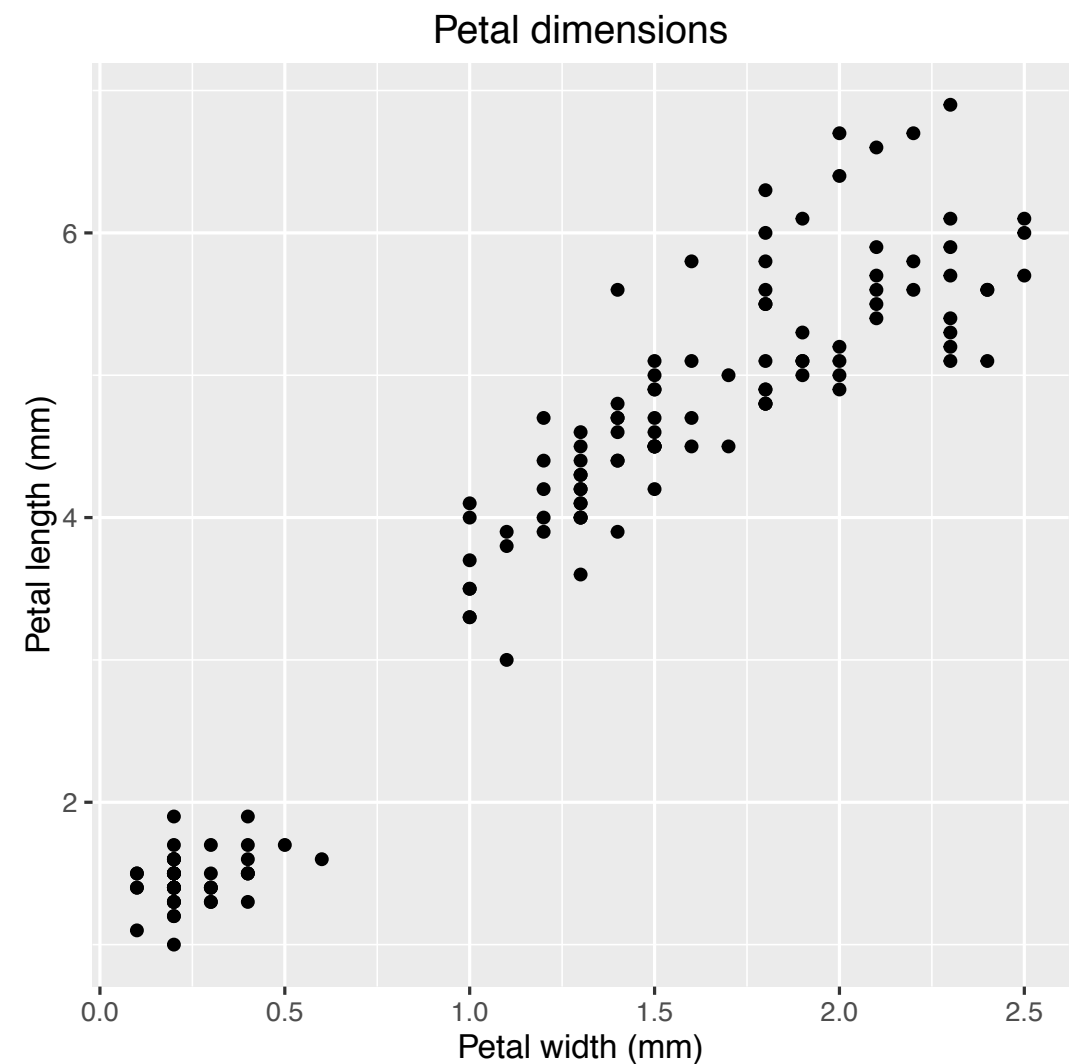
- It works similar to plot().

```
qplot(data = mydata,  
      y = var_y,  
      x = var_x, # could be optional  
      xlab = "x",  
      ylab = "y",  
      main = "a quick plot")
```

First plot with ggplot2: qplot

- It works similar to plot().

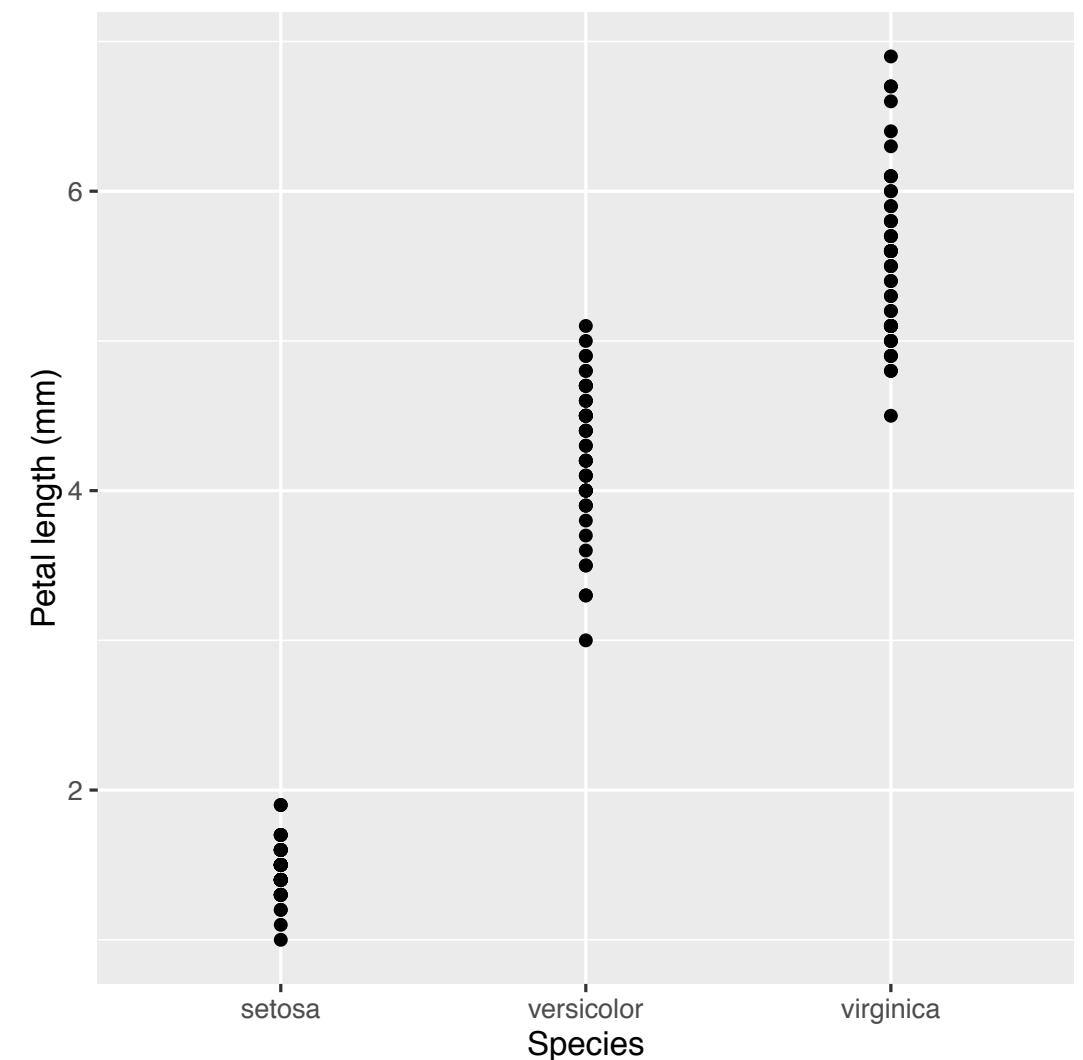
```
qplot(data = iris,  
      y = Petal.Length,  
      x = Petal.Width,  
      xlab = "Petal width (mm)",  
      ylab = "Petal length (mm)",  
      main = "Petal dimensions")
```



