

# Data Visualization in R

## 4. ggplot2

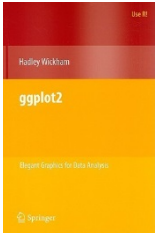


Michael Friendly  
SCS Short Course  
Sep/Oct, 2018



<http://www.datavis.ca/courses/RGraphics/>

# Resources: Books

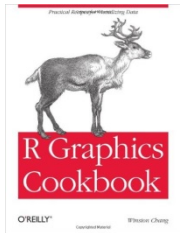


Hadley Wickham, *ggplot2: Elegant graphics for data analysis*, 2nd Ed.

1st Ed: Online, <http://ggplot2.org/book/>

ggplot2 Quick Reference: <http://sape.inf.usi.ch/quick-reference/ggplot2/>

Complete ggplot2 documentation: <http://docs.ggplot2.org/current/>

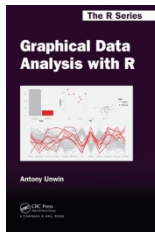


Winston Chang, *R Graphics Cookbook: Practical Recipes for Visualizing Data*

Cookbook format, covering common graphing tasks; the main focus is on ggplot2

R code from book: <http://www.cookbook-r.com/Graphs/>

Download from: <http://ase.tufts.edu/bugs/guide/assets/R%20Graphics%20Cookbook.pdf>

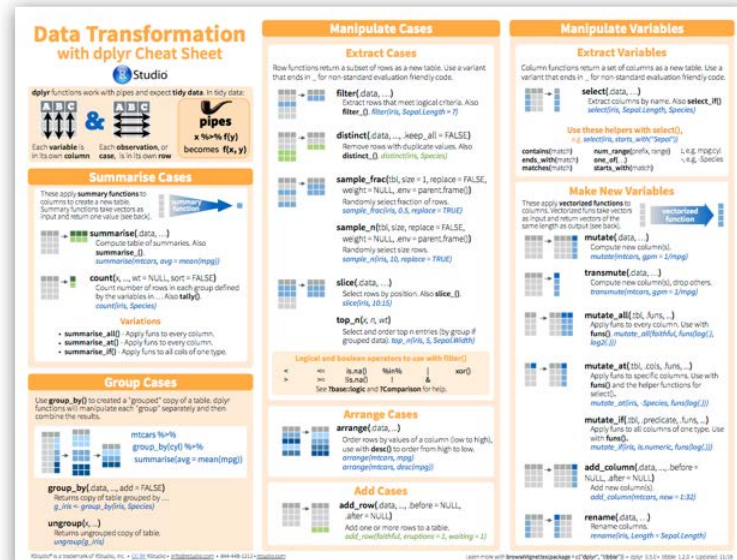
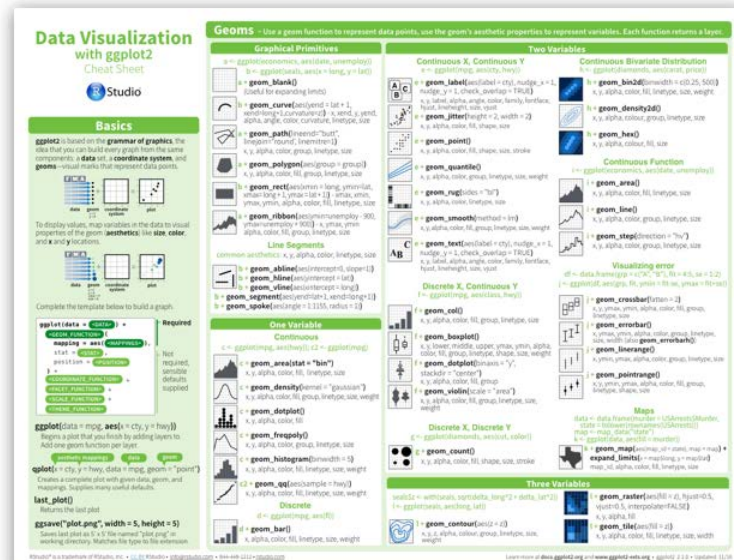


Antony Unwin, *Graphical Data Analysis with R*

R code: <http://www.gradaanwr.net/>

# Resources: Cheat sheets

- Data visualization with ggplot2:  
<https://www.rstudio.com/wp-content/uploads/2016/11/ggplot2-cheatsheet-2.1.pdf>
- Data transformation with dplyr:  
<https://github.com/rstudio/cheatsheets/raw/master/source/pdfs/data-transformation-cheatsheet.pdf>



# What is ggplot2?

- ggplot2 is Hadley Wickham's R package for producing “elegant graphics for data analysis”
  - It is an implementation of many of the ideas for graphics introduced in Lee Wilkinson's *Grammar of Graphics*
  - These ideas and the syntax of ggplot2 help to think of graphs in a new and more general way
  - Produces pleasing plots, taking care of many of the fiddly details (legends, axes, colors, ...)
  - It is built upon the “grid” graphics system
  - It is open software, with a large number of gg\_ extensions. See: <http://www.ggplot2-exts.org/gallery/>

# Follow along

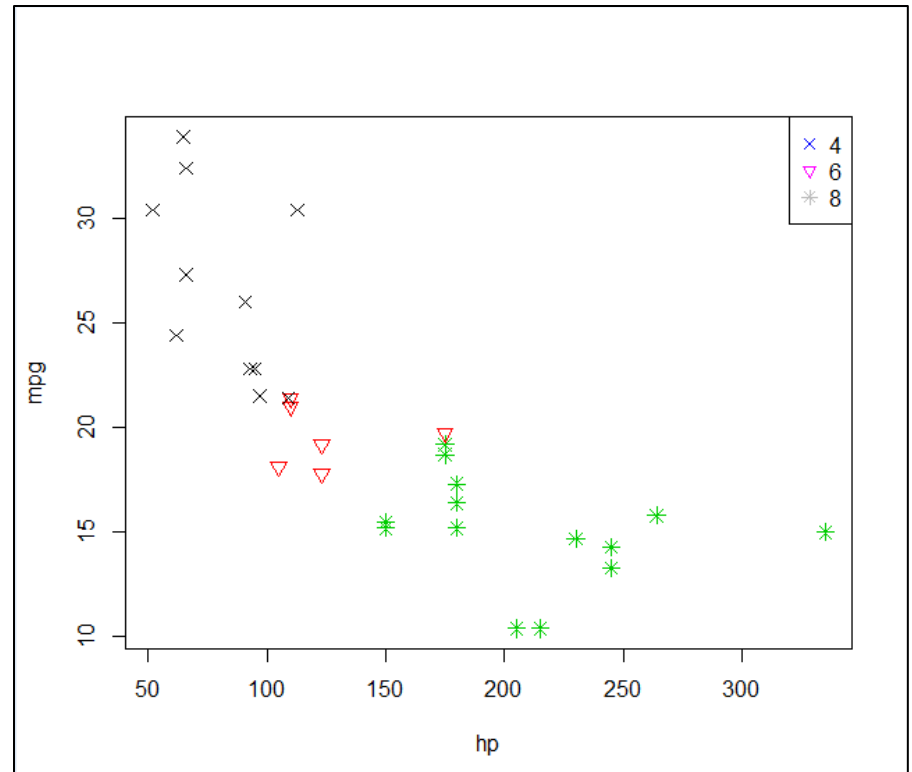
- From the course web page, click on the script gg-cars.R, <http://www.datavis.ca/courses/RGraphics/R/gg-cars.R>
- Select all (ctrl+A) and copy (ctrl+C) to the clipboard
- In R Studio, open a new R script file (ctrl+shift+N)
- Paste the contents (ctrl+V)
- Run the lines (ctrl+Enter) to along with me

# ggplot2 vs base graphics

Some things that should be simple  
are harder than you'd like in base  
graphics

Here, I'm plotting gas mileage (mpg) vs. horsepower and want to use color and shape for different # of cylinders.

But I don't quite get it right!



```
mtcars$cyl <- as.factor(mtcars$cyl)
plot(mpg ~ hp, data=mtcars,
      col=cyl, pch=c(4,6,8)[mtcars$cyl], cex=1.2)
legend("topright", legend=levels(mtcars$cyl),
      pch = c(4,6,8),
      col=levels(mtcars$cyl))
```

colors and point symbols work differently in plot() and legend()

# ggplot2 vs base graphics

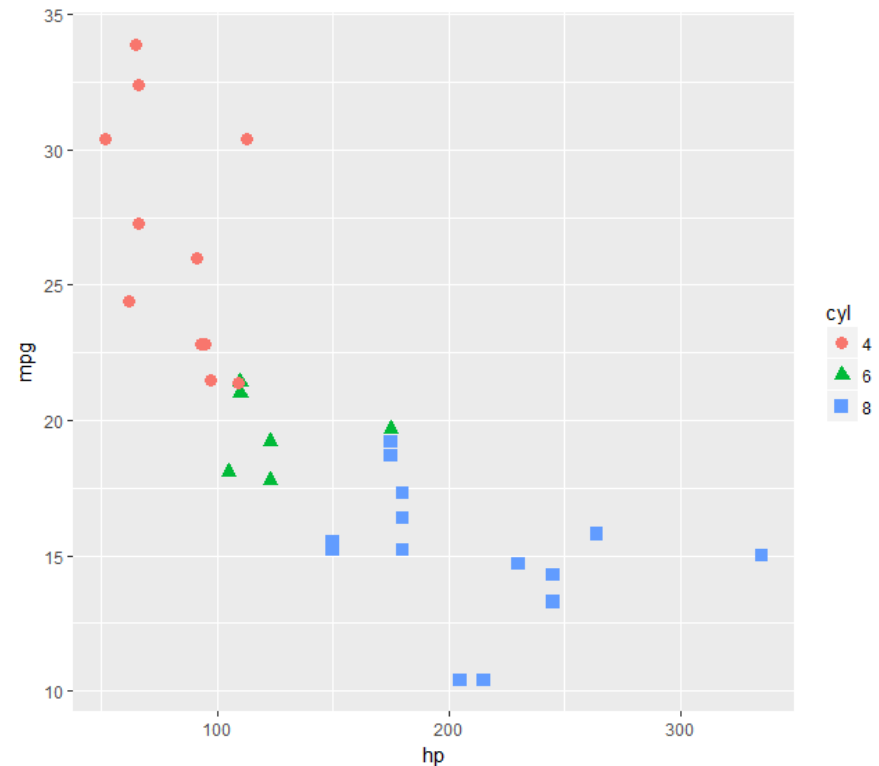
In ggplot2, just map the data variables to aesthetic attributes

