

Plotting continuous data in R with ggplot2

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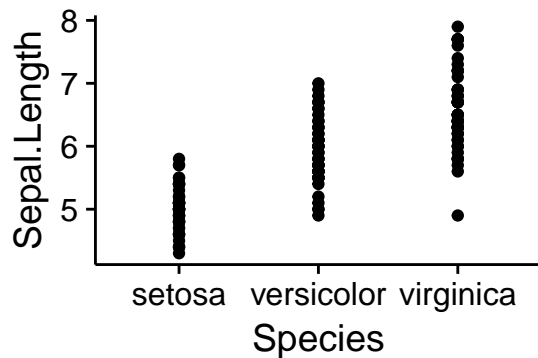
Introduction to ggplot using qplot

- We're going to plot some data using the qplot command
- We'll need to have 2 packages loaded
- ggplot2, which has the function qplot()
- cowplot, which provides some nice defaults
- We'll use the iris dataset that comes with R

A basic plot in ggplot using qplot()

- Unless told otherwise, qplot plots dots

```
qplot(y = Sepal.Length,  
      x = Species,  
      data = iris)
```

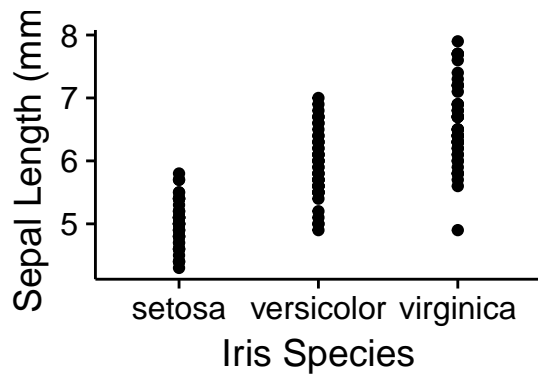


Box plot with labels

- R will usually generate labels for the x and y axes based on the command.
- These can be changed by adding another command after the qplot() command
- Add the command + **xlab**("...") sets the labels for the x-axis, + **ylab**("...") for the y axis.
- Text for the labels goes in quotes (ie, "Iris species").
- The use of the "+" is different than for most other R packages
- Forgetting the quotes will cause the code to fail.
- Note that units (mm) are included for the y axis.

```
qplot(y = Sepal.Length,  
      x = Species,  
      data = iris) +
```

```
xlab("Iris Species") +
ylab ("Sepal Length (mm)" )
```

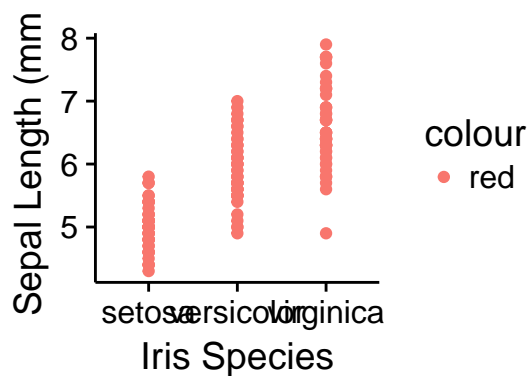


Changing colors in R plots

Changing colors in R plots part 1

- If we wanted we could change the color of the dots using the argument “**col =**”. This code can be used to change the color of most types of plots in R.
- This doesn’t increase the information content of the figure but maybe makes it nicer to look at.

```
qplot(y = Sepal.Length,
      x = Species,
      data = iris,
      color = "red") +
xlab("Iris Species") +
ylab ("Sepal Length (mm)" )
```



Changing colors in R plots part 2

```
#dopt w/color changes
qplot(y = Sepal.Length,
      x = Species,
      data = iris,
      color = Species) +
```

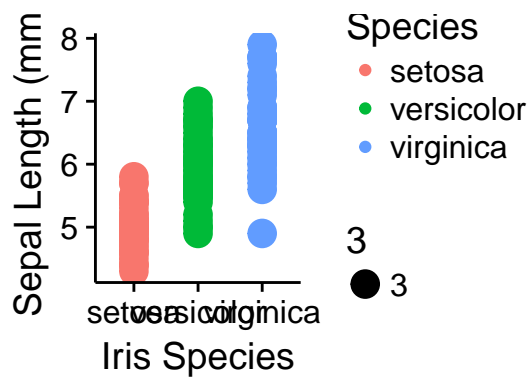
```
xlab("Iris Species") +
ylab ("Sepal Length (mm)")
```



Tweaking plots: changing the point size

Run the code below, Can you see what changed?

```
#dopt w/color changes
qplot(y = Sepal.Length,
      x = Species,
      data = iris,
      color = Species,
      size = 3) +
xlab("Iris Species") +
ylab ("Sepal Length (mm)")
```

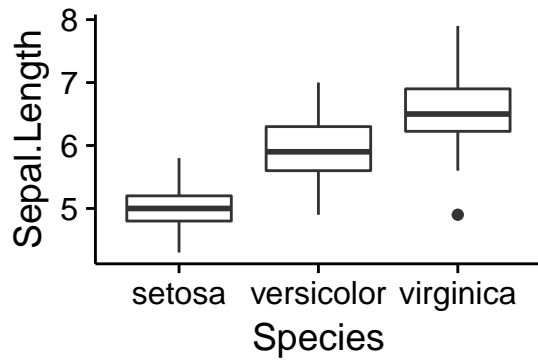


Boxplot with qplot

Basic boxplot with qplot

- note use of argument “geom = ...”