Lets try now with a categorical variable

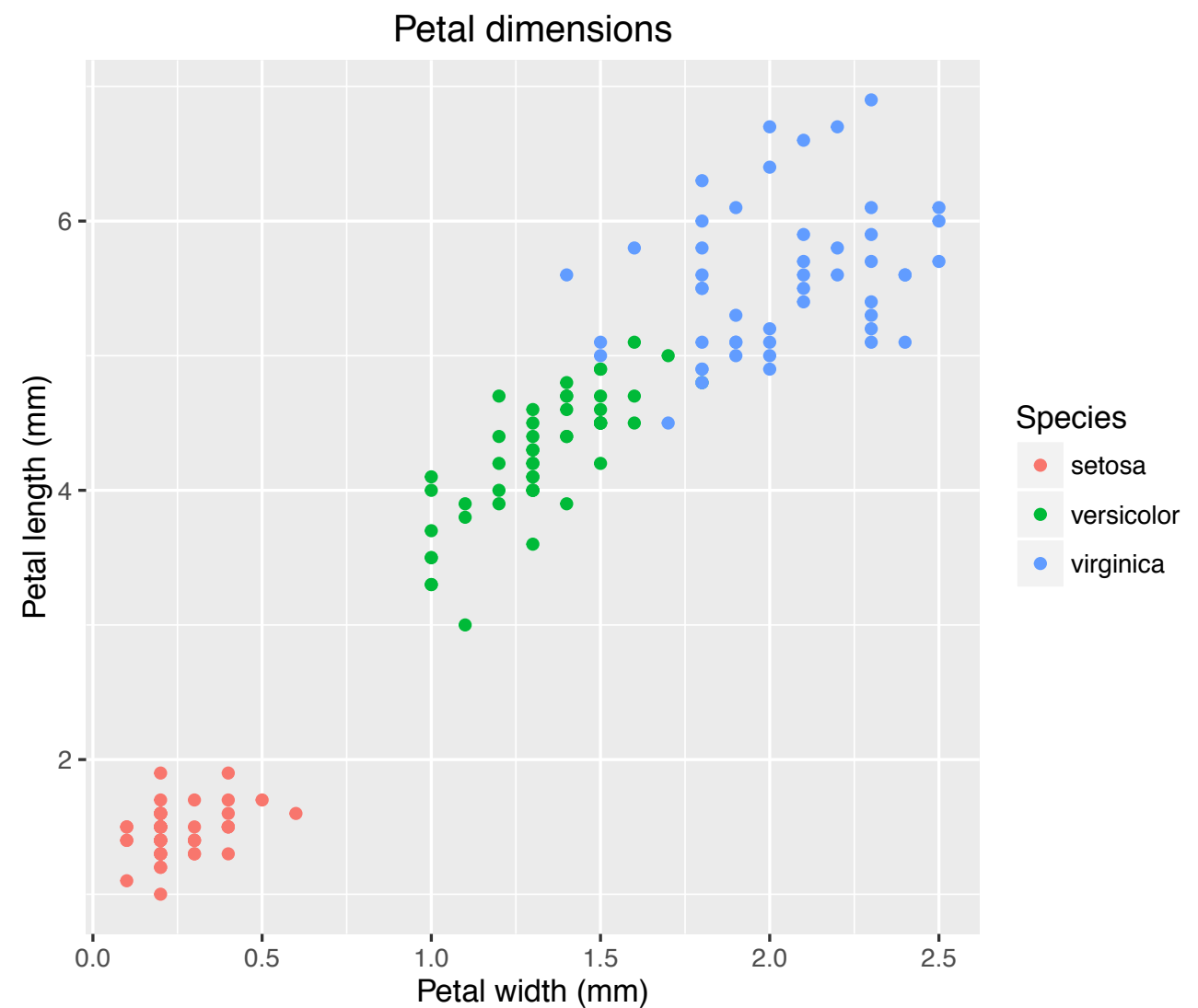
- It works similar to plot().

```
qplot(data = iris,  
      y = Petal.Length,  
      x = Species,  
      xlab = "Species",  
      ylab = "Petal length (mm)")
```



Lets pimp this a bit. (scale it!)

```
qplot(data = iris,  
      y = Petal.Length,  
      x = Petal.Width,  
      xlab = "Petal width (mm)",  
      ylab = "Petal length (mm)",  
      main = "Petal dimensions",  
      colour = Species)
```

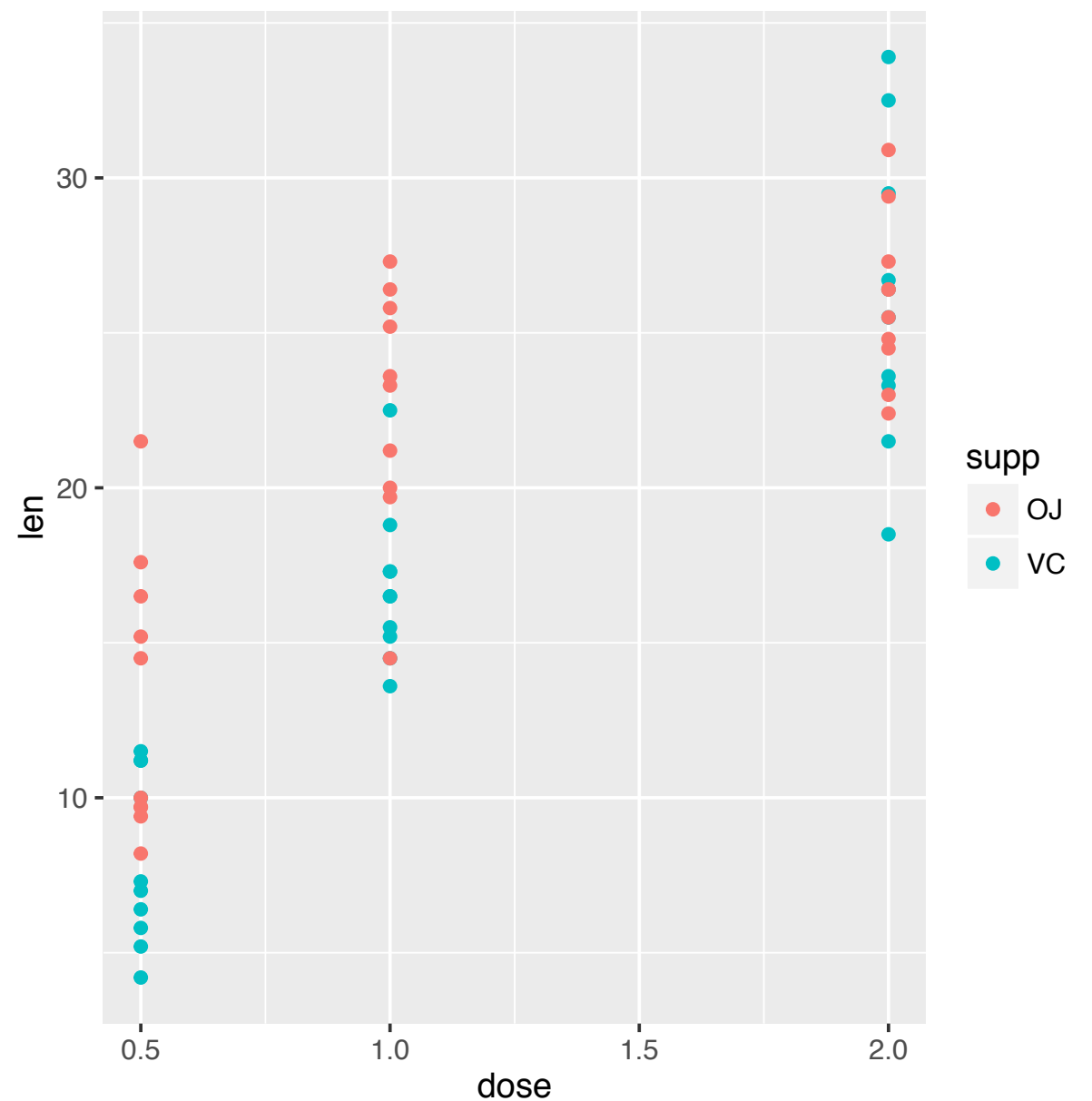


Try it yourself!