`aes(x, y, shape, color, size, ...)`

`ggplot()` takes care of the rest

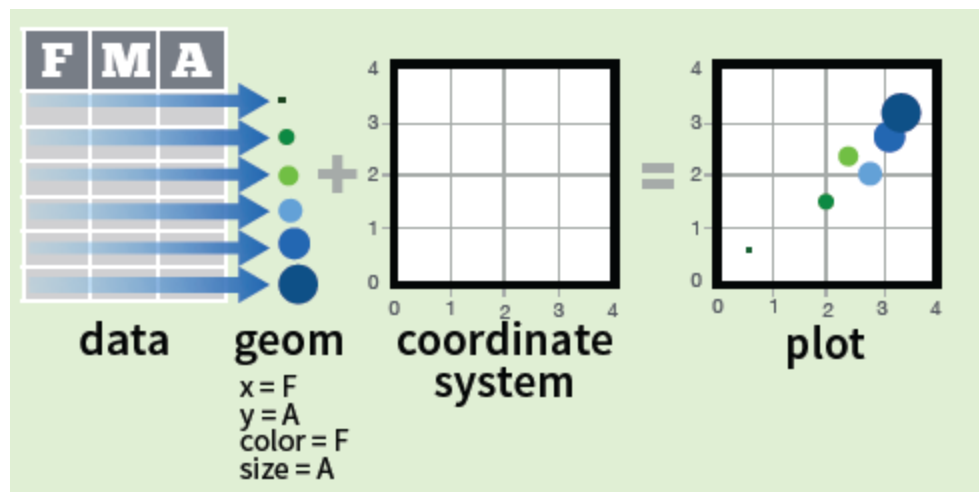
`aes()` mappings set in the call to `ggplot()` are passed to `geom_point()` here

```
library(ggplot2)
ggplot(mtcars, aes(x=hp, y=mpg, color=cyl, shape=cyl)) +
  geom_point(size=3)
```



# Grammar of Graphics

- Every graph can be described as a combination of independent building blocks:
  - **data**: a data frame: quantitative, categorical; local or data base query
  - **aes**thetic mapping of variables into visual properties: size, color, x, y
  - **geom**etric objects (“geom”): points, lines, areas, arrows, ...
  - **coord**inate system (“coord”): Cartesian, log, polar, map,





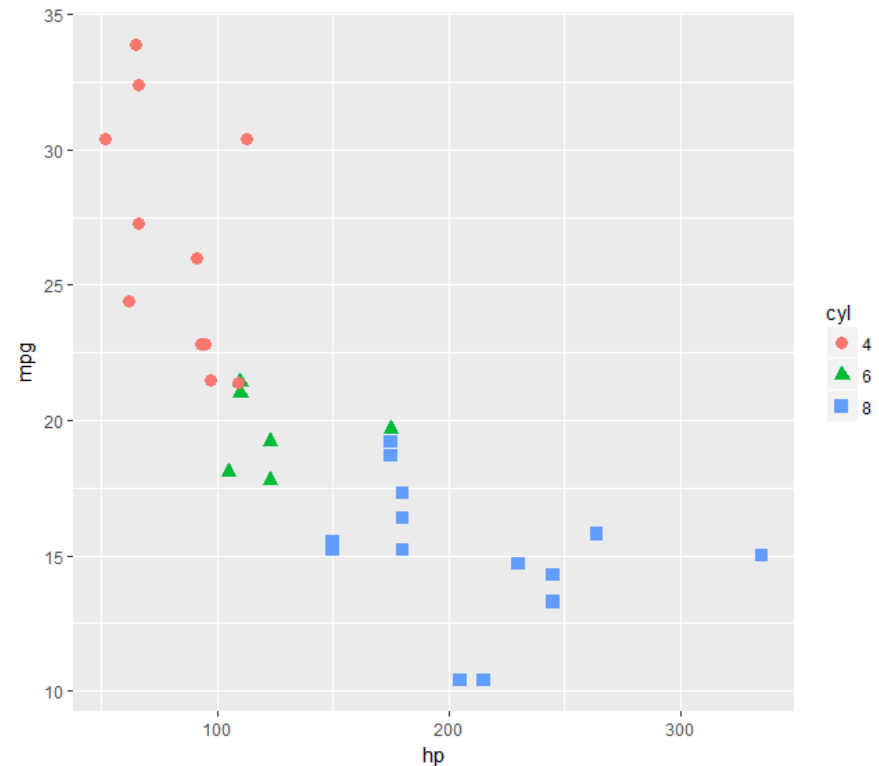
# ggplot2: data + geom -> graph

```
ggplot(data=mtcars,  
       aes(x=hp, y=mpg,  
           color=cyl, shape=cyl)) +  
  geom_point(size=3)
```

1  
2  
3  
4

In this call,

1. data=mtcars: data frame
2. aes(x=hp, y=mpg): plot variables
3. aes(color, shape): attributes
4. geom\_point(): what to plot
  - the coordinate system is taken to be the standard Cartesian (x,y)



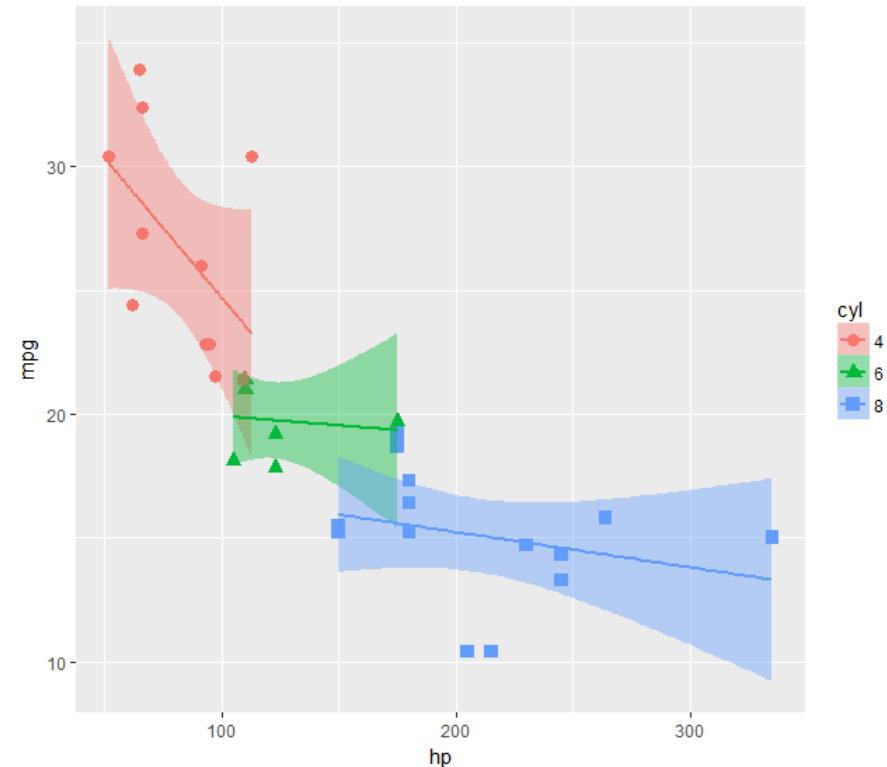
# ggplot2: geoms

Wow! I can really see something there.

How can I enhance this visualization?

Easy: add a `geom_smooth()` to fit linear regressions for each level of `cyl`

More generally: think of adding new layers to make a plot more useful.

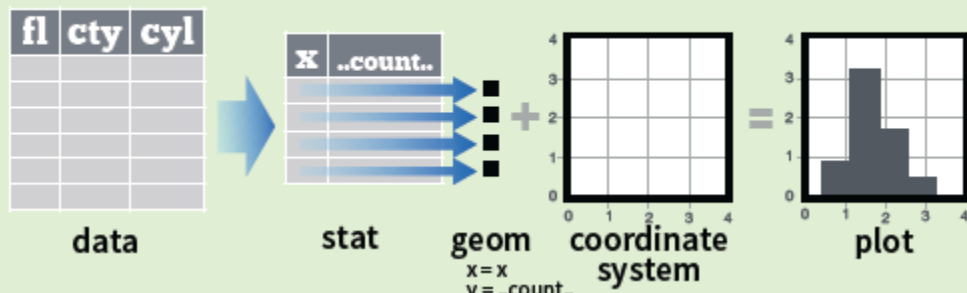


```
ggplot(mtcars, aes(x=hp, y=mpg, color=cyl, shape=cyl)) +  
  geom_point(size=3) +  
  geom_smooth(method="lm", aes(fill=cyl))
```

# Grammar of Graphics

- Other GoG building blocks:
  - **stat**istical transformations (“stat”) -- data summaries: mean, sd, binning & counting, ...
  - **scale**: legends, axes to allow reading data from a plot

A stat builds new variables to plot (e.g., count, prop).



# Grammar of Graphics

- Other GoG building blocks:
  - **position** adjustments: jitter, dodge, stack, ...
  - **faceting**: small multiples or conditioning to break a plot into subsets.

