Introduction to ggplot2

R Pruim

July, 2014

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Goals

What I will try to do

- give a tour of ggplot2
- explain how to think about plots the ggplot2 way

► prepare/encourage you to learn more later

What I can't do in one session

- show every bell and whistle
- make you an expert at using ggplot2

The Births78 data set - revised edition

```
require(dplyr)
require(mosaic)
require(lubridate)
Births2 <- Births78 %>%
  mutate(
    date = mdy(date) - years(100), # y2k fix
    wd = wday(date), # as a number
    wday = wday(date, label=TRUE, abbr=TRUE) # as text (4)
    head(Births2, 2)
```

##		date	births	dayofyear	wd	wday
##	1	1978-01-01	7701	1	1	Sun
##	2	1978-01-02	7527	2	2	Mon

The grammar of graphics

geom: the geometric "shape" used to display data (glyph)

► bar, point, line, ribbon, text, etc.

aesthetic: an attribute controlling how geom is displayed

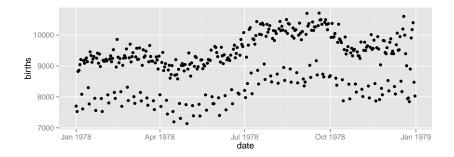
- ► x position, y position, color, fill, shape, size, etc.
- stat: a transformation applied to data before geom gets it
 - example: histograms work on binned data

scale: conversion of raw data to visual display

► particular assignment of colors, shapes, sizes, etc.

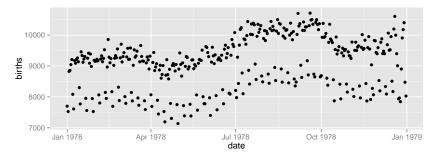
guide: helps user convert visual data back into raw data (legends, axes)

How do we make this plot?



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How do we make this plot?



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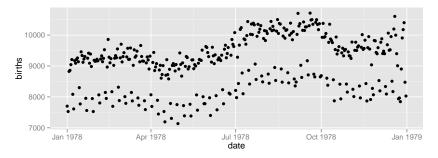
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What does R need to know?

How do we make this plot?



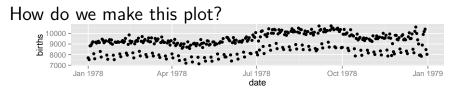
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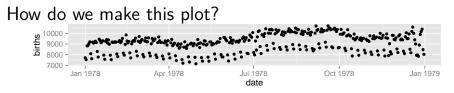
What does R need to know?

- data source
- ► aesthetics
- ▶ geom dots



What does R need to know?



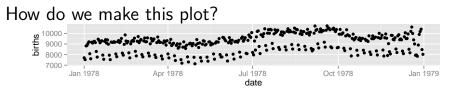


What does R need to know?

data frame containing the data: ggplot(data=)

```
ggplot(data=Births2)
```





What does R need to know?

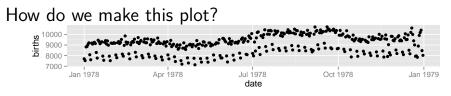
data frame containing the data: ggplot(data=)

```
ggplot(data=Births2)
```

* how we want to map our aesthetics: aes()

ggplot(data=Births2, aes(x=date, y=births))

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What does R need to know?

• data frame containing the data: ggplot(data=)

```
ggplot(data=Births2)
```

* how we want to map our aesthetics: aes()

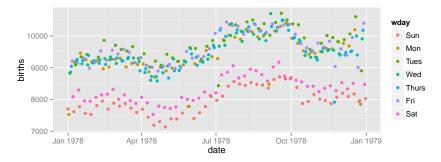
ggplot(data=Births2, aes(x=date, y=births))

what geom we want to use: + geom point()

ggplot(data=Births2, aes(x=date, y=births)) + geom point() ・ロト ・ 日 ト ・ 日 ト

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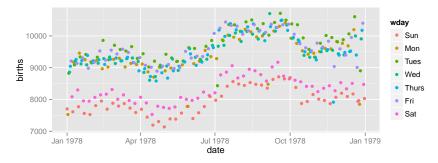


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What information has changed?