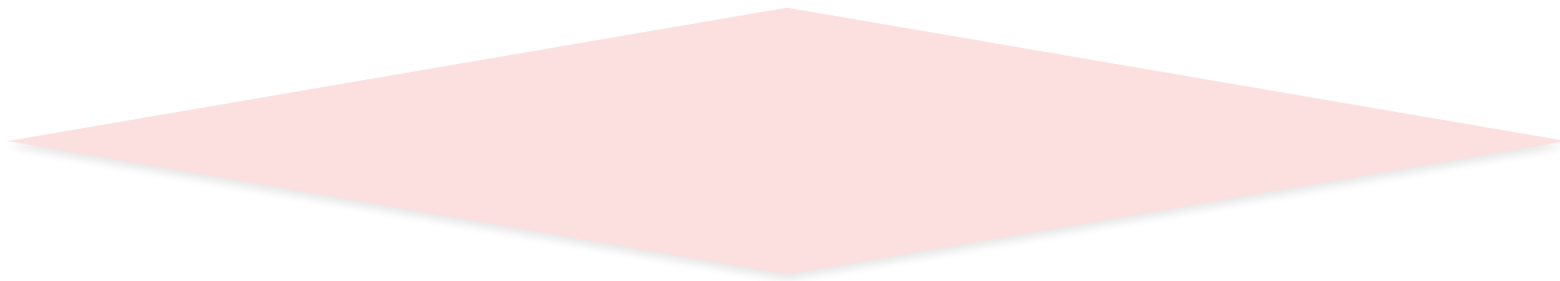
- Load the dataset ToothGrowth and produce a plot.

Try it yourself!

- Load the dataset ToothGrowth and produce a plot.

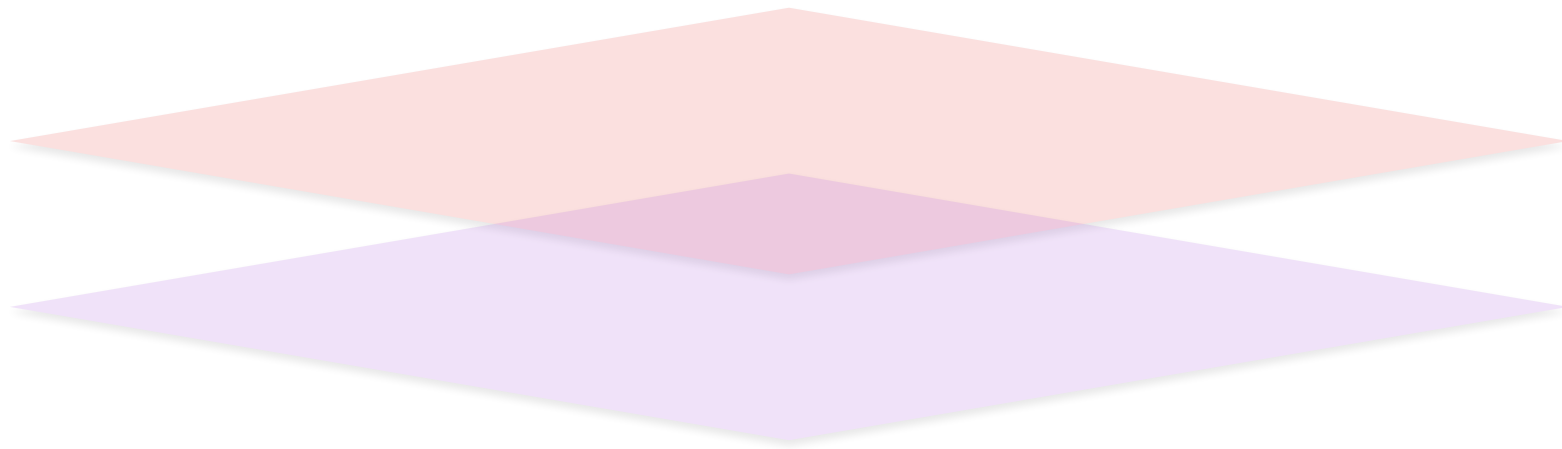


Layering in ggplot



Data

Layering in ggplot



Data

Mapping

Layering in ggplot



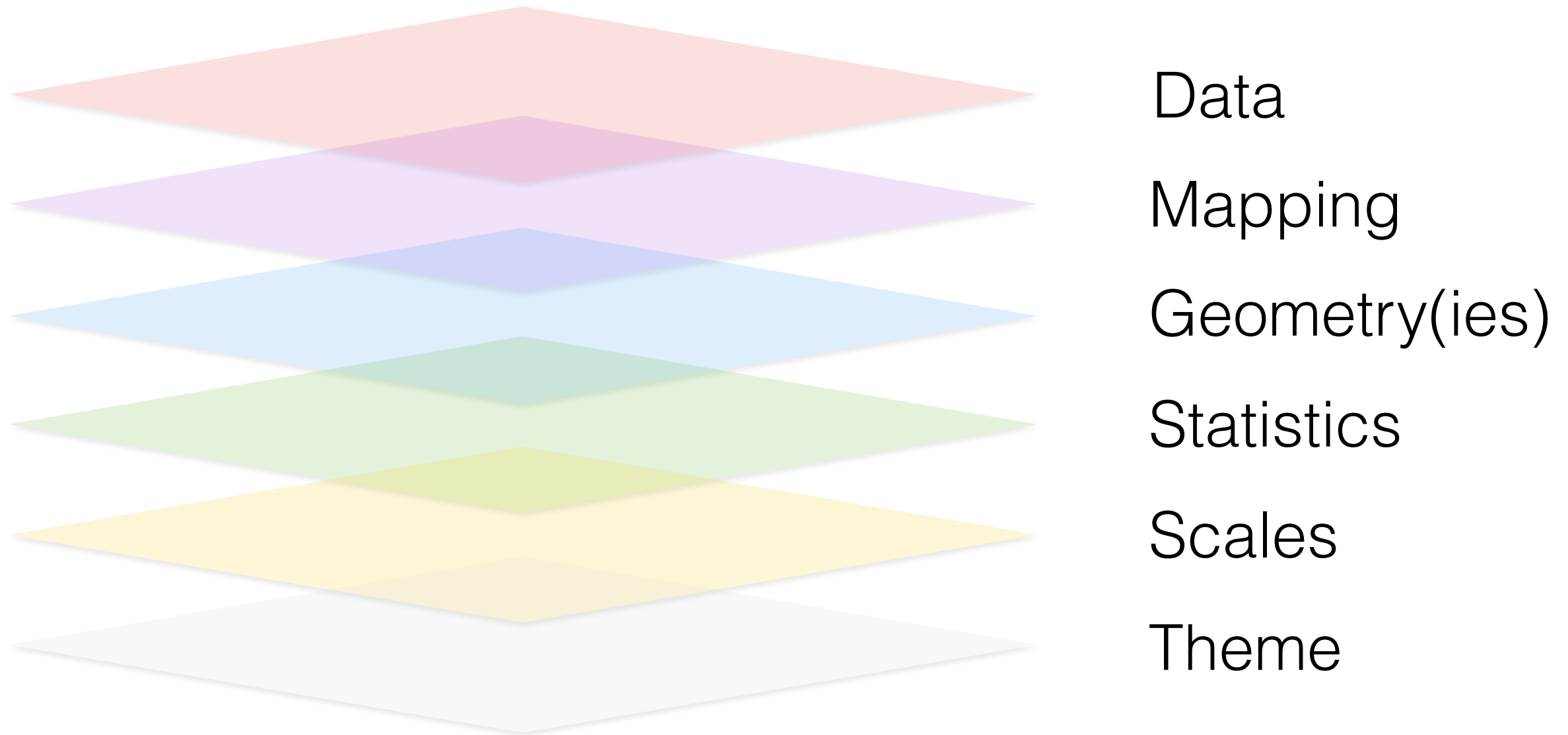
Data

Mapping

Geometry(ies)

Statistics

Layering in ggplot



How does it translate to R?

