















Richard Upton 4 August 2015





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Plotting is an important part of modeling

- For data exploration before modeling (graphical analysis).
- For assessing model performance (Visual Predictive Checks).
- For showing the predictions of a model (Simulation).
- For publishing modeling results.













Why use R for plotting?

- Of the statistical software environments, it's the most versatile.
- It's combination of data manipulation and plotting makes it a "one stop shop".
- It facilliates Reproducible Research and Open Science.













The base package

Pros - quick & easy

Cons - legends, scales and plotting by factor difficult

The lattice package

Pros - allows plotting by factor

Cons - fussy syntax, legends and scales difficult

The ggplot2 package

Pros - allows plotting by factor, automatic legends and scale Cons - slow for large datasets

The grid package

Low level drawing functions (lines, points, text)











Plotting devices in R

When R draws a plot, it is written to a device.

Plots can be sent to more than 1 device.

Plot appearance depends on device settings (e.g. dimensions, dpi).

The screen plot may look weird when a plot is formated for other devices.

Avaliable devices (?devices):

Screen (can copy to clipboard in Windows)

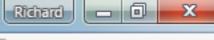
PDF file

PNG bitmap file

JPEG bitmap file

Typically, write to a file then import in Word or Powerpoint.















A Grammar of Graphics

Wilkinson (2005): A grammar for the components of a graphic:

- data and aesthetic mappings
- geometric objects
- scales
- facet specification
- statistical transformations
- coordinate system (usually cartersian for us)

Instead of "draw a scatterplot for these data".













ggplot2

ggplot2 - plotting in R based on A Grammar of Graphics. http://ggplot2.org (Hadley Wickham).

Most popular ploting in package in R (at the moment). The core functions are **qplot** and **ggplot**. Now forget about **qplot**.

The specifications for a graph and data are contained in a R object.

The plotting object can be built in layers.









Data and Aesthetic Mappings

Aesthetics for the plot of data x and y are mapped against two explanatory variables:

X	у	ID	TRT	aesthetic1	aesthetic2
0	2	Subject1	Baseline	red	circle
10	3	Subject1	Treated	red	triangle
0	4	Subject2	Baseline	blue	circle
		Subject2			triangle

Plot data (x, y) can be *Discrete* (a factor) or *Continuous*. Explanatory variables (ID, TRT) can be *Discrete* or Continuous.

There are aesthetics for: Colour, Shape, Size and Linetype.













Data and Aesthetic Mappings

```
> exampledata <- read.csv("data_aes.csv")
> exampledata
```

```
ΧΥ
1 0 2 Subject1 Baseline
2 10 3 Subject1 Treated
3 0 4 Subject2 Baseline
4 10 6 Subject2 Treated
```

```
> plotobj <- ggplot(exampledata)</pre>
> plotobj <- plotobj + geom_point(aes(x=X, y=Y, shape=TRT, colour=ID))</pre>
```





