Data Visualization with R & ggplot2

Karthik Ram

April 8, 2013

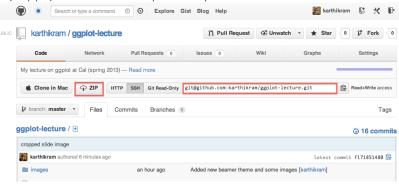
Some housekeeping

Install some packages

```
install.packages("ggplot2", dependencies = TRUE)
```

Download this PDF

github.com/karthikram/ggplot-lecture https://speakerdeck.com/karthik/



Base graphics

- Ugly, laborious, and verbose
- There are better ways to describe statistical visualizations.

Why ggplot2?

- Follows a grammar, just like any language.
- It defines basic components that make up a sentence. In this case, the grammar defines components in a plot.
- Grammar of graphics originally coined by Lee Wilkinson

Why ggplot2?

- Supports a continuum of expertise.
- Get started right away but with practice you can effortless build complex, publication quality figures.

Basics

Some terminology

- ggplot The main function where you specify the dataset and variables to plot
- geoms geometric objects
 - geom_point(), geom_bar(), geom_density(), geom_line(), geom_area()
- aes aesthetics
 - shape, transparency, color, fill.
- scales Define how your data will be plotted
 - continuous, discrete, log

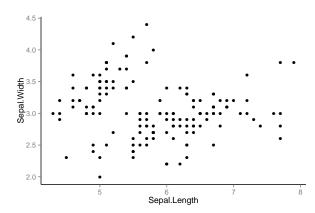
Assembling your first ggplot

The iris dataset

```
head(iris)
    Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##
## 1
             5.1
                         3.5
                                      1.4
                                                 0.2 setosa
## 2
             4.9
                         3.0
                                      1.4
                                                 0.2 setosa
## 3
             4.7
                         3.2
                                     1.3
                                                 0.2 setosa
## 4
             4.6
                         3.1
                                     1.5
                                                 0.2 setosa
## 5
            5.0
                         3.6
                                     1.4
                                                 0.2 setosa
## 6
             5.4
                         3.9
                                      1.7
                                                 0.4 setosa
```

Let's try an example

```
ggplot(data = iris, aes(x = Sepal.Length, y = Sepal.Width)) +
geom_point()
```



Basic structure

```
ggplot(data = iris, aes(x= Sepal.Length, y= Sepal.Width))
    + geom_point()
myplot <- ggplot(data = iris, aes(x = Sepal.Length, y = Sepal.Width))
myplot + geom_point()</pre>
```

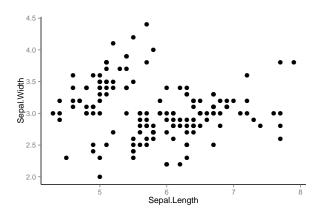
- Specify the data and variables inside the ggplot function.
- Anything else that goes in here becomes a global setting.
- Then add layers of geometric objects, statistical models, and panels.

Quick note

- Never use qplot short for quick plot.
- You'll end up unlearning and relearning a good bit.

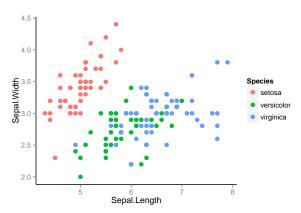
Increase the size of points

```
ggplot(data = iris, aes(x = Sepal.Length, y = Sepal.Width)) +
geom_point(size = 3)
```



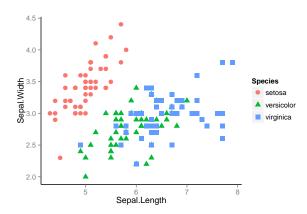
Add some color

```
ggplot(iris, aes(Sepal.Length, Sepal.Width, color = Species)) +
geom_point(size = 3)
```



Differentiate points by shape

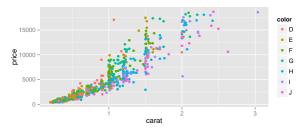
```
ggplot(iris, aes(Sepal.Length, Sepal.Width, color = Species)) +
geom_point(aes(shape = Species), size = 3)
```



Exercise 1

```
# Make a small sample of the diamonds dataset
d2 <- diamonds[sample(1:dim(diamonds)[1], 1000), ]</pre>
```

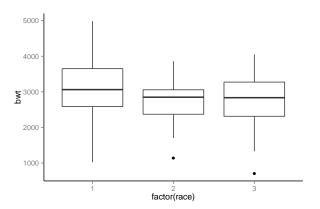
Then generate this plot below.