1. Create a plot object

```
myplot <- ggplot(data, aes(x,y))
```

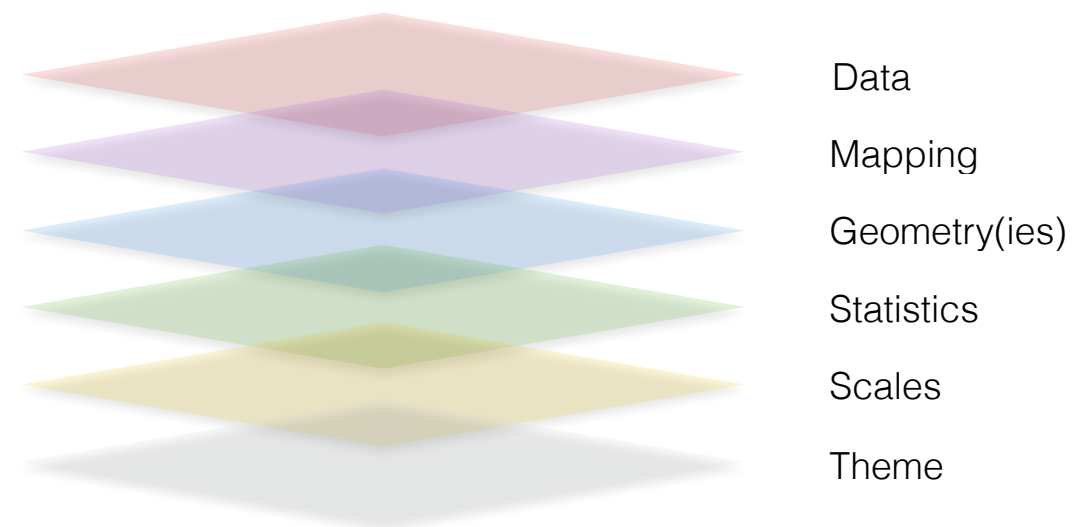
2. Add layers to it

```
myplot <- myplot + layer()
```

3. Repeat step 2 until you have all the layers that you need

4. Print your plot

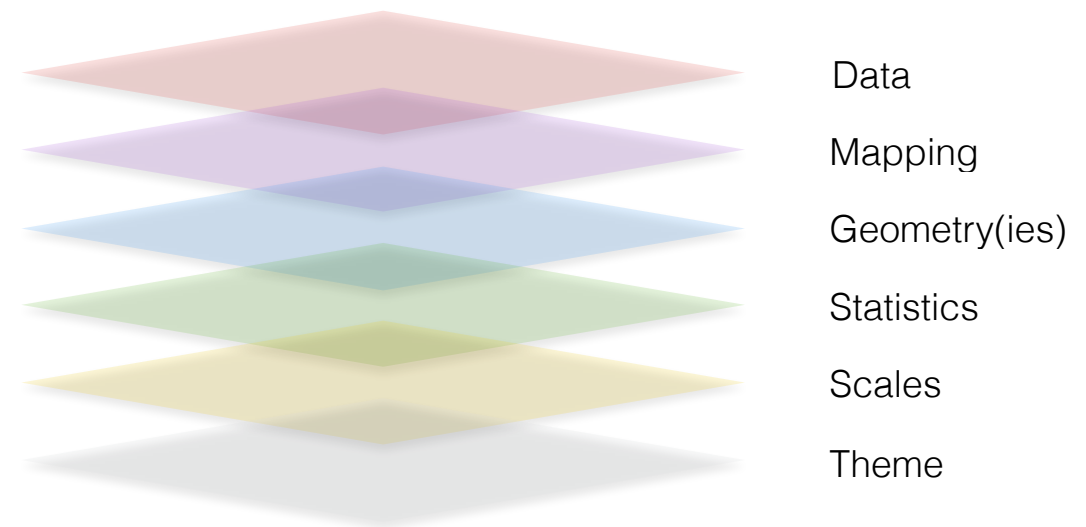
```
print(myplot)
```



Plotting with ggplot()

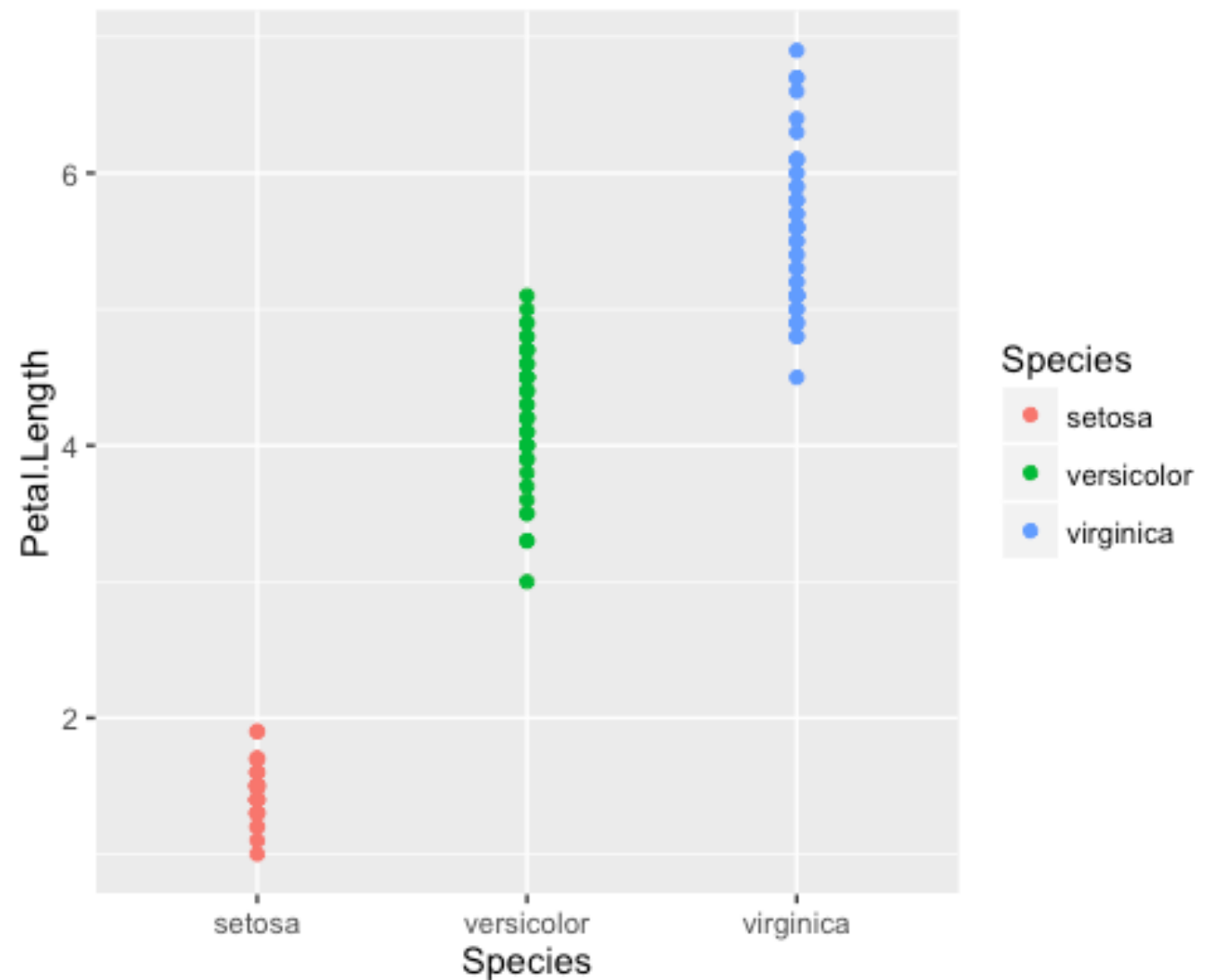
```
> p <- ggplot(data = iris,  
              aes(y = Petal.Length,  
                  x = Species,  
                  colour = Species))
```

```
> p + geom_point()
```