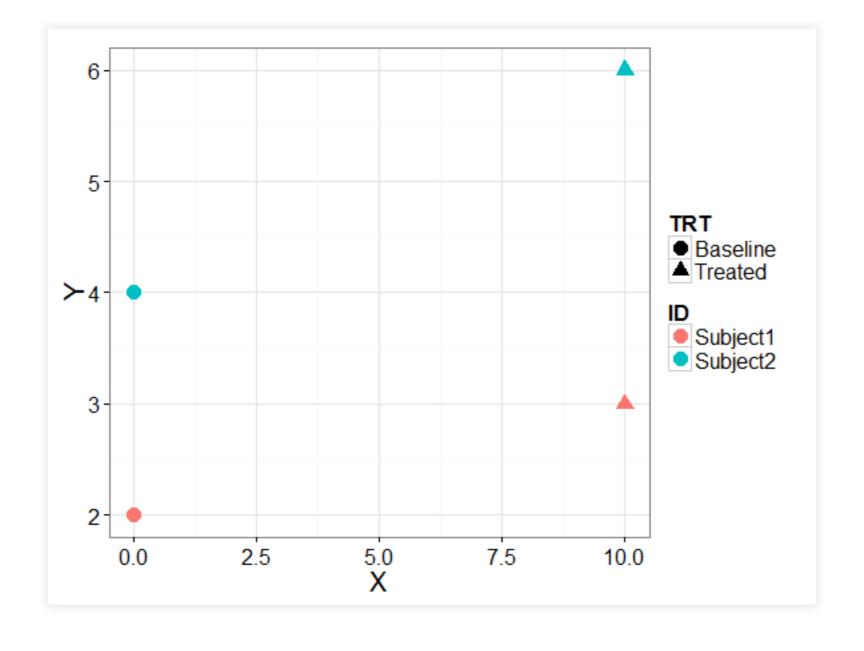








Data and Aesthetic Mappings













Geometric Objects

The plot of x and y can be made using a different geometries: There is a large toolkit of geometries This allows great flexibility in plotting Not all combinations of data and geometries are possible

- points
- lines
- boxplot
- ribbon
- errorbar
- step etc.











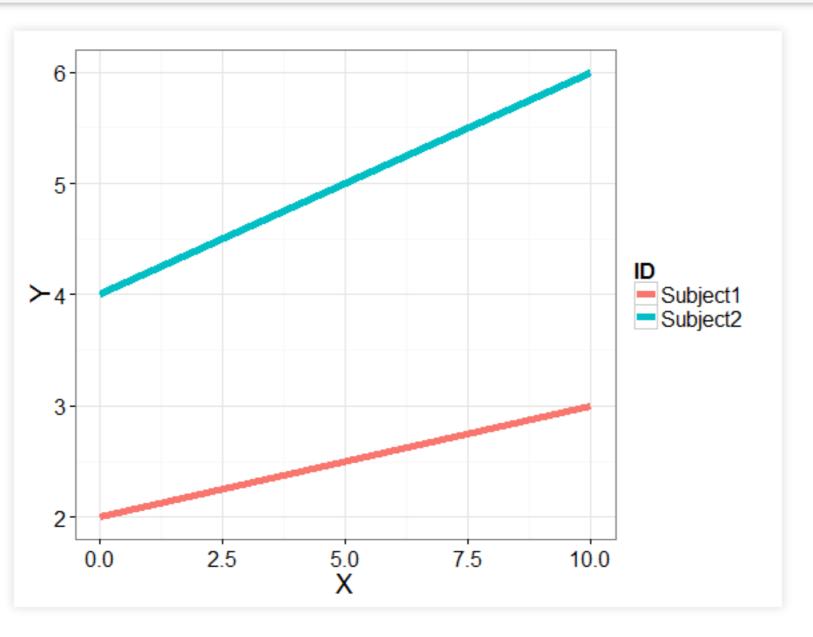


3

Geometric Objects

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```
> plotobj <- ggplot(exampledata)
> plotobj <- plotobj + geom_line(aes(x=X, y=Y, colour=ID))</pre>
```













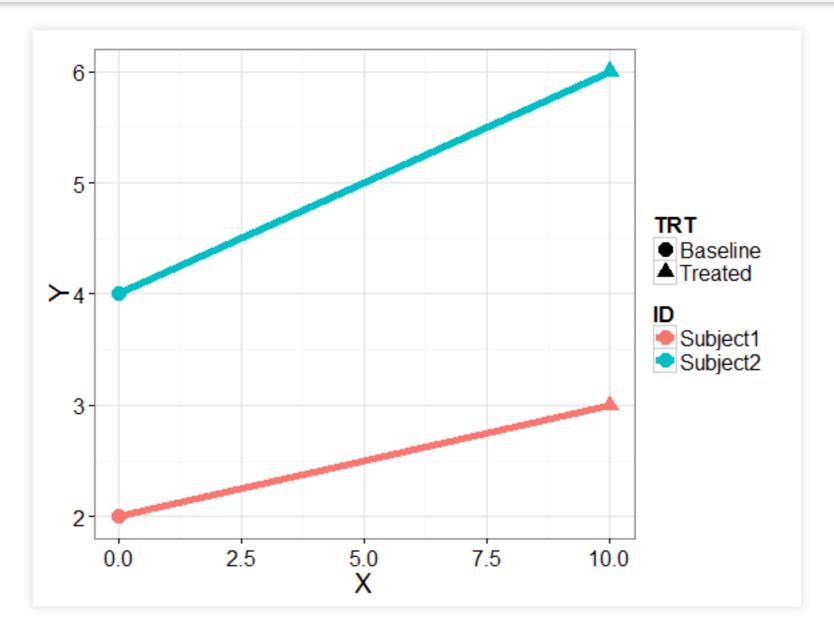




Plots can be built in layers

Two different geometries used in the same plot:

```
> plotobj <- ggplot(exampledata)</pre>
> plotobj <- plotobj + geom_point(aes(x=X, y=Y, shape=TRT, colour=ID))</pre>
> plotobj <- plotobj + geom_line(aes(x=X, y=Y, colour=ID))</pre>
```











Scales

The plot of x and y can be made using a different axis scales:

- scale_x_continuous
- scale_y_discrete
- scale_x_log10
- scale_x_reverse
- scale_x_datetime etc.

Within a scale there are arguments for:

- axis label (label)
- axis limits (lim)
- axis tick breaks (breaks)



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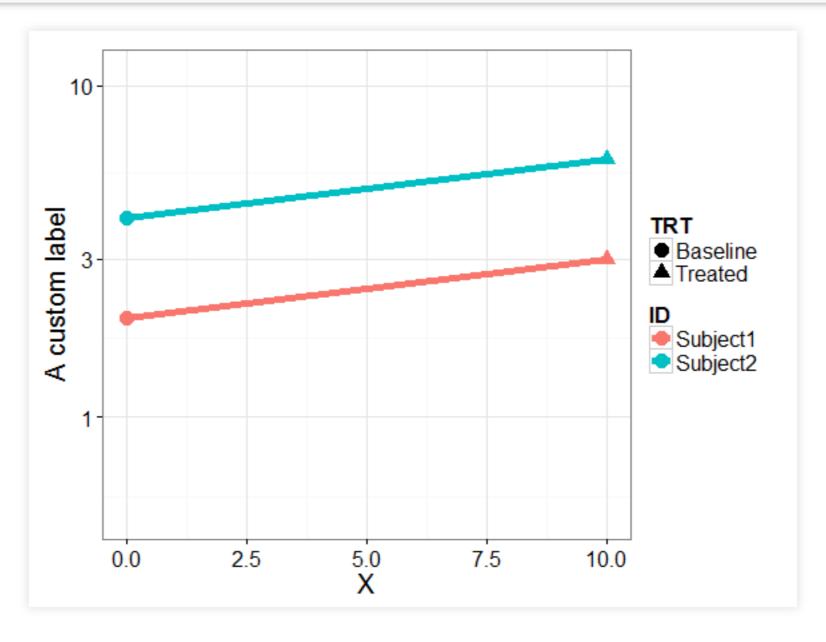








```
> plotobj <- ggplot(exampledata)</pre>
> plotobj <- plotobj + geom_point(aes(x=X, y=Y, shape=TRT, colour=ID))
> plotobj <- plotobj + geom_line(aes(x=X, y=Y, colour=ID))
> plotobj <- plotobj + scale_y_log10("A custom label")</pre>
```















Scales

Scales can also apply to aesthetics:

Example - a continuous colour scale of rainbow colors

Example - a discrete colour scale of red, blue & green

- scale_colour_continuous
- scale_colour_discrete
- scale_linetype etc.











Facetting

The plot of x and y can be facetted into sub-plots: This is powerful way to visualize subsets of the data

- facet_wrap
 - ~factor1+factor2
 - a linear series of plots displayed as ncol, rnow
- facet_grid
 - factor1 ~ factor2
 - a grid of plots displayed as factor vs factor









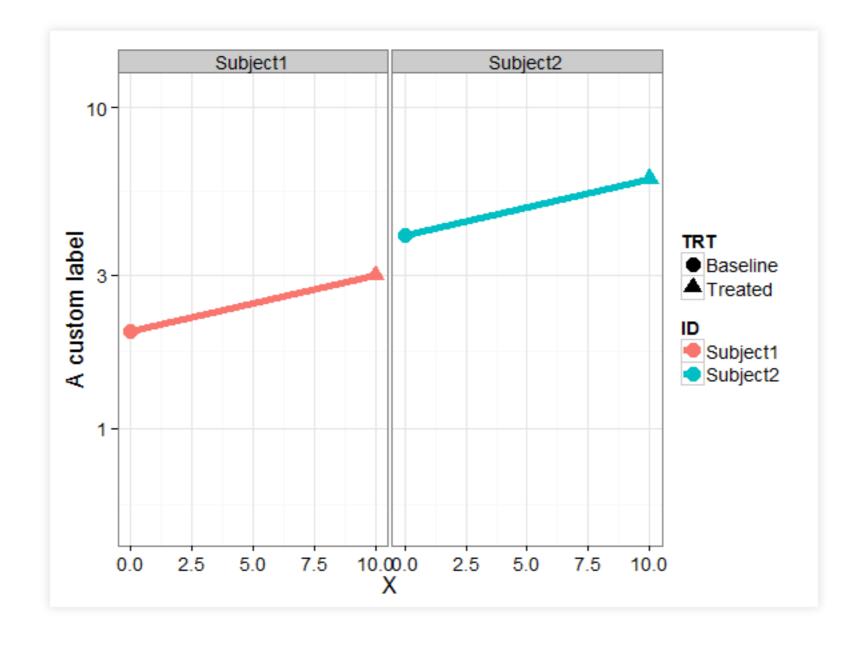