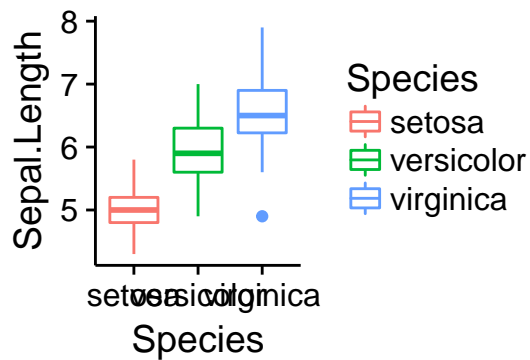
```
qplot(y = Sepal.Length,
      x = Species,
      data = iris,
      geom = "boxplot")
```



Basic boxplot with colors

- same as before, using “color =”

```
#dopt w/color changes
qplot(y = Sepal.Length,
      x = Species,
      data = iris,
      geom = "boxplot",
      color = Species)
```



Basic boxplot lables

- now use + xlab() and + ylab()

```
qplot(y = Sepal.Length,
      x = Species,
      data = iris,
      geom = "boxplot",
      color = Species) +
xlab("Iris Species") +
ylab("Sepal Length (mm)")
```



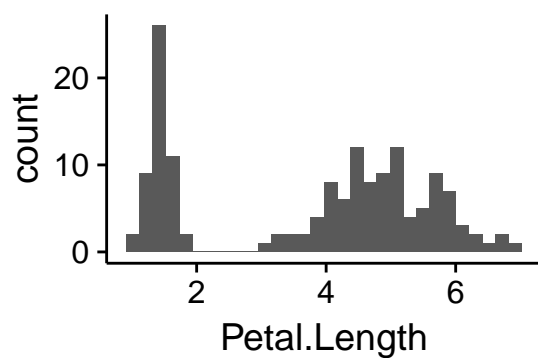
Histograms using qplot

- made with `geom = "histogram"` argument
- very very easy to make in R with `ggplot`
- very very very hard to make in Excel
- You should make them all the time for you data!

Histograms of iris data

- This code makes a histogram of one of the iris species' `Petal.Length`.
- Note that you don't have "`y =`" or "`x =`" for a histogram!

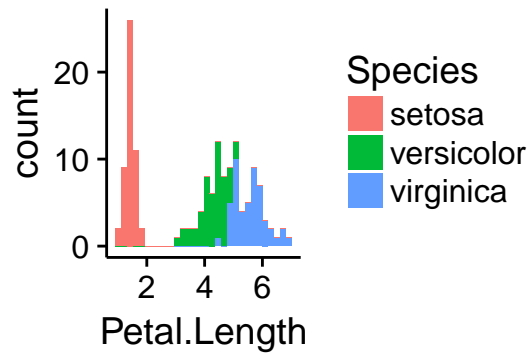
```
qplot(Petal.Length,
      data = iris)
```



Histogram with colors

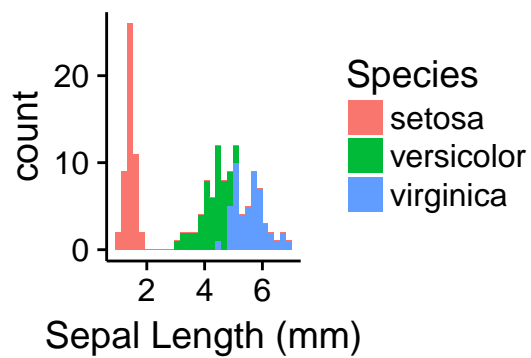
What does this show?

```
qplot(Petal.Length,
      data = iris,
      fill = Species)
```