Plotting with ggplot()

```
> p <- ggplot(data = iris,  
  aes(y = Petal.Length,  
    x = Species,  
    colour = Species))  
  
> p + geom_point()
```

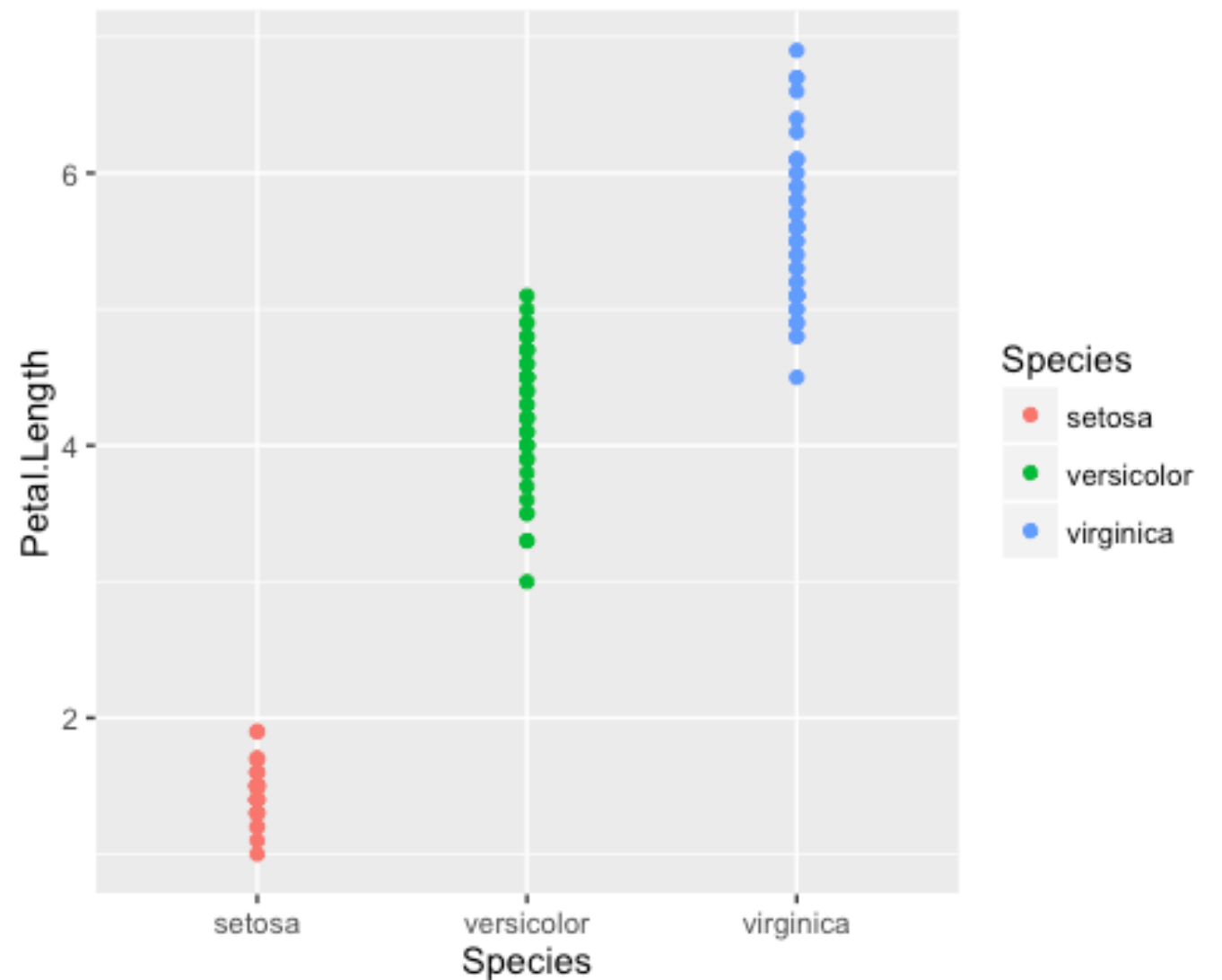


Plotting with ggplot()

```
> p <- ggplot(data = iris,
```

```
  aes(y = Petal.Length,  
      x = Species,  
      colour = Species)
```

```
> p + geom_point()
```



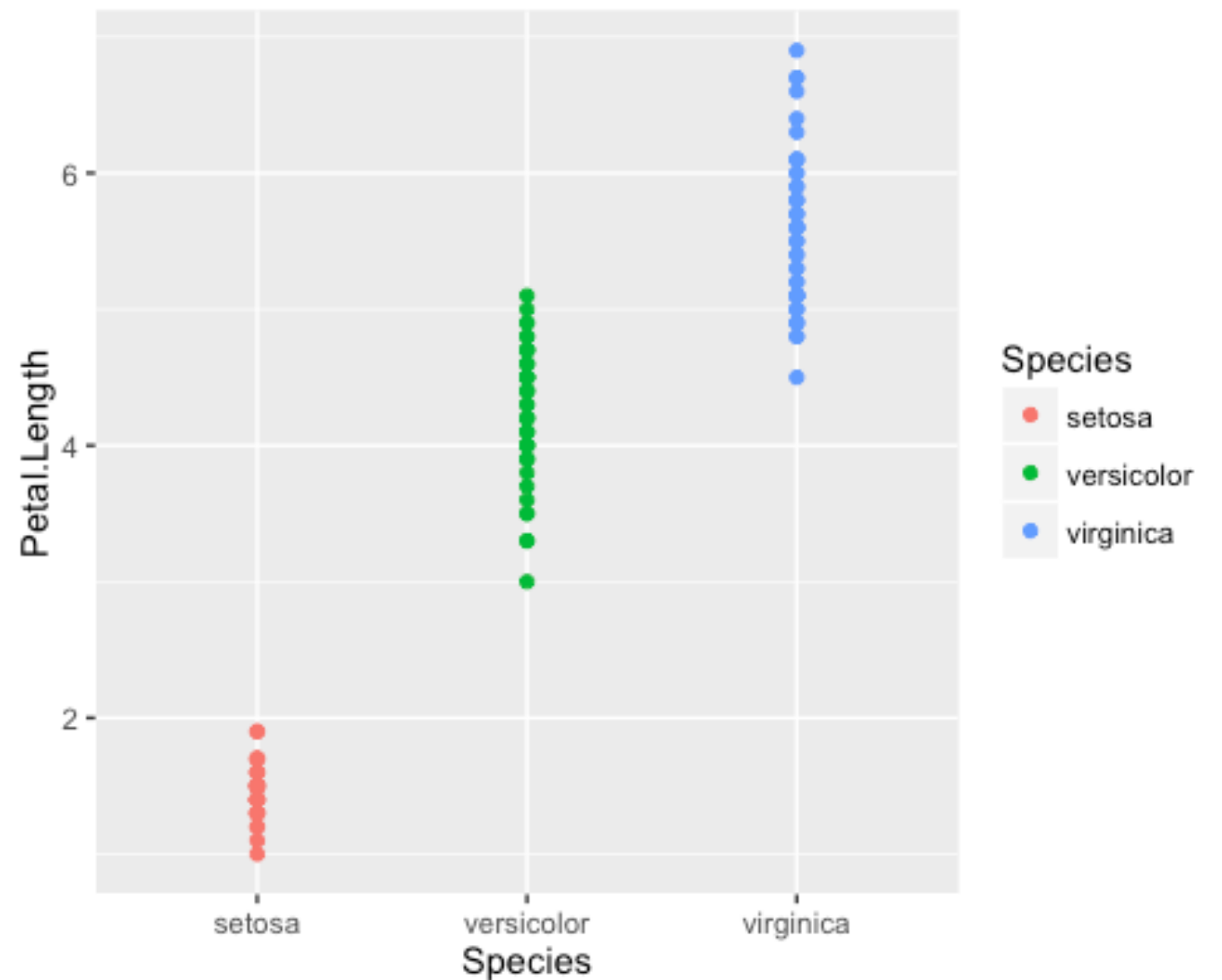
Aesthetics (mapping)