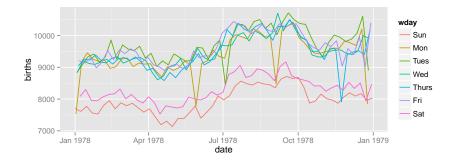


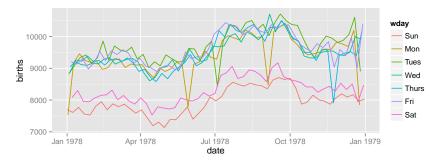
What information has changed?

new aesthetic: mapping color to day of week

ggplot(data=Births2, aes(x=date, y=births, color=wday)) +
geom_point()

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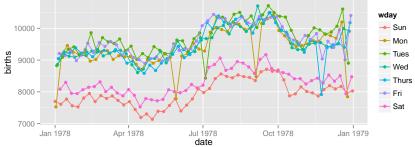
This time we use lines instead of dots

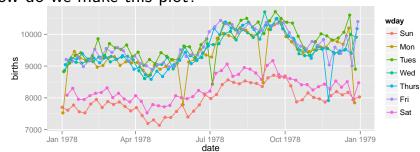
ggplot(data=Births2, aes(x=date, y=births, color=wday)) +
geom_line()

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How do we make this plot?

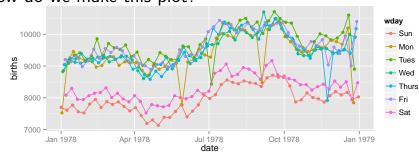




This time we have two layers, one with points and one with lines

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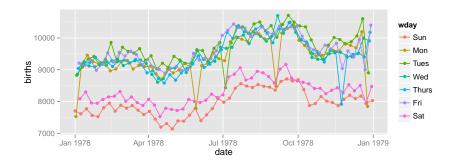


This time we have two layers, one with points and one with lines

► The layers are placed one on top of the other: the points are below and the lines are above. Sometimes the order of the layers can be important because of overplotting.

Alternative Syntax

```
Births2 %>%
ggplot(aes(x=date, y=births, color=wday)) +
geom_point() +
geom_line()
```

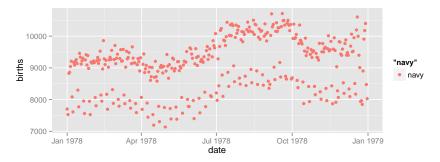


What does this do?

```
Births2 %>%
ggplot(aes(x=date, y=births, color="navy")) +
geom_point()
```

What does this do?

Births2 %>%
ggplot(aes(x=date, y=births, color="navy")) +
geom_point()

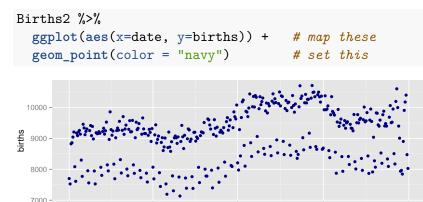


This is *mapping* the color aesthetic to a new variable with only one value ("navy"). So all the dots get set to the same color, but it's not navy.

Setting vs. Mapping

Jan 1978

If we want to *set* the color to be navy for all of the dots, we do it this way:



Apr 1978

► Note that color = "navy" is now outside of the aesthetics list. That's how ggplot2 distinguishes between mapping and setting.

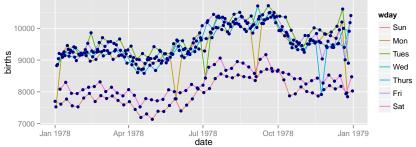
Jul 1978

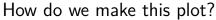
date

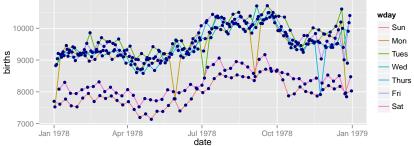
Oct 1978

lan 1979

How do we make this plot?