## Position Adjustments

Position adjustments determine how to arrange geoms that would otherwise occupy the same space.

```
s <- ggplot(mpg, aes(fl, fill = drv))
```



```
s + geom_bar(position = "dodge")
```

Arrange elements side by side



```
s + geom_bar(position = "fill")
```

Stack elements on top of one another, normalize height



```
e + geom_point(position = "jitter")
```

Add random noise to X and Y position of each element to avoid overplotting



```
e + geom_label(position = "nudge")
```

Nudge labels away from points



```
s + geom_bar(position = "stack")
```

Stack elements on top of one another

Each position adjustment can be recast as a function with manual **width** and **height** arguments

```
s + geom_bar(position = position_dodge(width = 1))
```

## Faceting

Facets divide a plot into subplots based on the values of one or more discrete variables.

```
t <- ggplot(mpg, aes(cty, hwy)) + geom_point()
```



```
t + facet_grid(. ~ fl)
```

facet into columns based on fl



```
t + facet_grid(year ~ .)
```

facet into rows based on year



```
t + facet_grid(year ~ fl)
```

facet into both rows and columns



```
t + facet_wrap(~ fl)
```

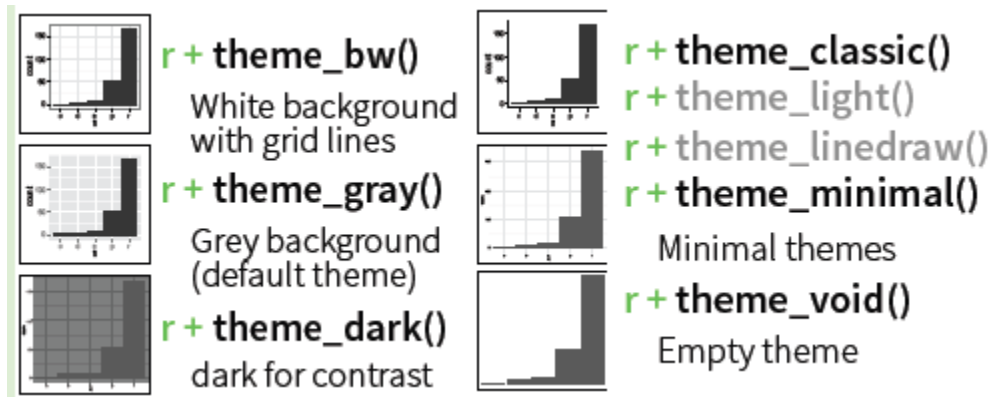
wrap facets into a rectangular layout

# ggplot2: GoG -> graphic language

- The implementation of GoG ideas in ggplot2 for R created a more expressive language for data graphs
  - **layers**: graph elements combined with “+” (read: “and”)

```
ggplot(mtcars, aes(x=hp, y=mpg)) +  
  geom_point(aes(color = cyl)) +  
  geom_smooth(method = "lm") +
```

- **theme**: change graphic elements consistently



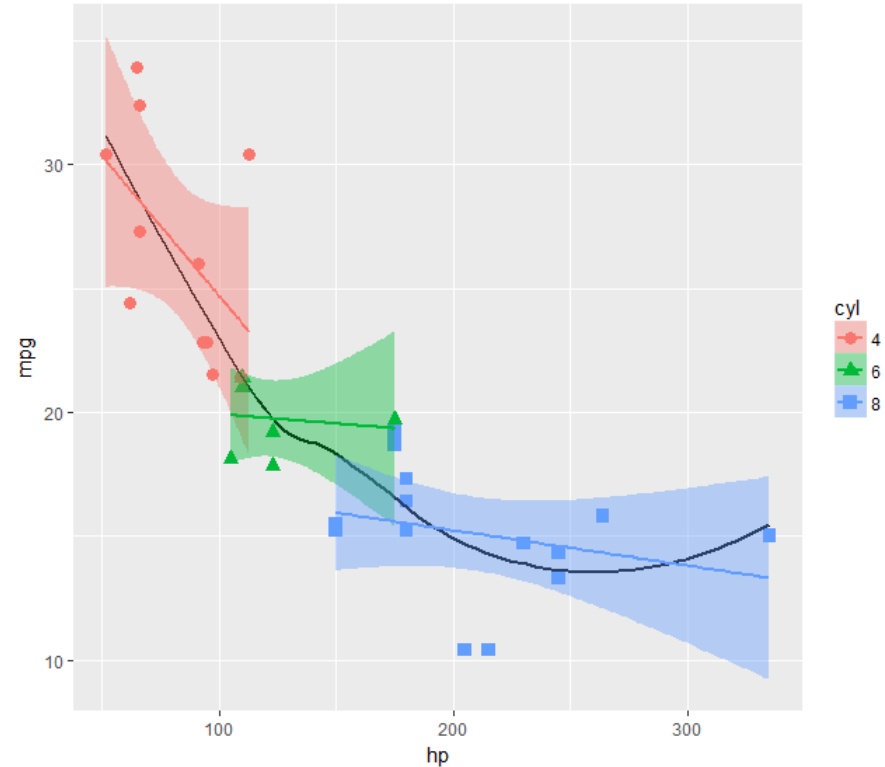
# ggplot2: layers & aes()

Aesthetic attributes in the `ggplot()` call are passed to `geom_()` layers

Other attributes can be passed as constants (`size=3, color="black"`) or with `aes(color=, ...)` in different layers

This plot adds an overall loess smooth to the previous plot

Specifying `color=` overrides other layers



```
ggplot(mtcars, aes(x=hp, y=mpg)) +  
  geom_point(size=3, aes(color=cyl, shape=cyl)) +  
  geom_smooth(method="lm", aes(color=cyl, fill=cyl)) +  
  geom_smooth(method="loess", color="black", se=FALSE)
```

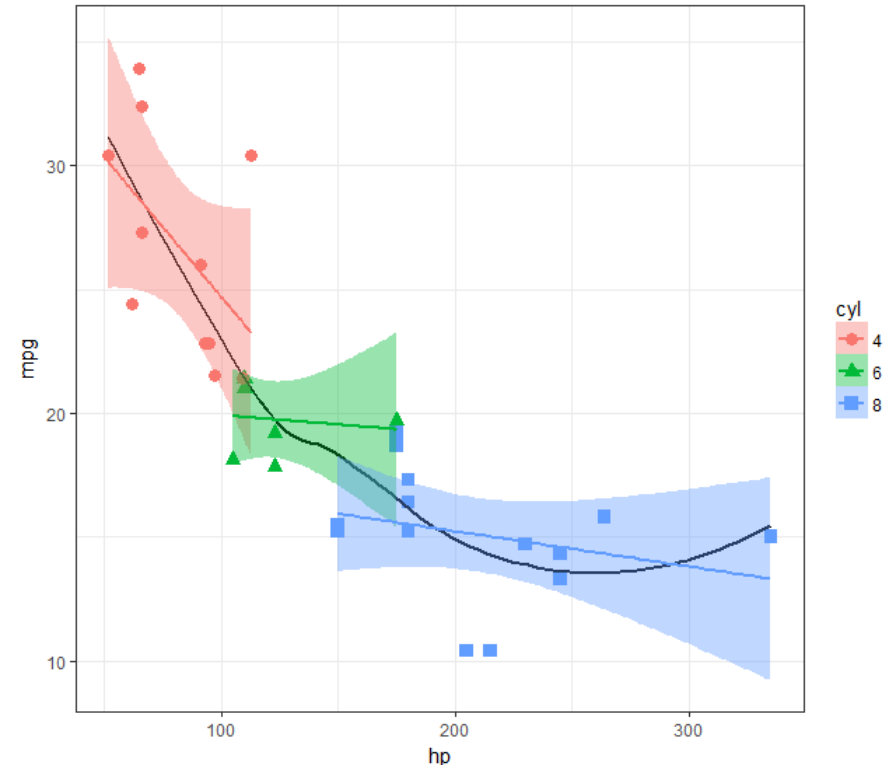
# ggplot2: themes

All the graphical attributes of ggplot2 are governed by themes – settings for all aspects of a plot

A given plot can be rendered quite differently just by changing the theme

If you haven't saved the ggplot object, `last_plot()` gives you something to work with further

```
last_plot() + theme_bw()
```

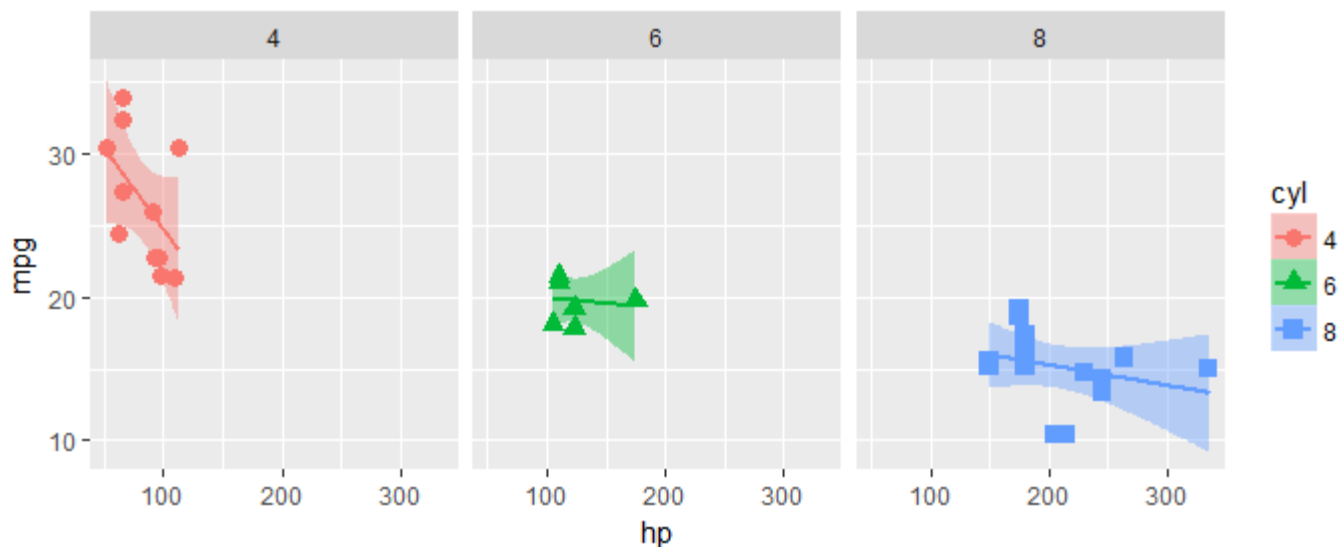


# ggplot2: facets

Facets divide a plot into separate subplots based on one or more discrete variables

```
plt <-  
ggplot(mtcars, aes(x=hp, y=mpg, color=cyl, shape=cyl)) +  
  geom_point(size=3) +  
  geom_smooth(method="lm", aes(fill=cyl))  
  
plt + facet_wrap(~cyl)
```