- x, y: positions of your data
- colour: colour of your points (or each point)
- group: assign groups to each point (e.g. treatment)
- shape: the form of the point
- linetype: the type of line to be used
- size: size of point or line
- alpha: transparency of the points

Plotting with ggplot()

```
> p <- ggplot(data = iris,  
              aes(y = Petal.Length,  
                  x = Species,  
                  colour = Species))
```

```
> p + geom_point()
```

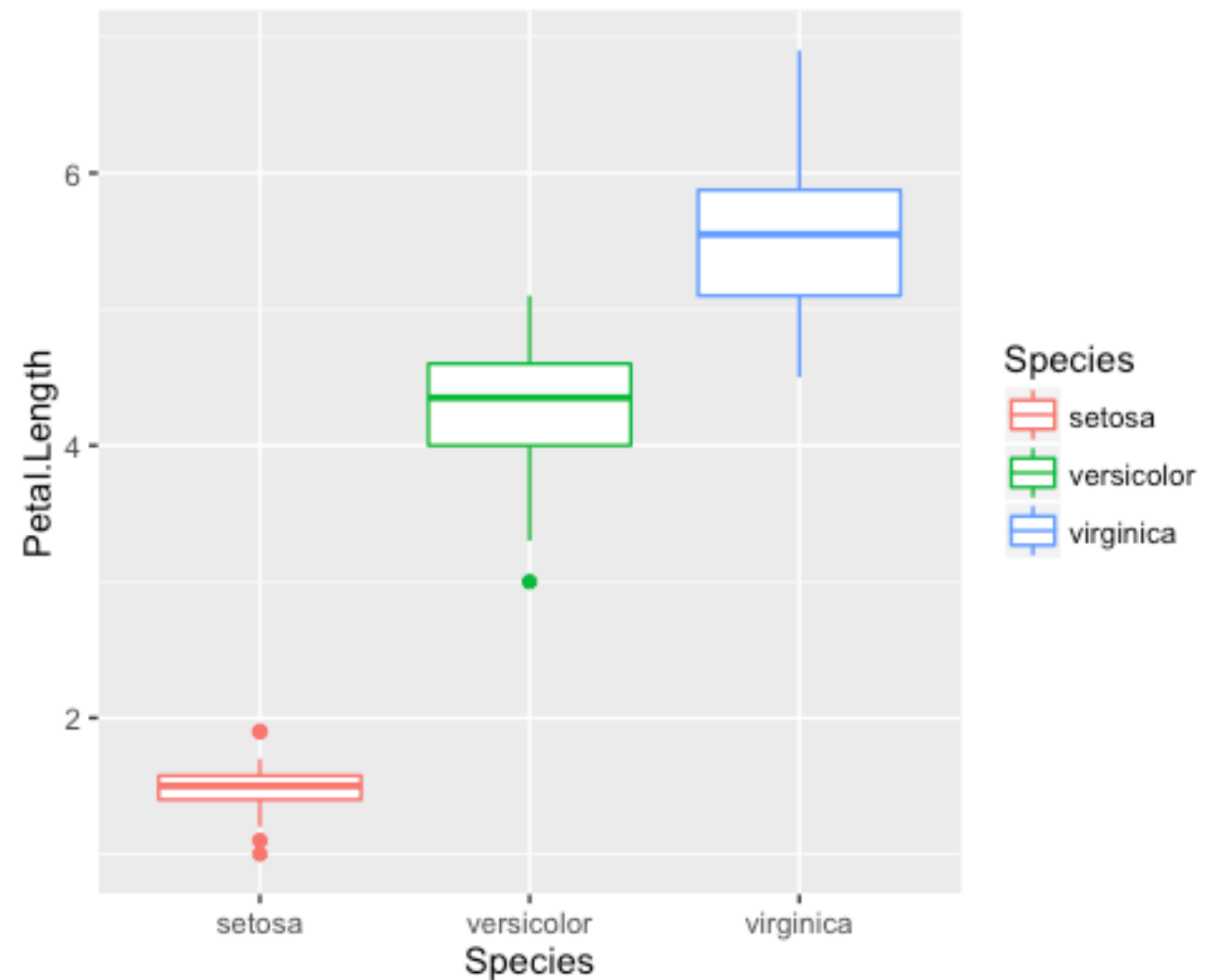


Geometry (geom_XX)

- points: scatterplot
- line: line plot (line increases with x)
- path: line plot, where line follows order of observations
- boxplot: box- and whisker-plots, very useful for categorical data on the y axis
- bar: barplots (but we rarely like those!)
- histogram: histograms (when you have only one variable)