Births2 %>%
ggplot(aes(x=date, y=births)) +
geom_line(aes(color=wday)) + # map color here
geom_point(color="navy") # set color here

- ggplot() establishes the default data and aesthetics for the geoms, but each geom may change these defaults.
- ▶ good practice: put into ggplot() the things that affect all (or most) of the layers; rest in geom_blah(), above ab

Other geoms

apropos("^geom_")

- [1] "geom_abline"
- [4] "geom_bin2d"

[10] "geom_density2d"

[13] "geom_errorbarh"

[16] "geom_histogram"

[19] "geom_line"

[22] "geom_path"

[31] "geom_rug"

[34] "geom_step"

[25] "geom_polygon"

[28] "geom_raster"

- [7] "geom_contour"

[37] "geom_tufteboxplot" "geom_violin"

help pages will tell you their aesthetics and default stats

- "geom_area" "geom_blank"

 - "geom_crossbar"

 - "geom_dotplot"

 - "geom_freqpoly"

"geom_linerange"

"geom_quantile"

"geom_segment"

"geom_bar"

"geom_hex"

"geom_map"

"geom_boxplot

"geom_density

"geom_errorba

"geom_jitter"

"geom_pointra

"geom_rangefra

"geom_ribbon"

"geom_smooth"

"geom_tile"

"geom_vline"

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"geom_hline"

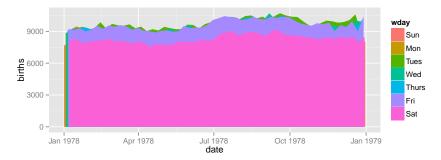
"geom_point"

"geom_rect"

"geom_text"

Let's try geom_area

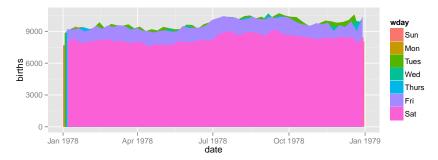
```
Births2 %>%
ggplot(aes(x=date, y=births, fill=wday)) +
geom_area()
```



This is not a good plot

Let's try geom_area

```
Births2 %>%
ggplot(aes(x=date, y=births, fill=wday)) +
geom_area()
```



This is not a good plot

- overplotting is hiding much of the data
- ► extending y-axis to 0 may or may not be desirable.

Side note: what makes a plot good?

Most (all?) graphics are intended to help us make comparisons

- How does something change over time?
- ► Do my treatments matter? How much?
- Do men and women respond the same way?

Key plot metric: Does my plot make the comparisions I am interested in

- easily, and
- ► accurately?

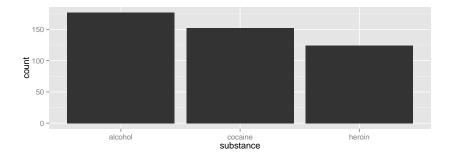
Time for some different data

HELPrct: Health Evaluation and Linkage to Primary care randomized clinical trial

?HELPrct

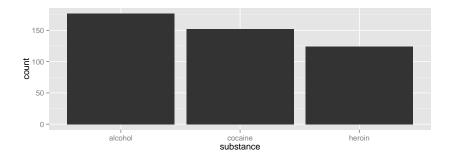
Why are these people in the study?

```
HELPrct %>%
ggplot(aes(x=substance)) +
geom_bar()
```



Why are these people in the study?

```
HELPrct %>%
ggplot(aes(x=substance)) +
geom_bar()
```

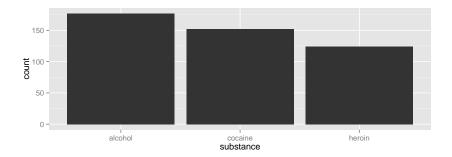


► Hmm. What's up with y?

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Why are these people in the study?

```
HELPrct %>%
ggplot(aes(x=substance)) +
geom_bar()
```



- ► Hmm. What's up with y?
 - stat_bin() is being applied to the data before the geom_bar() gets to do its thing. Binning creates the y values.