Faceting is most useful with other variables, not used in the main plot





# labeling points: geom\_text()

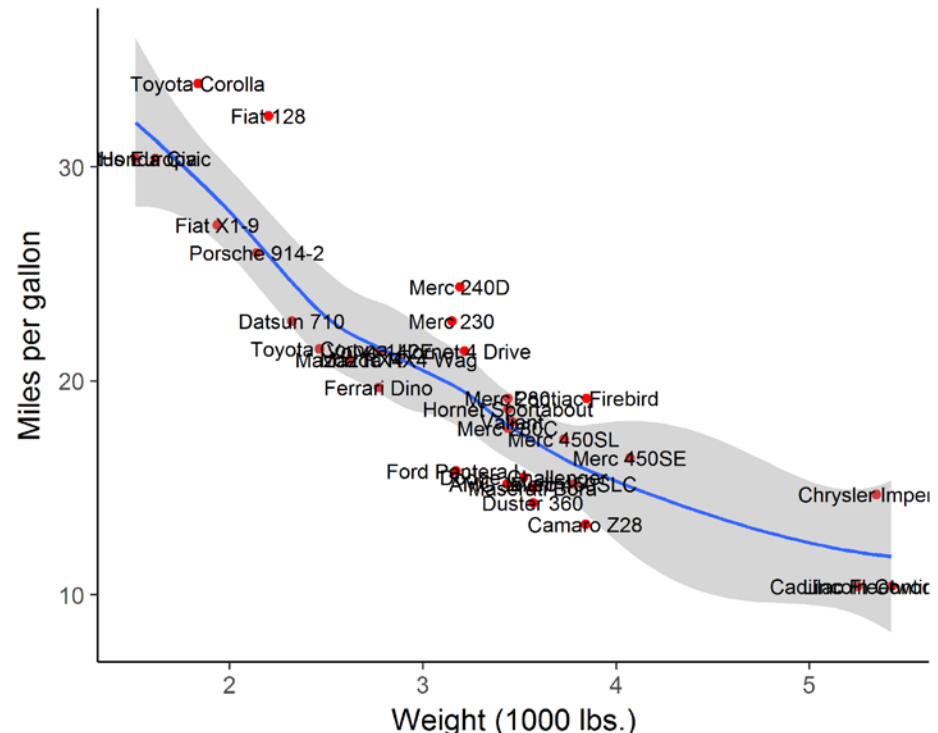
```
plt2 <- ggplot(mtcars, aes(x=wt, y=mpg)) +  
  geom_point(color = 'red', size=2) +  
  geom_smooth(method="loess") +  
  labs(y="Miles per gallon", x="Weight (1000 lbs.)") +  
  theme_classic(base_size = 16)
```

```
plt2 + geom_text(aes(label = rownames(mtcars)))
```

Note the use of `theme_classic()`  
and better axis labels

Sometimes it is useful to label  
points to show their identities.

`geom_text()` usually gives messy,  
overlapping text

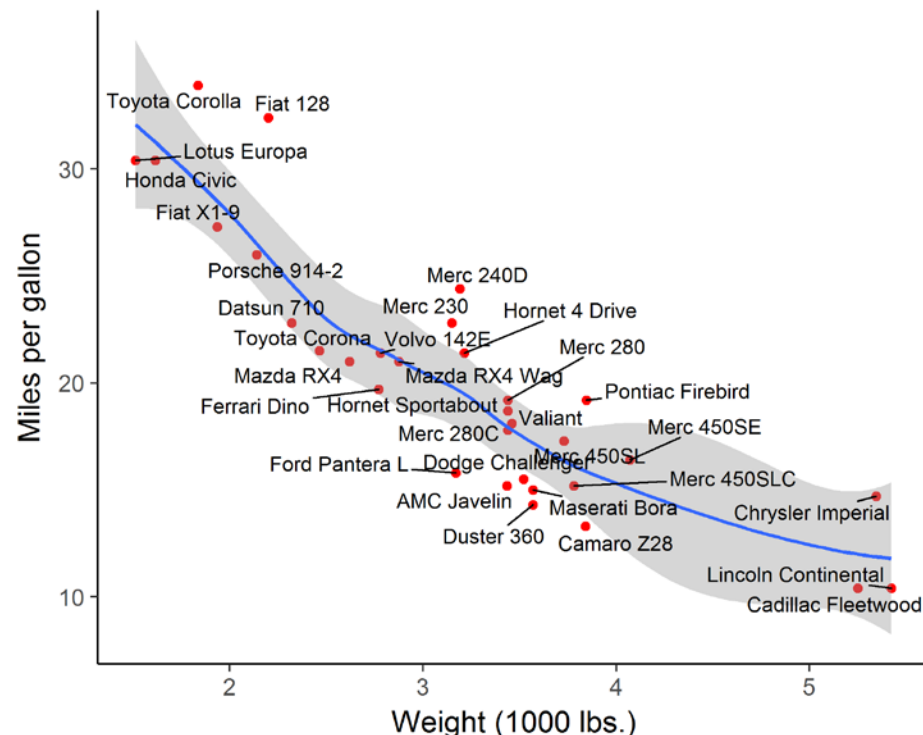


# labeling points: geom\_text\_repel()

```
install.packages("ggrepel")  
library(ggrepel)  
plt2 +  
  geom_text_repel(aes(label = rownames(mtcars)))
```

`geom_text_repel()` in the `ggrepel` package assigns repulsive forces among points and labels to assure no overlap

Some lines are drawn to make the assignment clearer



# labeling points: selection

It is easy to label points selectively, using some criterion to assign labels to points

```
mod <- loess( mpg ~ wt, data=mtcars)
resids <- residuals(mod)
mtcars$label <- ifelse(abs(resids) > 2.5,
                      rownames(mtcars), "")
```

```
plt2 + geom_text_repel(aes(label = mtcars$label))
```

1

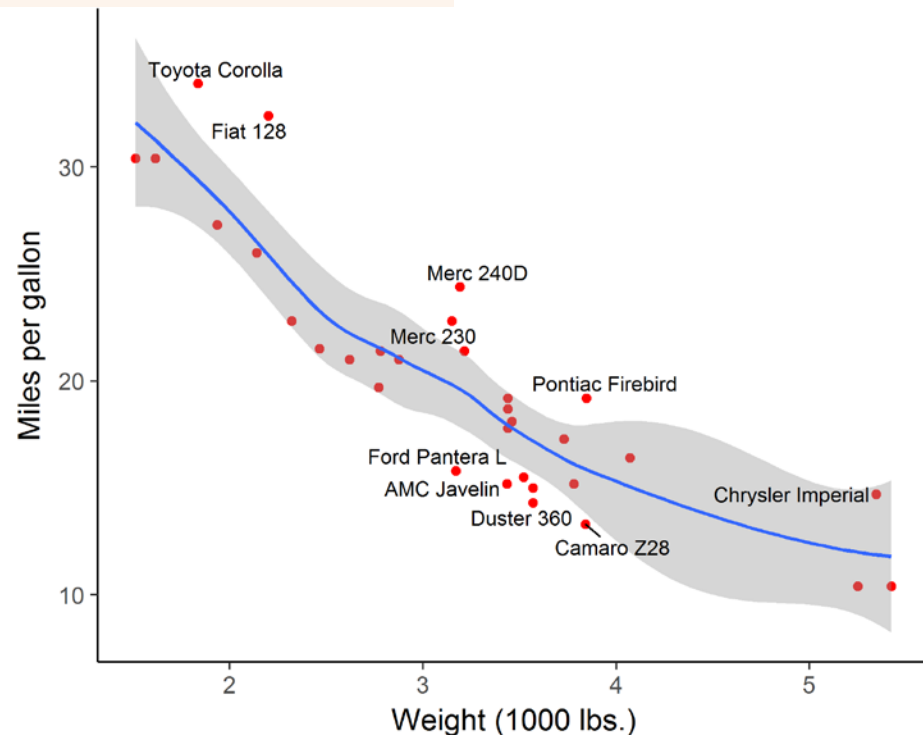
2

3

4

Here, I:

1. fit the smoothed loess curve,
2. extract residuals,  $r_i$
3. assign labels where  $|r_i| > 2.5$
4. add the text layer