```
> plotobj <- ggplot(exampledata)</pre>
> plotobj <- plotobj + geom_point(aes(x=X, y=Y, shape=TRT, colour=ID))</pre>
> plotobj <- plotobj + geom_line(aes(x=X, y=Y, colour=ID))</pre>
> plotobj <- plotobj + scale_y_log10("A custom label")</pre>
> plotobj <- plotobj + facet_wrap(~ID)</pre>
```



6

Facetting















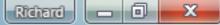
Statistical Summaries

Statistical summaries can be added to the plot of x and y:

Summary statistics are calculated inside the plot This is handy and time-efficient

- stat_summary
- stat density
- stat_qq
- stat_smooth (Loess, polynomial, linear)

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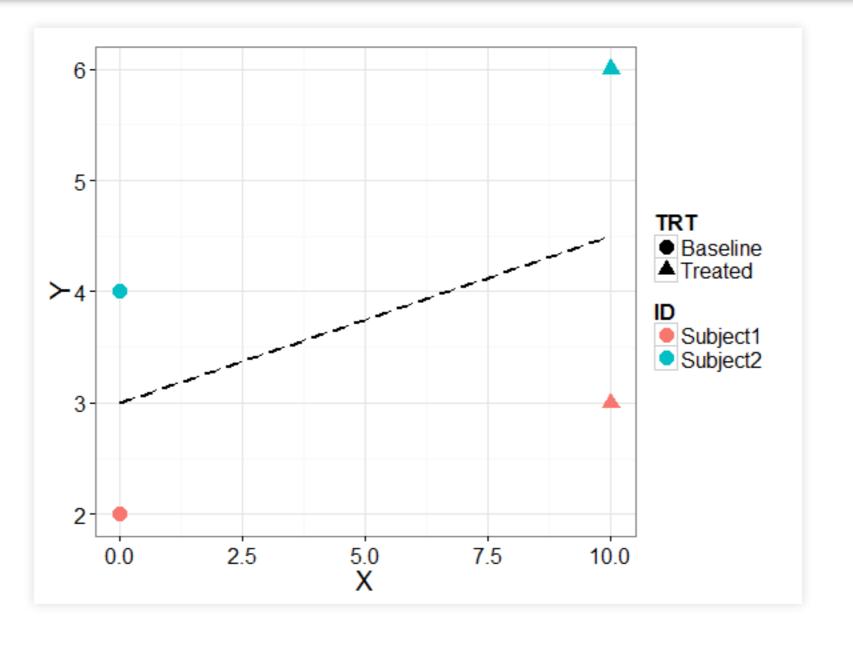