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Data Flow

org data $\xrightarrow{\text{stat}}$ statified $\xrightarrow{\text{aesthetics}}$ aesthetic data $\xrightarrow{\text{scales}}$ scaled data Simplifications:

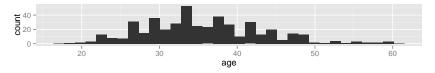
- aesthetics get computed twice, once before the stat and again after. Examples: bar charts, histograms
- ► item we need to look at the aesthetics to figure out which variable to bin
 - then the stat does the binning
 - bin counts become part of the aesthetics for geom: y=..count..
- ► This process happens in each layer
- stat_identity() is the "do nothing" stat.

How old are people in the HELP study?

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How old are people in the HELP study?

```
HELPrct %>%
ggplot(aes(x=age)) +
geom_histogram()
```

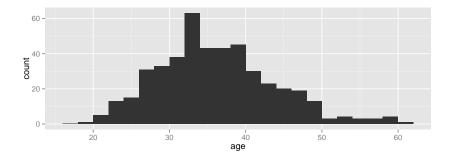


Notice the messages

- stat_bin: Histograms are not mapping the raw data but binned data.
 stat_bin() performs the data transformation.
- binwidth: a default binwidth has been selected, but we should really choose our own.

Setting the binwidth manually

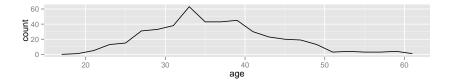
```
HELPrct %>%
ggplot(aes(x=age)) +
geom_histogram(binwidth=2)
```



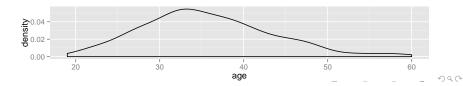
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How old are people in the HELP study? - Other geoms

```
HELPrct %>%
ggplot(aes(x=age)) +
geom_freqpoly(binwidth=2)
```



HELPrct %>%
ggplot(aes(x=age)) +
geom_density()

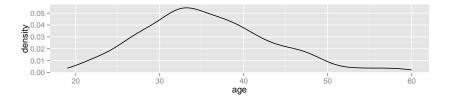


Selecting stat and geom manually

Every geom comes with a default stat

- for simple cases, the stat is stat_identity() which does
 nothing
- ▶ we can mix and match geoms and stats however we like

```
HELPrct %>%
ggplot(aes(x=age)) +
geom_line(stat="density")
```

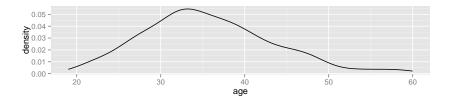


Selecting stat and geom manually

Every stat comes with a default geom

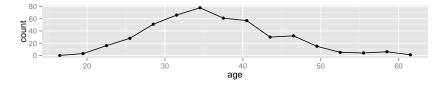
- ▶ we can specify stats instead of geom, if we prefer
- we can mix and match geoms and stats however we like

```
HELPrct %>%
ggplot(aes(x=age)) +
stat_density( geom="line")
```

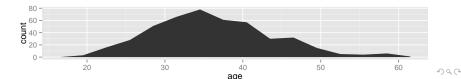


More combinations

```
HELPrct %>%
ggplot(aes(x=age)) +
geom_point(stat="bin", binwidth=3) +
geom_line(stat="bin", binwidth=3)
```



HELPrct %>%
ggplot(aes(x=age)) +
geom_area(stat="bin", binwidth=3)

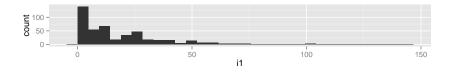


Your turn: How much do they drink? (i1)

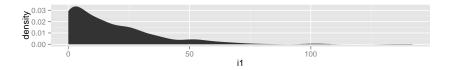
Create a plot that shows the distribution of the average daily alcohol consumption in the past 30 days (i2).

How much do they drink? (i1)

HELPrct %>%
ggplot(aes(x=i1)) + geom_histogram()



HELPrct %>%
ggplot(aes(x=i1)) + geom_area(stat="density")



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Covariates: Adding in more variables

Q. How does alcohol consumption (or age, your choice) differ by sex and substance (alcohol, cocaine, heroin)?

Decisions:

- How will we display the variables: i1 (or age), sex, substance
- What comparisons are we most interested in?