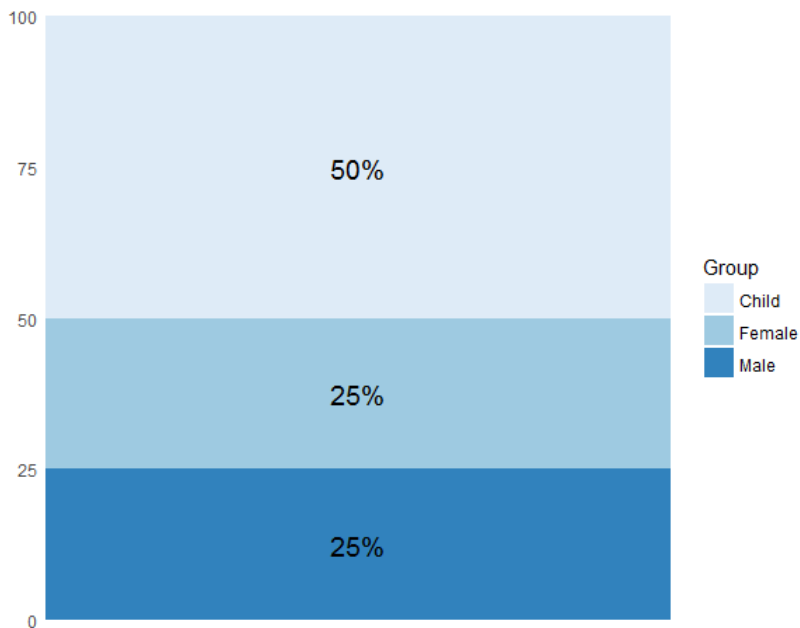


# ggplot2: coords

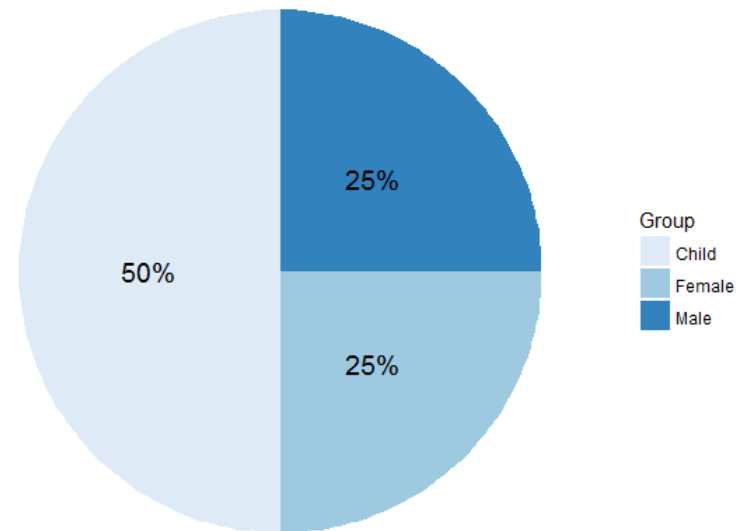
Coordinate systems, `coord_*()` functions, handle conversion from geometric objects to what you see on a 2D plot.

A pie chart is just a bar chart in polar coordinates!

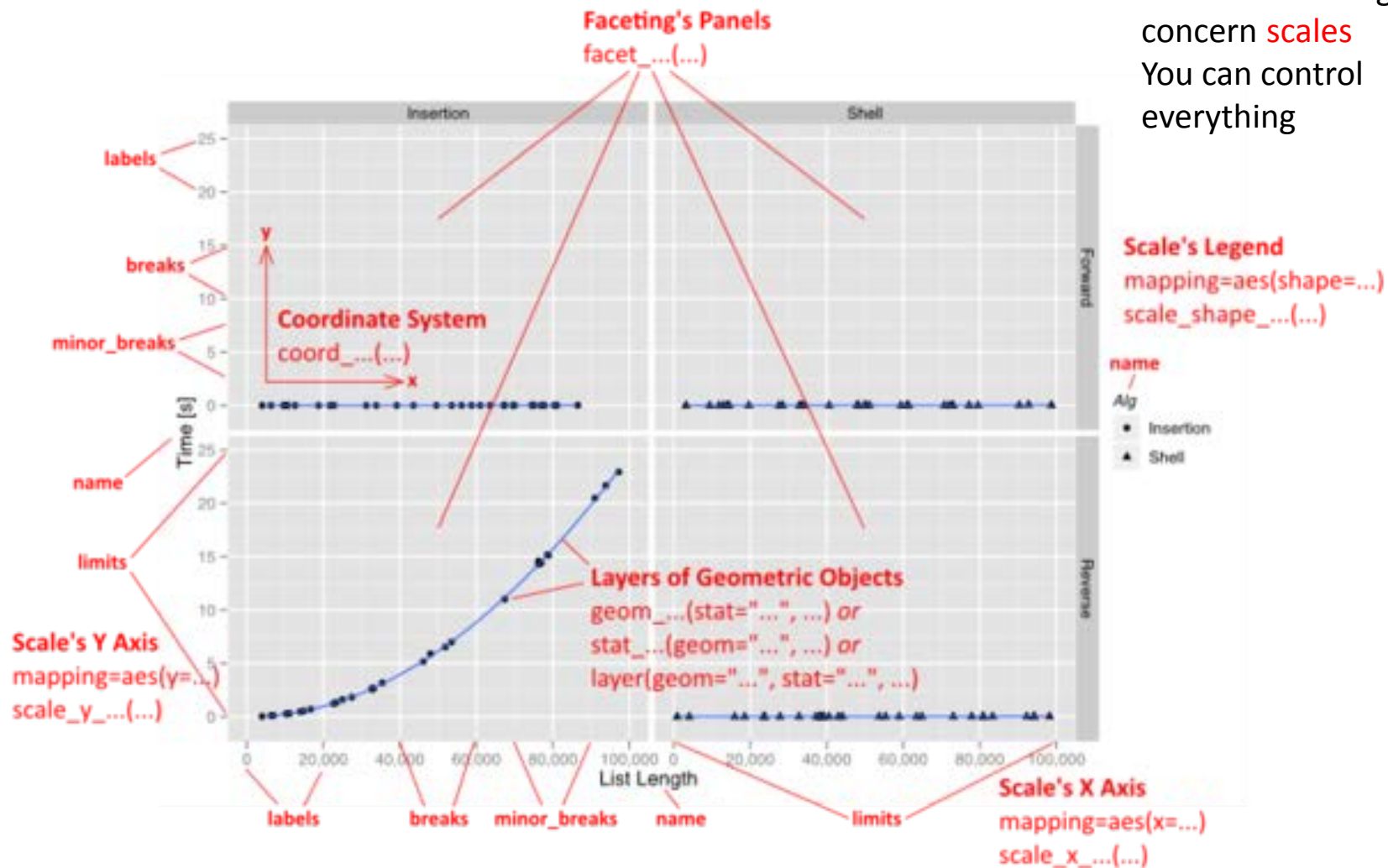
```
p <- ggplot(df, aes(x = "", y = value, fill = group)) +  
  geom_bar(stat = "identity")
```



```
p + coord_polar("y", start = 0)
```



# Anatomy of a ggplot



# ggplot objects

Traditional R graphics just produce graphical output on a device

However, `ggplot()` produces a “ggplot” object, a list of elements

```
> names(plt)
[1] "data"      "layers"    "scales"    "mapping"   "theme"     "coordinates"
[7] "facet"     "plot_env"  "labels"
> class(plt)
[1] "gg"  "ggplot"
```

What methods are available?

```
> methods(class="gg")
[1] +
```

This is what makes layers work with ‘+’

```
> methods(class="ggplot")
[1] grid.draw  plot  print  summary
```

Normal methods for plot-type objects.  
`summary()` gives some useful info

# Playfair: Balance of trade charts

In the *Commercial and Political Atlas*, William Playfair used charts of imports and exports from England to its trading partners to ask “How are we doing”?

Here is a re-creation of one example, using ggplot2. How was it done?

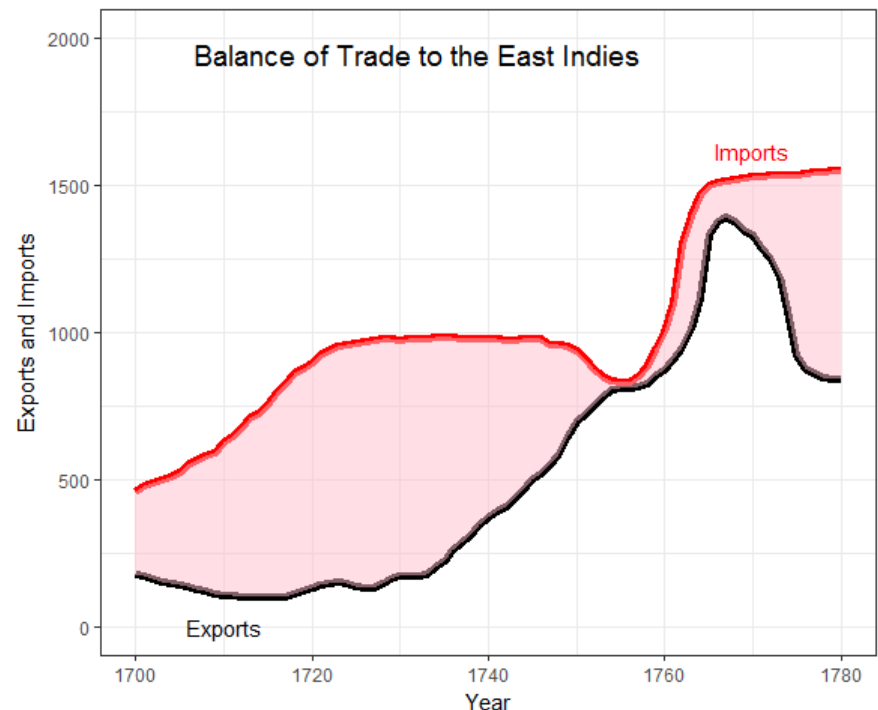
```
> data(EastIndiesTrade, package="GDAdata")
```

```
> head(EastIndiesTrade)
```

	Year	Exports	Imports
1	1700	180	460
2	1701	170	480
3	1702	160	490
4	1703	150	500
5	1704	145	510
6	1705	140	525
...	...	...	...

ggplot thinking:

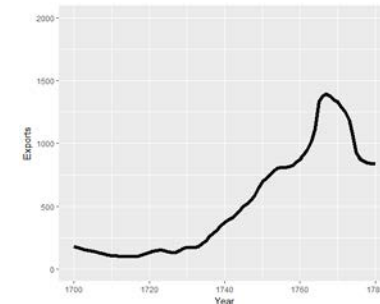
- what are the elements of this graph?
- how can I do them?