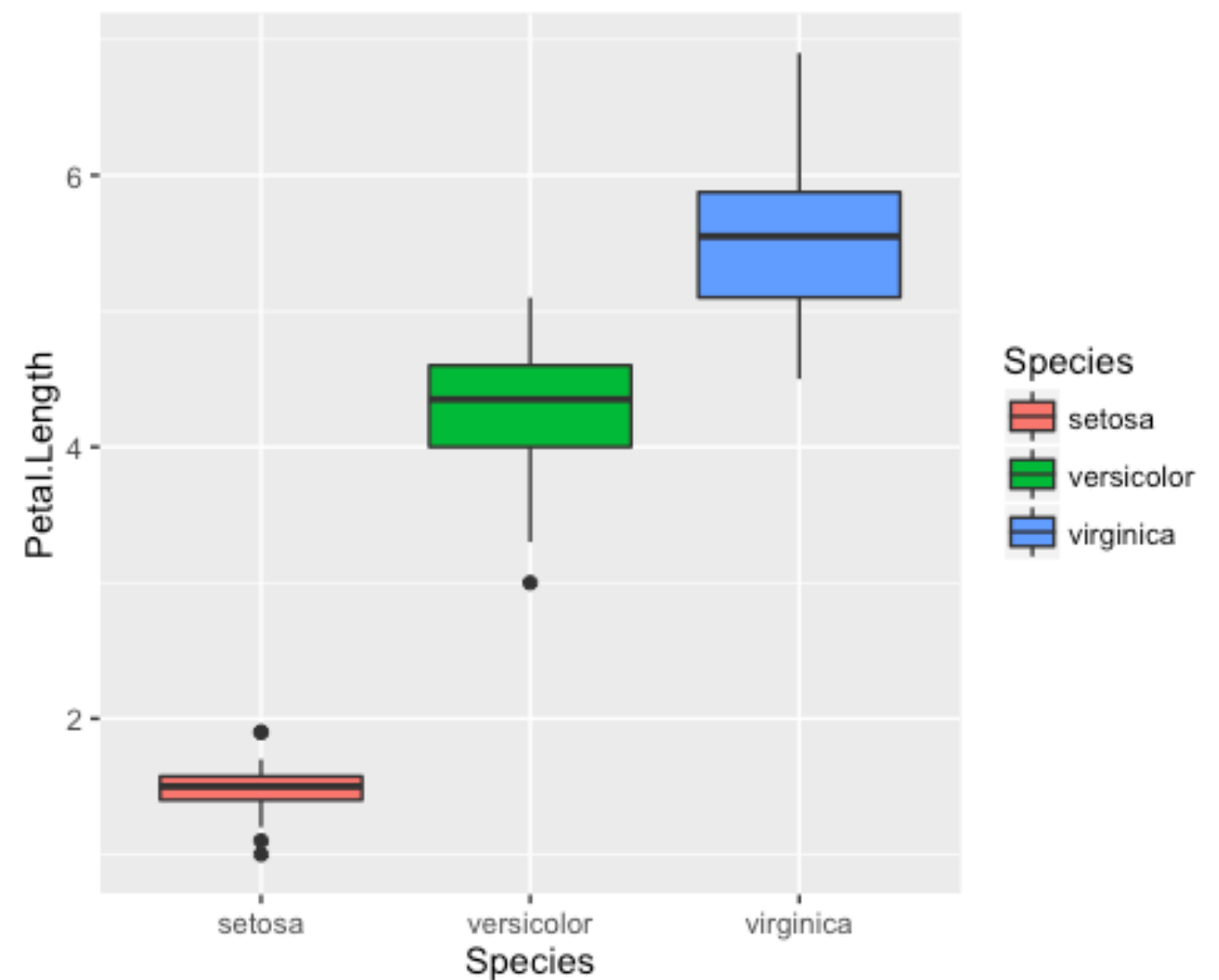
Plotting with ggplot()

```
> p <- ggplot(data = iris,  
  aes(y = Petal.Length,  
    x = Species,  
    colour = Species))  
  
> p + geom_boxplot()
```



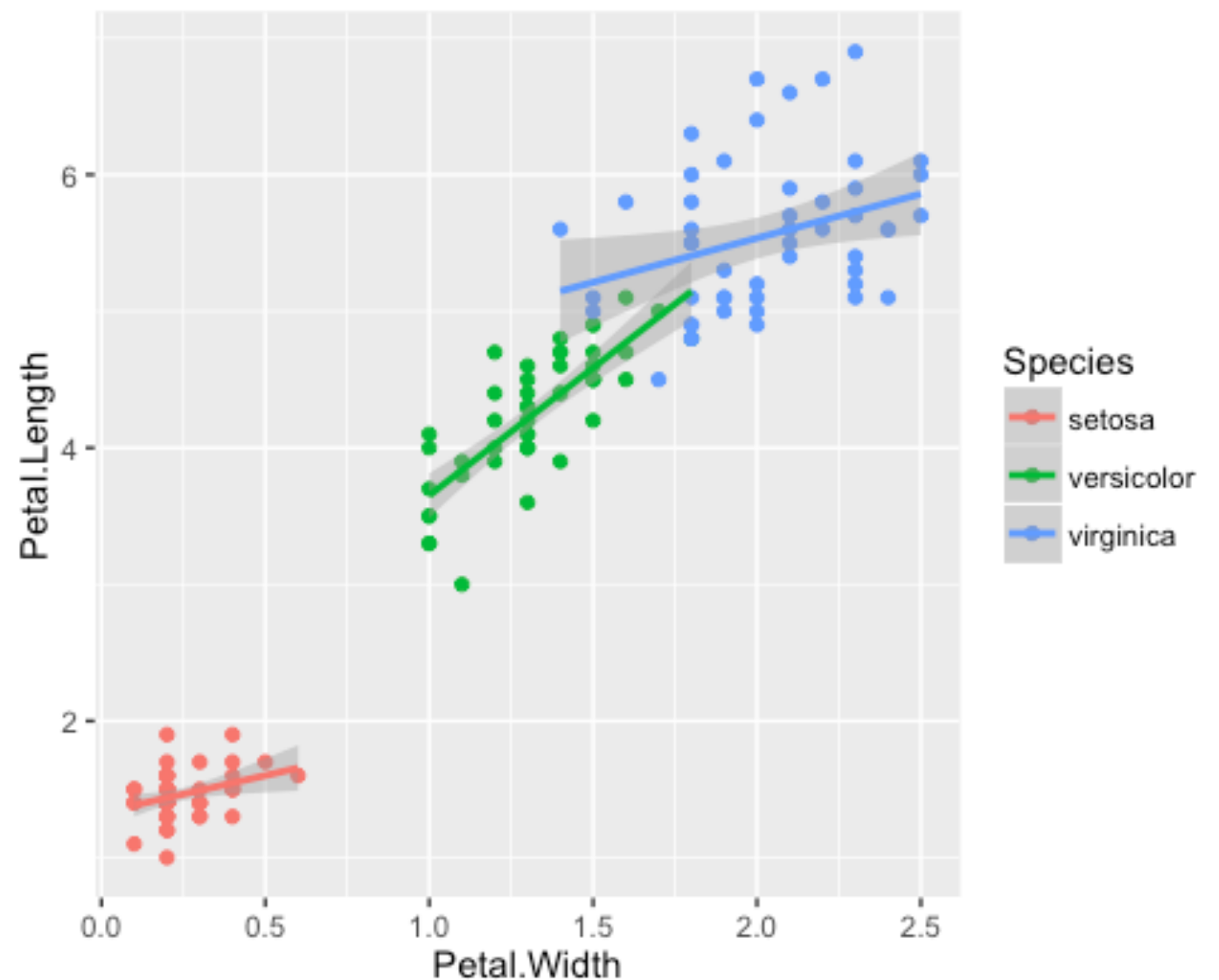
Plotting with ggplot()

```
> p <- ggplot(data = iris,  
  aes(y = Petal.Length,  
    x = Species,  
    fill = Species))  
  
> p + geom_boxplot()
```