Box plots

See ?geom_boxplot for list of options

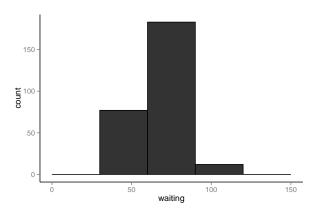
```
library(MASS)
ggplot(birthwt, aes(factor(race), bwt)) + geom_boxplot()
```



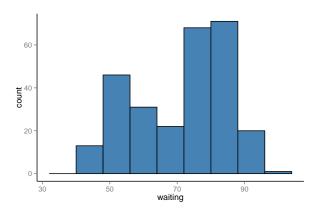
Histograms

See ?geom_histogram for list of options

```
h <- ggplot(faithful, aes(x = waiting))
h + geom_histogram(binwidth = 30, colour = "black")</pre>
```

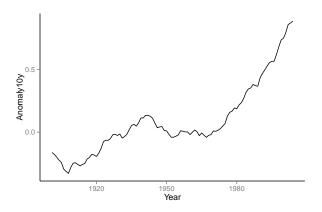


```
h <- ggplot(faithful, aes(x = waiting))
h + geom_histogram(binwidth = 8, fill = "steelblue", colour = "black")</pre>
```



Line plots

```
climate <- read.csv("climate.csv", header = T)
ggplot(climate, aes(Year, Anomaly10y)) +
geom_line()</pre>
```

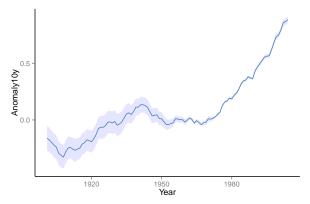


climate <- read.csv(text =

RCurl::getURL(https://raw.github.com/karthikram/ggplot-lecture/master/climate.csv))

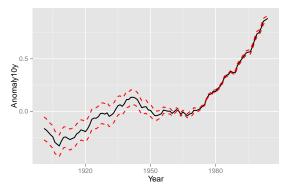
We can also plot confidence regions

```
ggplot(climate, aes(Year, Anomaly10y)) +
geom_ribbon(aes(ymin = Anomaly10y - Unc10y,
ymax = Anomaly10y + Unc10y),
fill = "blue", alpha = .1) +
geom_line(color = "steelblue")
```



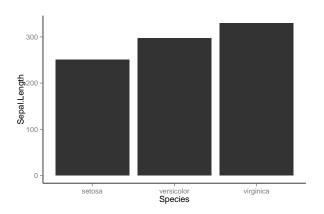
Exercise 2

• Modify the previous plot and change it such that there are three lines instead of one with a confidence band.

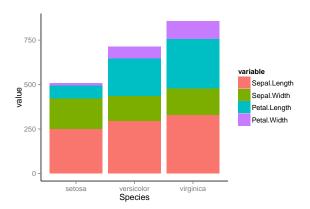


Bar plots

```
ggplot(iris, aes(Species, Sepal.Length)) +
geom_bar(stat = "identity")
```



```
df <- melt(iris, id.vars = "Species")
ggplot(df, aes(Species, value, fill = variable)) +
geom_bar(stat = "identity")</pre>
```



plyr and reshape are key for using R

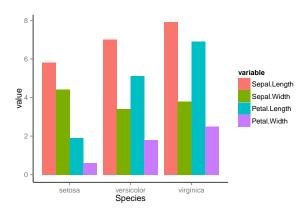
plyr and reshape

These two packages are the swiss army knives of R.