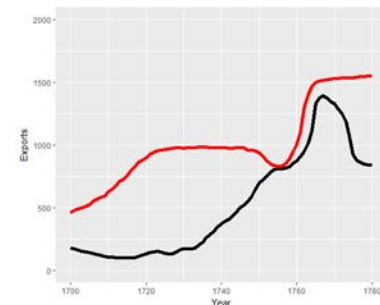
# ggplot thinking

I want to plot two time series, & fill the area between them

- Start with a line plot of Exports vs. Year: `geom_line()`
- Add a layer for the line plot of Imports vs. Year

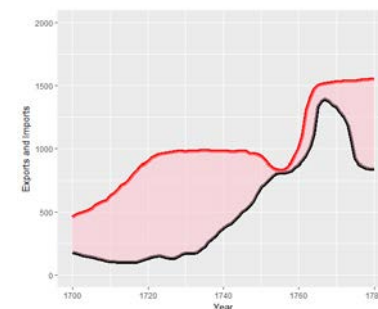


```
c1 <-  
ggplot(EastIndiesTrade, aes(x=Year, y=Exports)) +  
  ylim(0,2000) +  
  geom_line(colour="black", size=2) +  
  geom_line(aes(x=Year, y=Imports), colour="red", size=2)
```



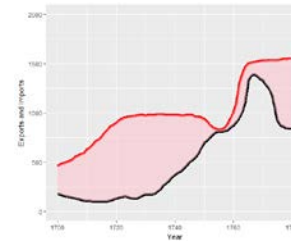
- Fill the area between the curves: `geom_ribbon()`
- change the Y label

```
c1 <- c1 +  
  geom_ribbon(aes(ymin=Exports, ymax=Imports), fill="pink") +  
  ylab("Exports and Imports")
```





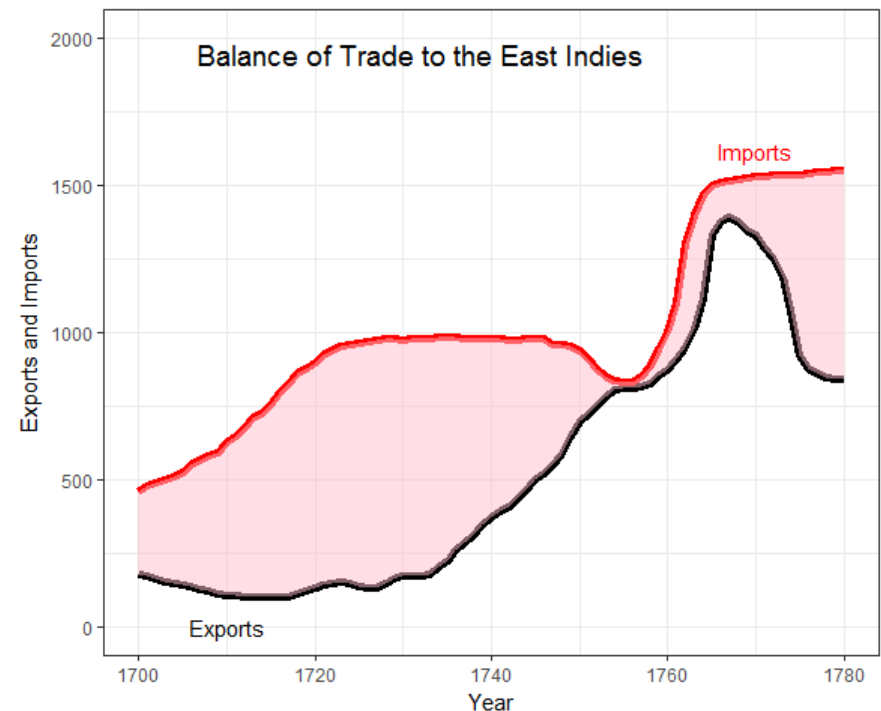
This looks pretty good.  
Add some text labels using `annotate()`



```
c1 <- c1 +  
  annotate("text", x = 1710, y = 0, label = "Exports", size=4) +  
  annotate("text", x = 1770, y = 1620, label = "Imports", color="red", size=4) +  
  annotate("text", x = 1732, y = 1950, label = "Balance of Trade to the East Indies", color="black", size=5)
```

Finally, change the theme to b/w

```
c1 <- c1 + theme_bw()
```



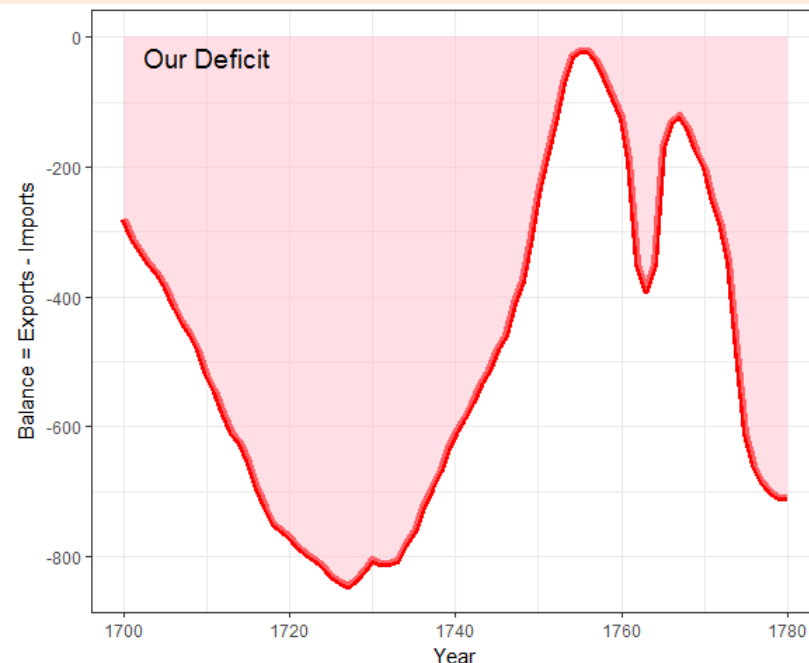
# Plot what you want to show

Playfair's goal was to show the balance of trade with different countries.  
Why not plot Exports – Imports directly?

```
c2 <-  
ggplot(EastIndiesTrade, aes(x=Year, y=Exports-Imports)) +  
  geom_line(colour="red", size=2) +  
  ylab("Balance = Exports - Imports") +  
  geom_ribbon(aes(ymin=Exports-Imports, ymax=0), fill="pink",alpha=0.5) +  
  annotate("text", x = 1710, y = -30, label = "Our Deficit", color="black", size=5) +  
  theme_bw()
```

## Questions:

- what are the basic plot variables?
- how did I make it shade above the curve?

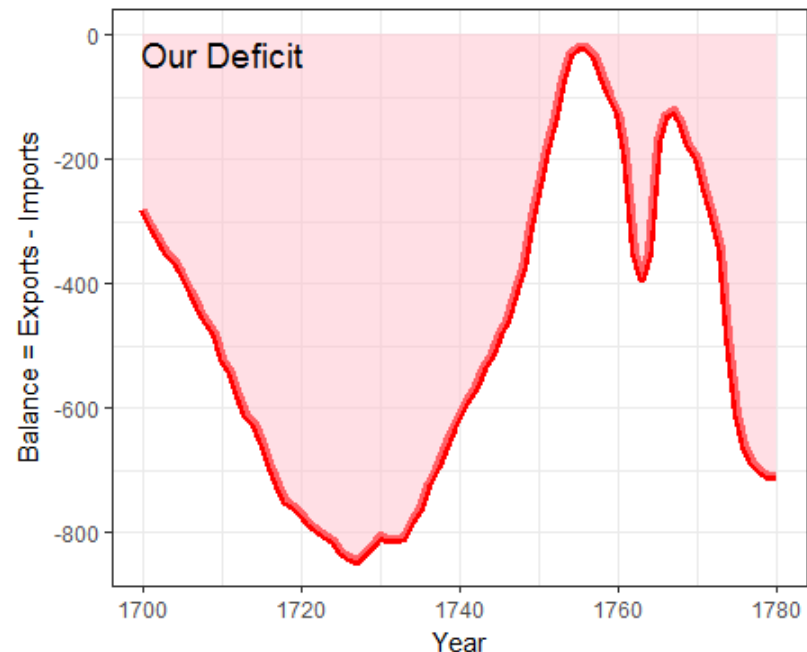
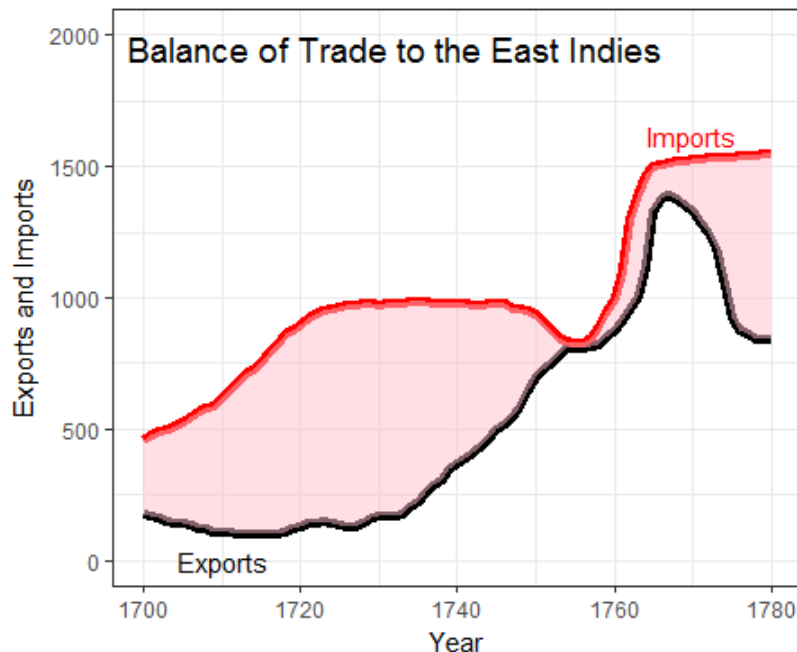


# Composing several plots

ggplot objects use grid graphics for rendering

The gridExtra package has functions for combining or manipulating grid-based graphs

```
library(gridExtra)  
grid.arrange(c1, c2, nrow=1)
```



# Saving plots: ggsave()

- If the plot is on the screen

```
ggsave("path/filename.png")
```

- If you have a plot object

```
ggsave(myplot, file="path/filename.png")
```

- Specify size:

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

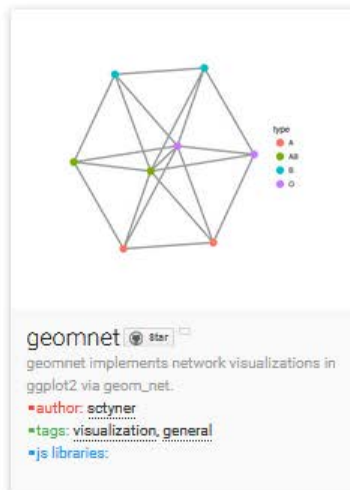
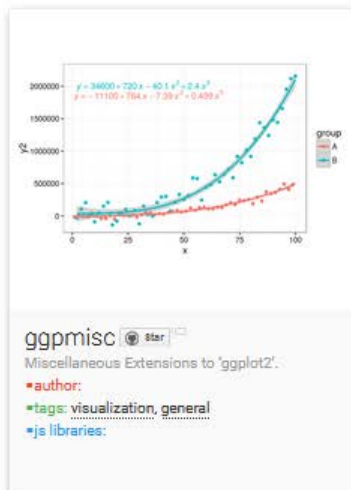
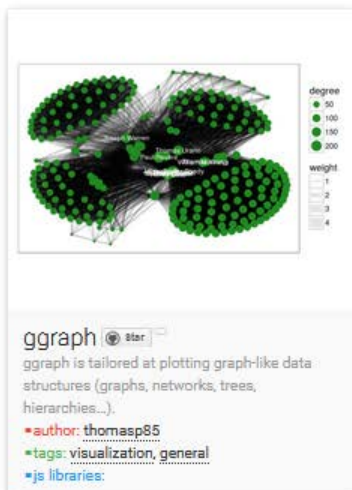
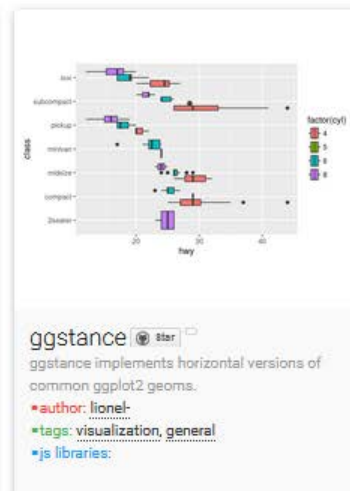
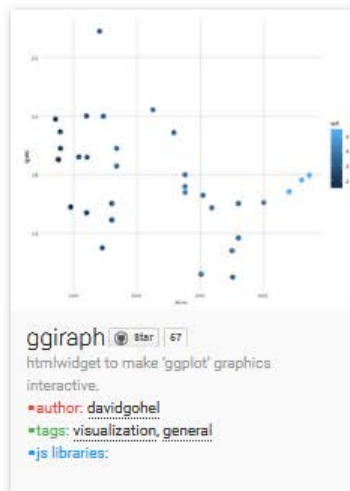
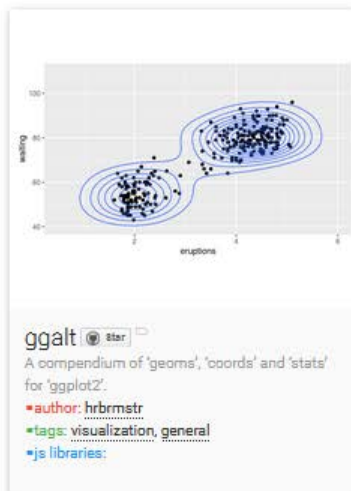
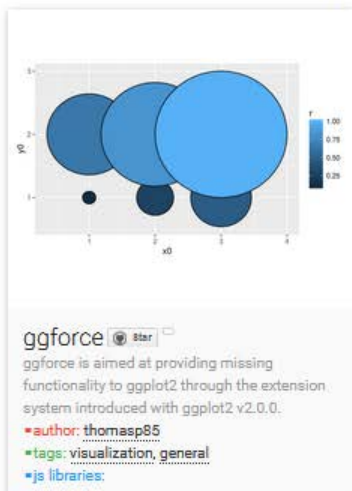
- any plot format (pdf, png, eps, svg, jpg, ...)

```
ggsave(myplot, file="path/filename.jpg")
```

```
ggsave(myplot, file="path/filename.pdf")
```

# ggplot extensions

There are a large number of ggplot extensions. See: <http://www.ggplot2-exts.org/>



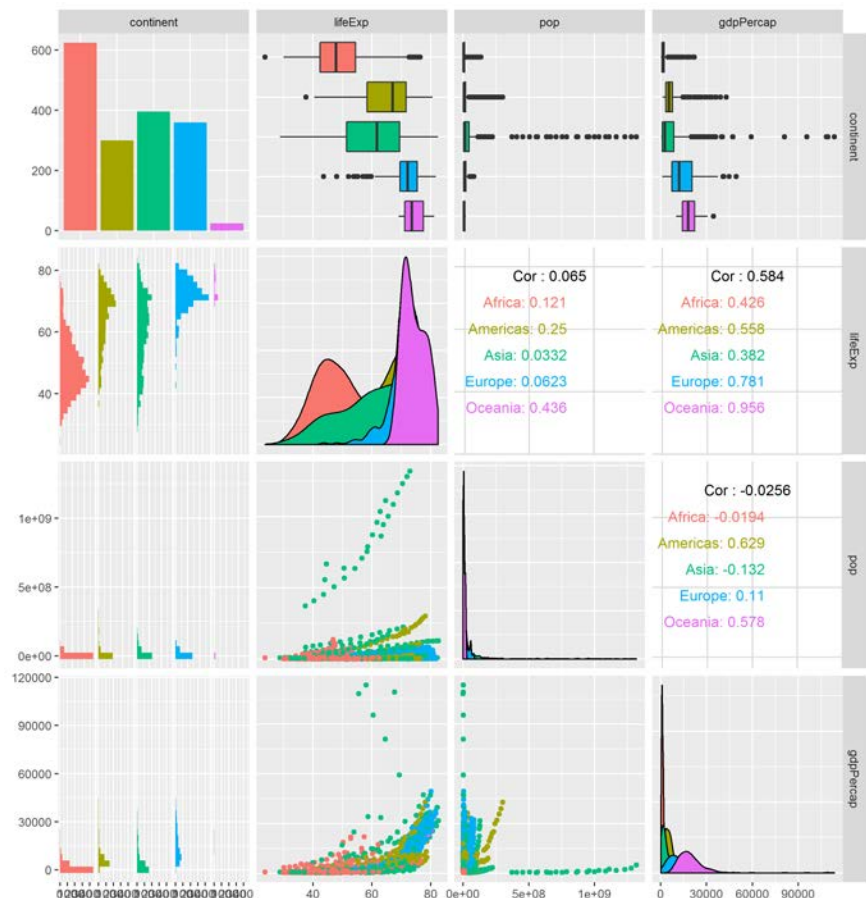
# ggplot extensions: GGally

GGally contains a large number of functions that extend `ggplot2` to multivariate data

`ggpairs()` produces generalized scatterplot matrices, with lots of options

```
library(GGally)
library(dplyr)
library(ggplot2)
library(gapminder)
```

```
gapminder %>%
  select(-country, -year) %>%
  ggpairs(aes(color=continent))
```



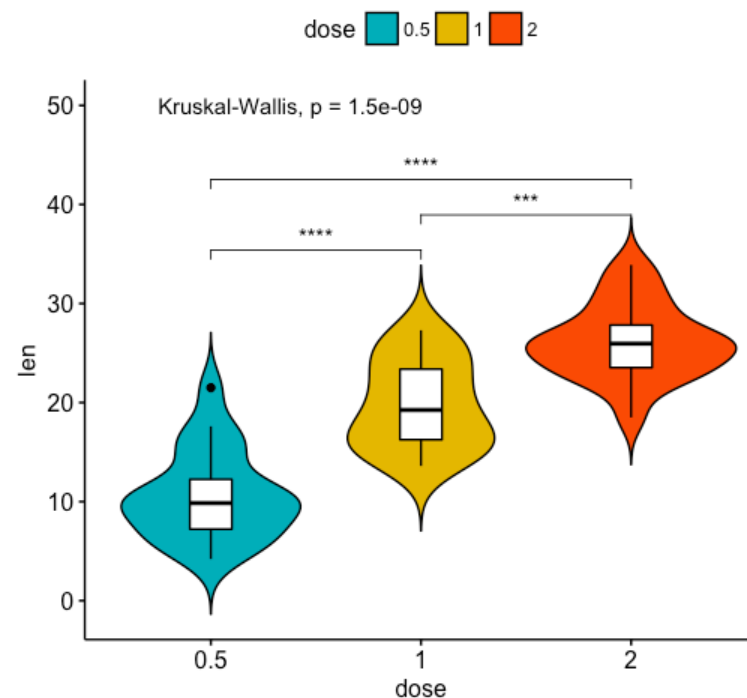
# ggpubr

The **ggpubr** package provides some easy-to-use functions for creating and customizing publication ready plots.

```
ggviolin(df, x = "dose", y = "len", fill = "dose",  
  palette = c("#00AFBB", "#E7B800", "#FC4E07"),  
  add = "boxplot", add.params = list(fill = "white")) +  
  stat_compare_means(comparisons = my_comparisons, label = "p.signif") +  
  stat_compare_means(label.y = 50)
```

see the examples at

<http://www.sthda.com/english/rpkgs/ggpubr/>

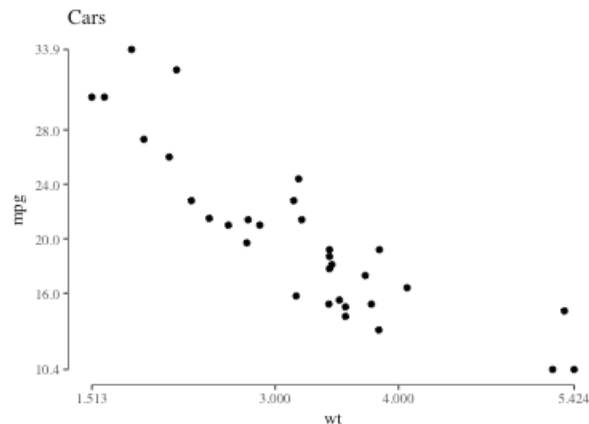


# ggthemes

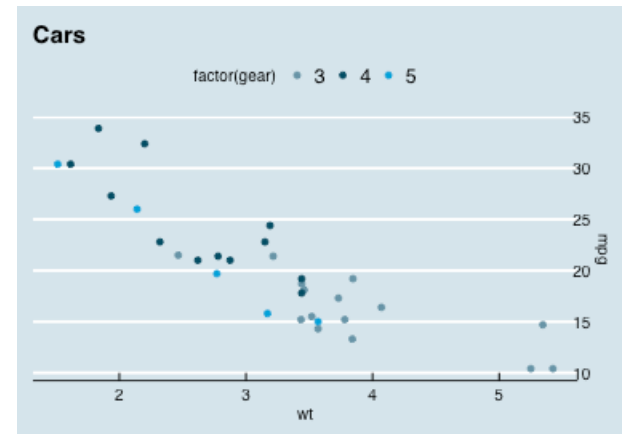
ggthemes provides a large number of extra  
geoms, scales, and themes for ggplot

```
install.packages('ggthemes', dependencies = TRUE)
```