Histogram with axes labels

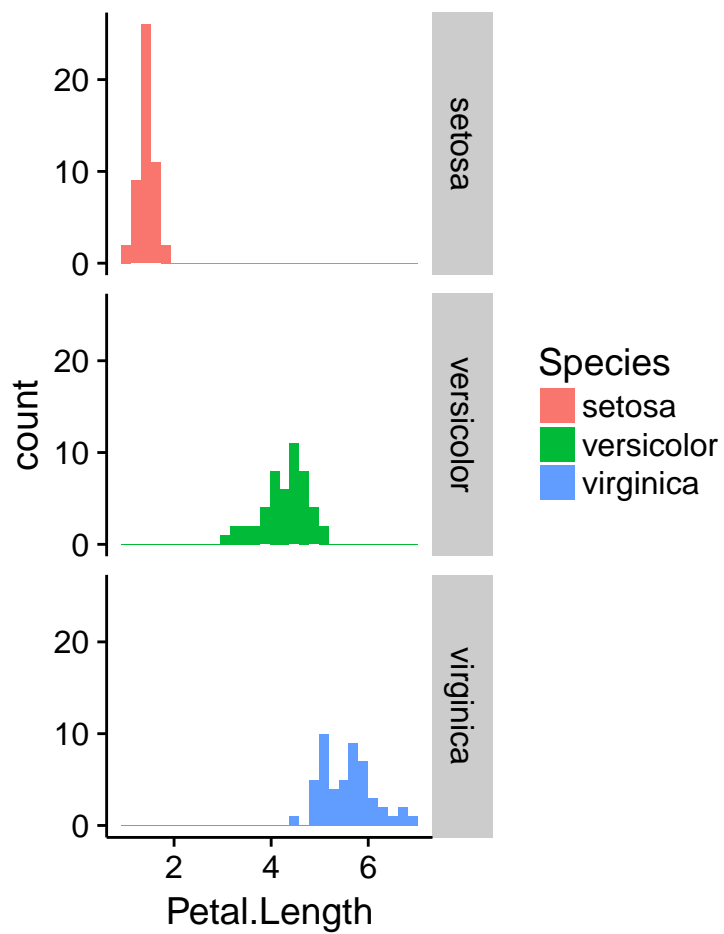
```
qplot(Petal.Length,
      data = iris,
      fill = Species) +
  xlab ("Sepal Length (mm)")
```



Multiple histograms: “Facets”

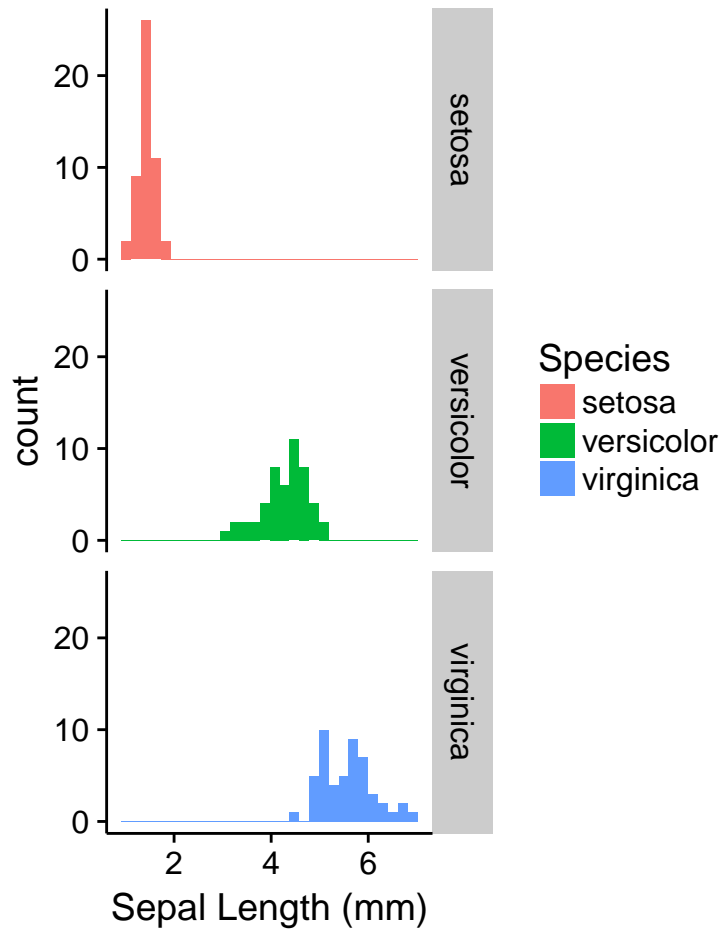
What does this show?

```
qplot(Petal.Length,
      data = iris,
      fill = Species,
      facets = Species ~.)
```



Add a label to x-axis

```
qplot(Petal.Length,  
      data = iris,  
      fill = Species,  
      facets = Species ~.) +  
xlab("Sepal Length (mm)")
```



Modifying histograms: titles with the main = argument

- Titles are good for your own personal use but actually are almost never used in figures published in papers and books.
- We can add a title like this using the argument “main =”

```
qplot(Petal.Length,  
      data = iris,  
      fill = Species,  
      main = "Iris species histograms",  
      facets = Species ~.) +  
xlab ("Sepal Length (mm)")
```