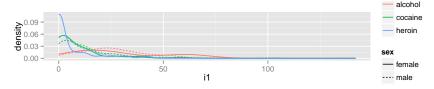
Give it a try.

 Note: I'm cheating a bit. You may want to do some things I haven't shown you yet. (Feel free to ask.)

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Covariates: Adding in more variables Using color and linetype:

```
HELPrct %>%
ggplot(aes(x=i1, color=substance, linetype=sex)) +
geom_line(stat="density")
```



Using color and facets

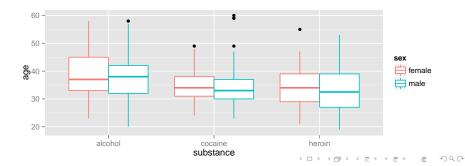
```
HELPrct %>%
ggplot(aes(x=i1, color=substance)) +
geom_line(stat="density") + facet_grid( . ~ sex )
```



Boxplots

Boxplots use stat_quantile() which computes a five-number summary (roughly the five quartiles of the data) and uses them to define a "box" and "whiskers". The quantitative variable must be y, and there must be an additional x variable.

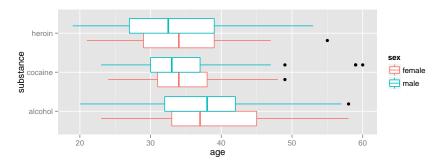
```
HELPrct %>%
ggplot(aes(x=substance, y=age, color=sex)) +
geom_boxplot()
```



Horizontal boxplots

Horizontal boxplots are obtained by flipping the coordinate system:

```
HELPrct %>%
ggplot(aes(x=substance, y=age, color=sex)) +
geom_boxplot() +
coord_flip()
```

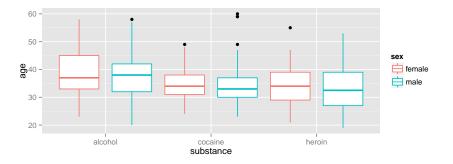


► coord_flip() may be used with other plots as well to reverse the roles of x and y on the plot.

Give me some space

We've triggered a new feature: dodge (for dodging things left/right). We can control how much if we set the dodge manually.

```
HELPrct %>%
ggplot(aes(x=substance, y=age, color=sex)) +
geom_boxplot(position=position_dodge(width=1))
```

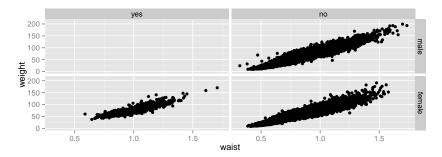


Issues with bigger data

dim(NHANES)

[1] 31126 53

NHANES %>% ggplot(aes(x=waist, y=weight)) +
geom_point() + facet_grid(sex ~ pregnant)



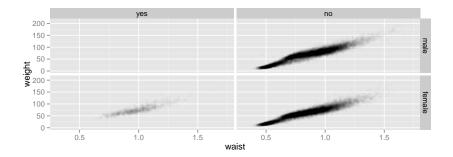
 Although we can see a generally positive association (as we would expect), the overplotting may be hiding information.

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Using alpha (opacity)

One way to deal with overplotting is to set the opacity low.

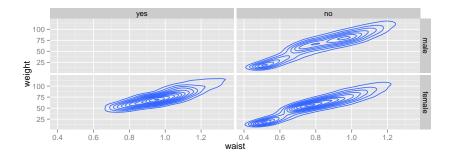
```
NHANES %>%
ggplot(aes(x=waist, y=weight)) +
geom_point(alpha=0.01) + facet_grid( sex ~ pregnant )
```



geom_density2d

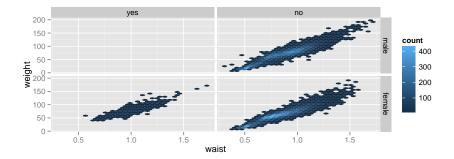
Alternatively (or simultaneously) we might prefere a different geom altogether.

```
NHANES %>%
ggplot(aes(x=waist, y=weight)) +
geom_density2d() + facet_grid( sex ~ pregnant )
```