- plyr
 - ddply
 - Ilply
 - join
- reshape.
 - 1 melt
 - dcast
 - 3 acast

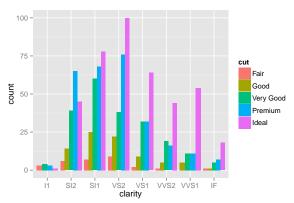
```
iris[1:2, ]
    Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##
## 1
        5.1
                      3.5 1.4 0.2 setosa
                      3.0
                                 1.4 0.2 setosa
## 2
           4.9
df <- melt(iris, id.vars = "Species")</pre>
df[1:2, ]
    Species variable value
##
## 1 setosa Sepal.Length 5.1
## 2 setosa Sepal.Length 4.9
```

```
ggplot(df, aes(Species, value, fill = variable)) +
geom_bar(stat = "identity", position = "dodge")
```



Exercise 3

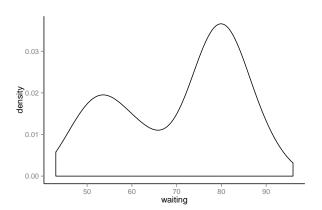
Using the d2 dataset you created earlier, generate this plot below. Take a quick look at the data first to see if it needs to be binned.



Density Plots

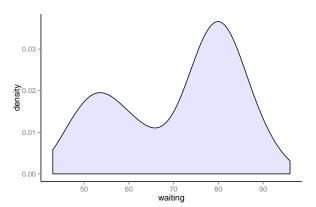
Density plots

ggplot(faithful, aes(waiting)) + geom_density()

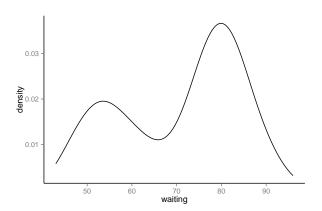


Density plots

```
ggplot(faithful, aes(waiting)) +
geom_density(fill = "blue", alpha = 0.1)
```



```
ggplot(faithful, aes(waiting)) +
geom_line(stat = "density")
```



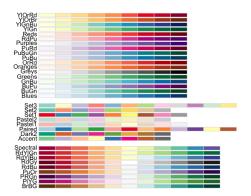
Mapping Variables to colors

Colors

```
aes(color = variable)
aes(color = "black")
# Or add it as a scale
scale_fill_manual(values = c("color1", "color2"))
```

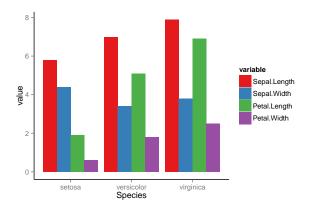
The RColorBrewer package

```
library(RColorBrewer)
display.brewer.all()
```



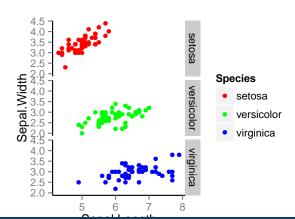
Using a color brewer palette

```
df <- melt(iris, id.vars = "Species")
ggplot(df, aes(Species, value, fill = variable)) +
geom_bar(stat = "identity", position = "dodge") +
scale_fill_brewer(palette = "Set1")</pre>
```



Manual color scale

```
ggplot(iris, aes(Sepal.Length, Sepal.Width, color = Species)) +
geom_point() +
facet_grid(Species ~ .) +
scale_color_manual(values = c("red", "green", "blue"))
```



Refer to a color chart for beautful visualizations

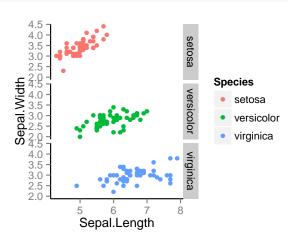
http://tools.medialab.sciences-po.fr/iwanthue/



Faceting

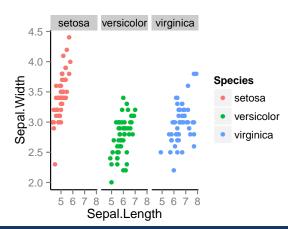
Faceting along columns

```
ggplot(iris, aes(Sepal.Length, Sepal.Width, color = Species)) +
geom_point() +
facet_grid(Species ~ .)
```