```
> plotobj <- ggplot(exampledata)</pre>
> plotobj <- plotobj + geom_point(aes(x=X, y=Y, shape=TRT, colour=ID))</pre>
> plotobj <- plotobj + stat_summary(aes(x=X, y=Y), fun.y=mean, geom="line",</pre>
linetype="dashed")
```

















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Themes control the overall look for a plot: The standard theme is influenced by Edward Tufte some prefer bolder colours also see the package "ggthemes" publication ready b+w is possible, but obscures information

Elements of a plot can be adjusted individually: fonts and font sizes legend positions plot margins





```
> plotobj <- ggplot(exampledata)
> plotobj <- plotobj + geom_point(aes(x=X, y=Y, shape=TRT, colour=ID))
> plotobj <- plotobj + scale_x_continuous("This is the X data")
> plotobj <- plotobj + ggtitle("My graph")
> plotobj <- plotobj + theme_gray()
> plotobj <- plotobj + theme(legend.position = "left")
> plotobj <- plotobj + theme(axis.title.x = element_text(colour = "red", face="italic"))</pre>
```







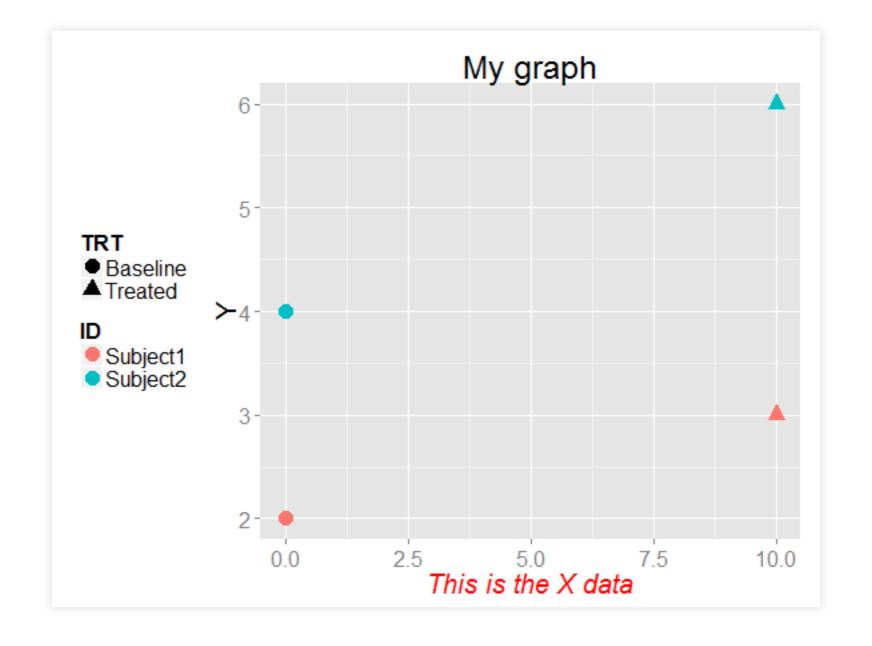




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Themes and Elements











Saving plots

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ggsave is a good place to start see also ?devices

ggsave writes the current plot object to a file file type is inferred from the extension set the height and width as appropriate

```
> plotobj <- ggplot(exampledata)</pre>
> plotobj <- plotobj + geom_point(aes(x=X, y=Y,</pre>
shape=TRT, colour=ID))
> ggsave("myplot.png", width=5,height=4)
```











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ggplot2 is evolving as a key method for data visualization

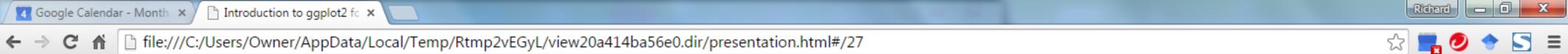
Pros:

It's reproducible Script based plotting makes complicated plots easy Quickly reshape plots to investigate relationships Code can be recycled

Cons:

It discourages "stupid" plots (pie charts, 2 y axes) Fine control over plots sometimes frustrating (subscripts, superscripts) There is a learning-curve folks































Installing packages

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This will be the first time we have used a package

Packages need to be *installed* once in your version of R Do this from the Packages\Install Package(s) menu Select a CRAN mirror Select the package by name They can also be installed from a downloaded zip file

Packages need to be *loaded* once in each session of R do this with the library function: library(ggplot2)

If a script can't find a function, it's package may not be loaded

You can review your installed packages with installed.packages()