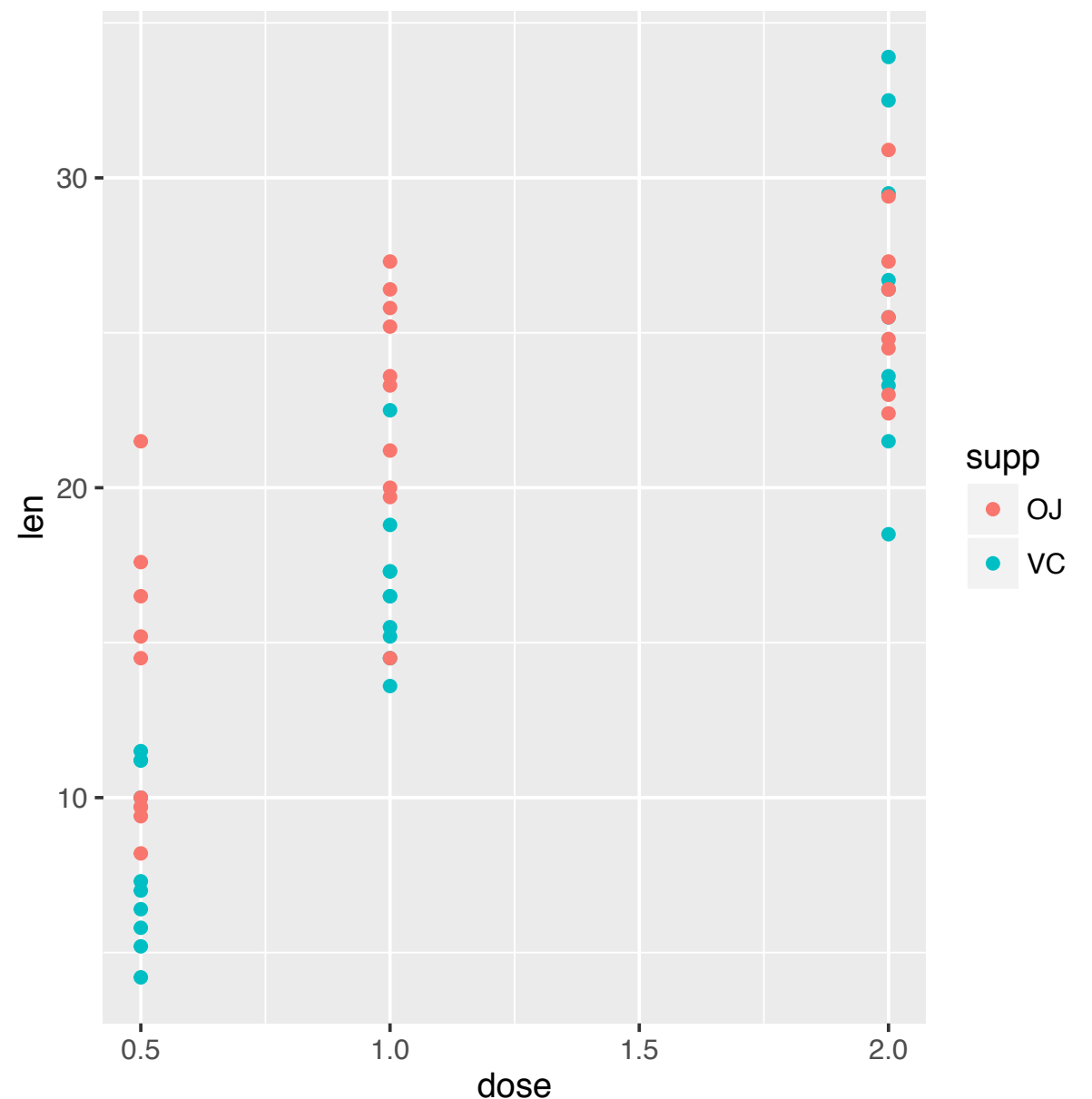
Using statistics

```
> p <- ggplot(data = iris,  
  aes(y = Petal.Length,  
    x = Petal.Width,  
    colour = Species))  
  
> p + geom_point() +  
  stat_smooth(method = "lm")
```



Try it yourself with ggplot()!

- Can you make this plot look even better using the command `ggplot()`.
- What layers would you choose and why?

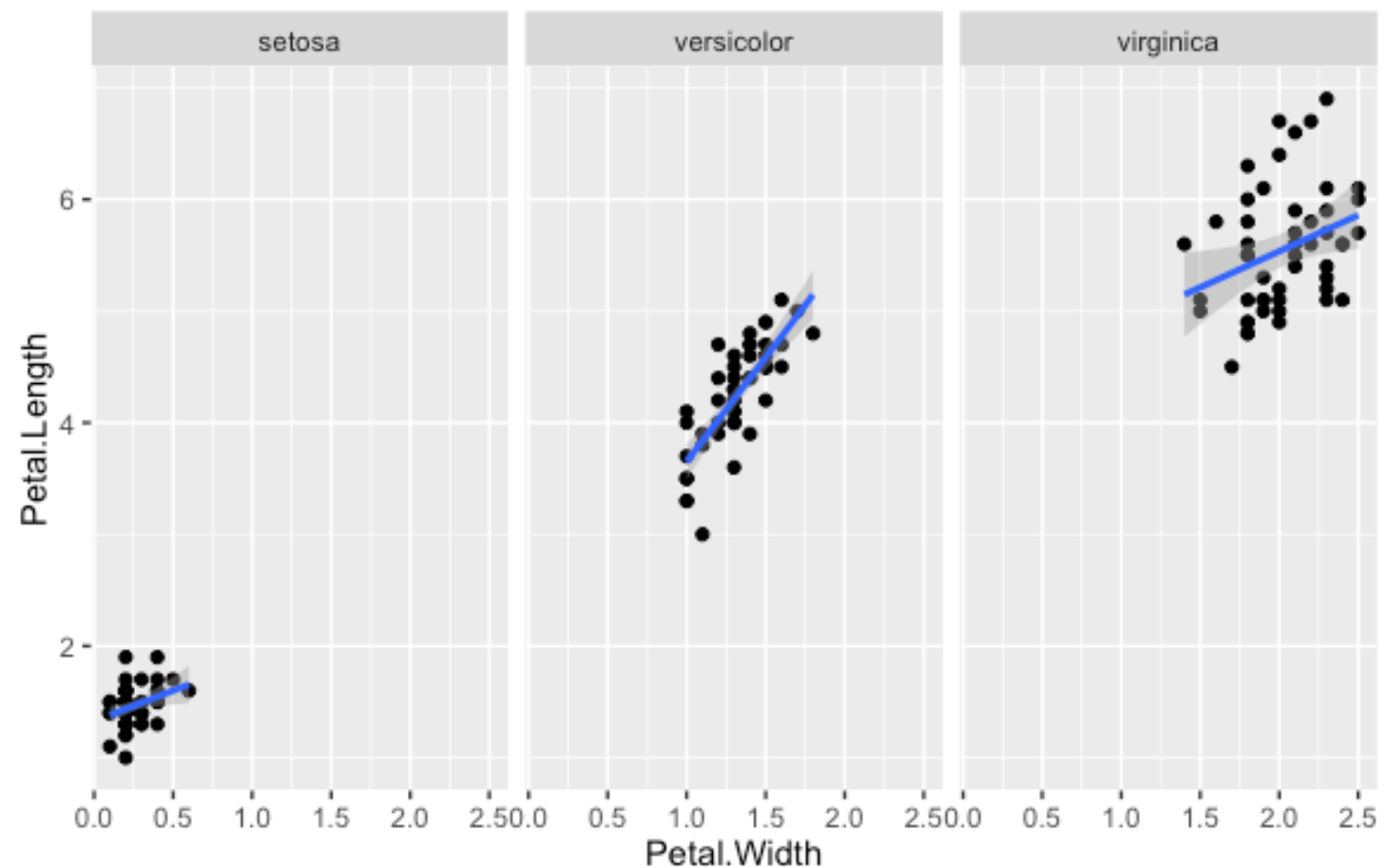


Facets

- Used to create a multipanel plot based on some grouping variable.
- Notation: `face_grid(rows~columns)`
- If you just want columns, then `~columns`, and vice versa.

Let's use facets to see things better

```
> p <- ggplot(data = iris,  
  aes(y = Petal.Length,  
    x = Petal.Width))  
  
p + geom_point() +  
  stat_smooth(method = "lm") +  
  facet_grid(.~Species)
```



Final retouches: scale_XX

Scales are the equivalent of make-up on your plot:

- Change your axis (e.g. `scale_x_continuous`)
- Labels (e.g. `labs`)
- Colours (e.g. `scale_colour_manual`)
- and lot more!

Ok, Enough! I know you are about to get a stroke! (me too)