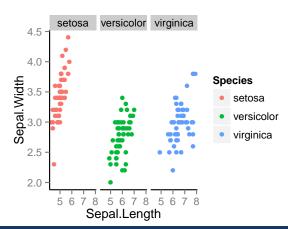
and along rows

```
ggplot(iris, aes(Sepal.Length, Sepal.Width, color = Species)) +
geom_point() +
facet_grid(. ~ Species)
```



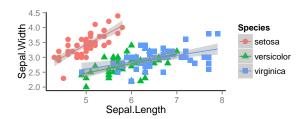
or just wrap your panels

```
ggplot(iris, aes(Sepal.Length, Sepal.Width, color = Species)) +
geom_point() +
facet_wrap( ~ Species)
```

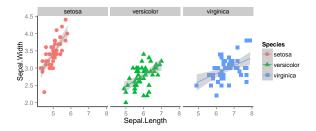


Adding smoothers

```
ggplot(iris, aes(Sepal.Length, Sepal.Width, color = Species)) +
geom_point(aes(shape = Species), size = 3) +
geom_smooth(method = "lm")
```



```
ggplot(iris, aes(Sepal.Length, Sepal.Width, color = Species)) +
geom_point(aes(shape = Species), size = 3) +
geom_smooth(method = "lm") +
facet_grid(. ~ Species)
```



Themes

Adding themes

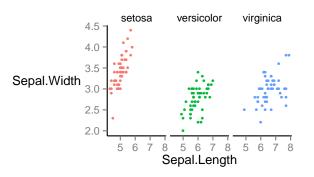
Themes are a great way to define custom plots.

```
theme()
# see ?theme() for more options
```

A themed plot

```
ggplot(iris, aes(Sepal.Length, Sepal.Width, color = Species)) +
geom_point(size = 1.2, shape = 16) +
facet_wrap( ~ Species) +
theme(legend.key = element_rect(fill = NA),
legend.position = "bottom",
strip.background = element_rect(fill = NA),
axis.title.y = element_text(angle = 0))
```

Adding themes



Species • setosa • versicolor • virginic

ggthemes library

```
install.packages("ggthemes")
library(ggthemes)
# Then add one of these themes to your plot
+theme_stata()
+theme_excel()
+theme_wsj()
+theme_solarized()
```

Create functions to automate your plotting

Write functions for day to day plots

```
my_custom_plot <- function(df) {
    ggplot(df, ...)
}</pre>
```

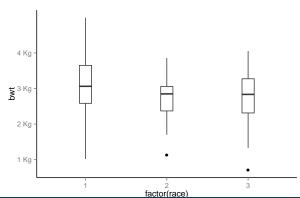
Scales

Commonly used scales

```
scale_fill_discrete(), scale_colour_discrete()
scale_fill_hue(), scale_color_hue()
scale_fill_manual(), scale_color_manual()
scale_fill_brewer(), scale_color_brewer()
scale_linetype(), scale_shape_manual()
```

Adding a continuous scale

```
library(MASS)
ggplot(birthwt, aes(factor(race), bwt)) +
geom_boxplot(width = .2) +
scale_y_continuous(labels = (paste0(1:4, " Kg")),
breaks = seq(1000, 4000, by = 1000))
```



Another example

```
# Assign the plot to an object
dd <- ggplot(iris, aes(Sepal.Length, Sepal.Width, color = Species)) +
geom_point(size = 4, shape = 16) +
facet_grid(. ~Species)
# Now add a scale
dd +
scale_y_continuous(breaks = seq(2, 8, by = 1),
labels = paste0(2:8, " cm"))</pre>
```

Publication quality figures

• If the plot is on your screen

```
ggsave("~/path/to/figure/filename.png")
```

• If your plot is assigned to an object

```
ggsave(plot1, file = "~/path/to/figure/filename.png")
```

Specify a size

```
ggsave(file = "/path/to/figure/filename.png", width = 6,
height =4)
```

• or any format (pdf, png, eps, svg, jpg)

```
ggsave(file = "/path/to/figure/filename.eps")
ggsave(file = "/path/to/figure/filename.jpg")
ggsave(file = "/path/to/figure/filename.pdf")
```

Further help

- You've just scratched the surface with ggplot2.
- Practice
- Read the docs (either locally in R or at http://docs.ggplot2.org/current/)
- Work together