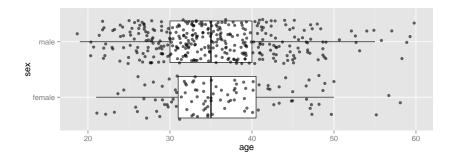
geom_hex

```
NHANES %>%
ggplot(aes(x=waist, y=weight)) +
geom_hex() + facet_grid( sex ~ pregnant )
```



Multiple layers

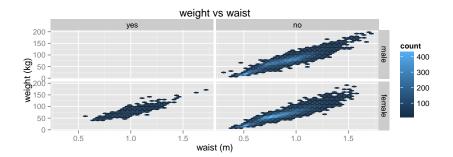
```
ggplot( data=HELPrct, aes(x=sex, y=age)) +
geom_boxplot(outlier.size=0) +
geom_jitter(alpha=.6) +
coord_flip()
```



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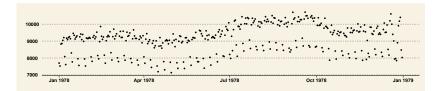
Labeling

```
NHANES %>%
ggplot(aes(x=waist, y=weight)) +
geom_hex() + facet_grid( sex ~ pregnant ) +
labs(x="waist (m)", y="weight (kg)", title="weight vs was
```



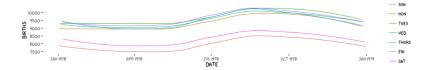
- scales (fine tuning mapping from data to plot)
- guides (so reader can map from plot to data)
- coords (coord_flip() is good to know about)
- themes (for customizing appearance)

require(ggthemes)
qplot(x=date, y=births, data=Births2) + theme_wsj()



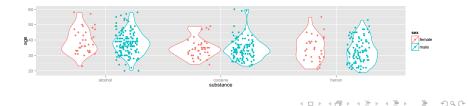
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- scales (fine tuning mapping from data to plot)
- guides (so reader can map from plot to data)
- coords (coord_flip() is good to know about)
- themes (for customizing appearance)



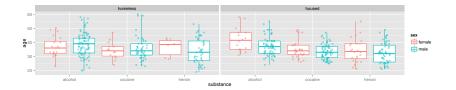
- scales (fine tuning mapping from data to plot)
- ▶ guides (so reader can map from plot to data)
- coords (coord_flip() is good to know about)
- themes (for customizing appearance)
- ▶ position (position_dodge() can be used for side by side bars)

ggplot(data=HELPrct, aes(x=substance, y=age, color=sex)) geom_violin(coef = 10, position=position_dodge()) +
geom_point(aes(color=sex, fill=sex), position=position_j;



- scales (fine tuning mapping from data to plot)
- guides (so reader can map from plot to data)
- ▶ themes (for customizing appearance)
- > position (position_dodge(), position_jitterdodge(), position_stack(), etc.)

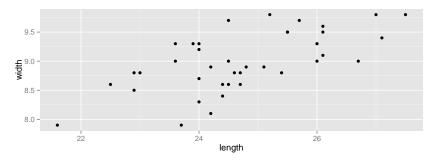
A little bit of everything



Some short cuts

1. qplot() provides "quick plots" for ggplot2

qplot(length, width, data=KidsFeet)



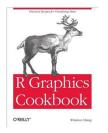
mplot(dataframe) provides an interactive plotting tool for both ggplot2 and lattice.

mplot(HELPrct)

► quickly make several plots from a data frame (D > (≥) (≥) (≥) (≥)

Want to learn more?

- ► docs.ggplot2.org/
- ► Winston Chang's: *R Graphics Cookbook*



What's around the corner?

ggvis

- ► dynamic graphics (brushing, sliders, tooltips, etc.)
- ▶ uses Vega (D3) to animate plots in a browser
- similar structure to ggplot2 but different syntax and names

version 0.3 just released to github

Dynamic documents

- ► combination of RMarkdown, ggvis, and shiny
- beta testing now