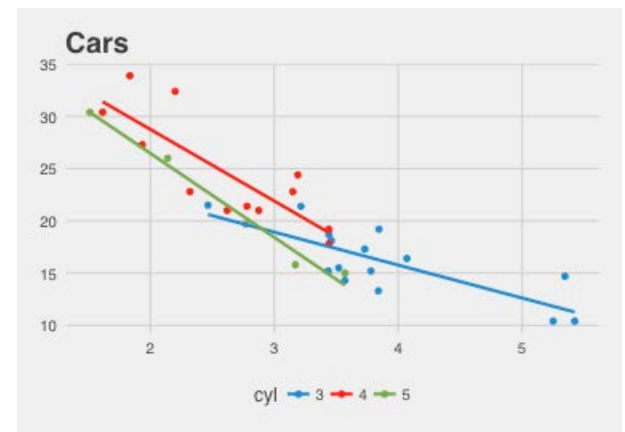
+ theme\_tufte()



+ theme\_economist()



+ theme\_fivethirtyeight()





# Tables in R

- Not a ggplot topic, but it is useful to know that you can also produce beautiful tables in R
- There are many packages for this: See the CRAN Task View on Reproducible Research, <https://cran.r-project.org/web/views/ReproducibleResearch.html>
  - xtable: Exports tables to LaTeX or HTML, with lots of control
  - stargazer: Well-formatted model summary tables, side-by-side
  - apaStyle: Generate APA Tables for MS Word
- Every time you cut & paste ...
  - ... God kills a kitten



# Tables in R: xtable

Just a few examples, stolen from xtable: vignette(“xtableGallery.pdf”)

```
fm1 <- aov(tlimth ~ sex + ethnicity + grade + disadvg, data = tli)
xtable(fm1)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
sex	1	75.37	75.37	0.38	0.5417
ethnicity	3	2572.15	857.38	4.27	0.0072
grade	1	36.31	36.31	0.18	0.6717
disadvg	1	59.30	59.30	0.30	0.5882
Residuals	93	18682.87	200.89		

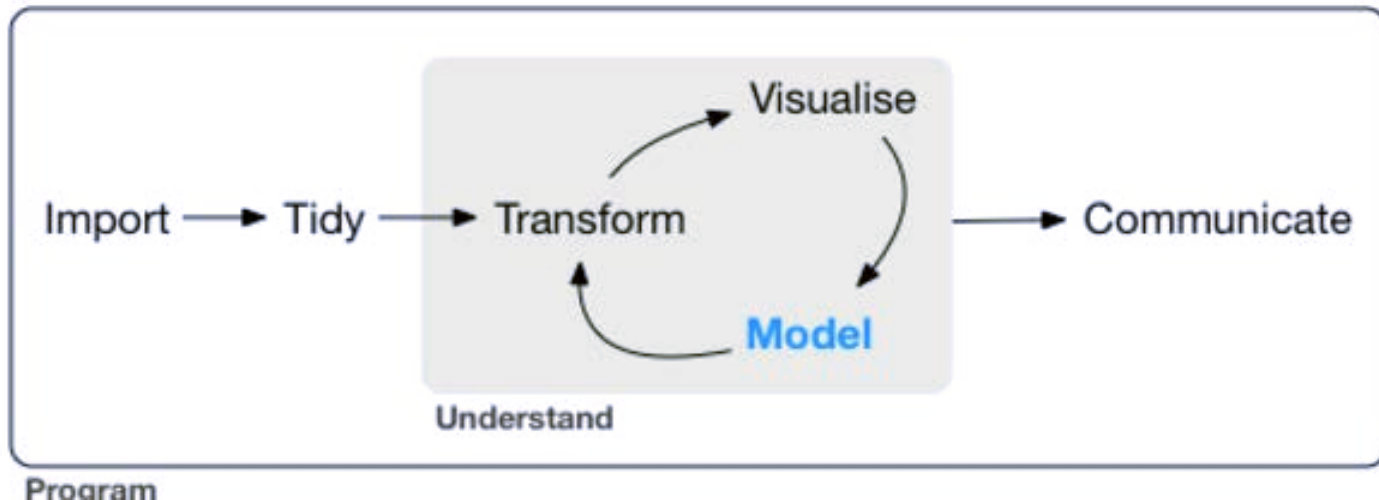
```
fm3 <- glm(disadvg ~ ethnicity*grade, data = tli, family = binomial)
xtable(fm3)
```

	Estimate	Std. Error	z value	Pr(> z )
(Intercept)	3.1888	1.5966	2.00	0.0458
ethnicityHISPANIC	-0.2848	2.4808	-0.11	0.9086
ethnicityOTHER	212.1701	22122.7093	0.01	0.9923
ethnicityWHITE	-8.8150	3.3355	-2.64	0.0082
grade	-0.5308	0.2892	-1.84	0.0665
ethnicityHISPANIC:grade	0.2448	0.4357	0.56	0.5742
ethnicityOTHER:grade	-32.6014	3393.4687	-0.01	0.9923
ethnicityWHITE:grade	1.0171	0.5185	1.96	0.0498

Too many decimals are used here, but you can control all that

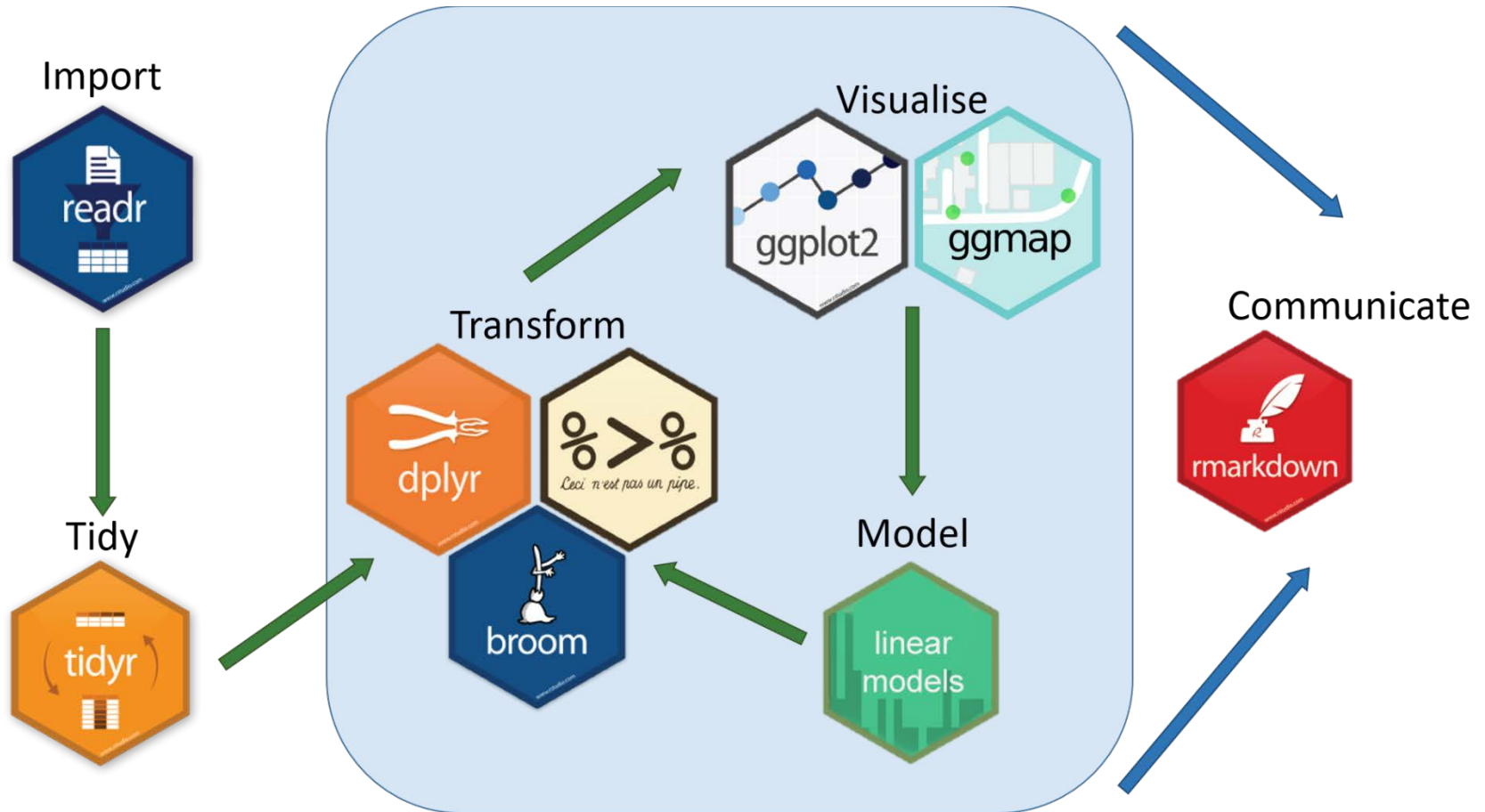
# A larger view: Data science

- Data science treats statistics & data visualization as parts of a larger process
  - Data import: text files, data bases, web scraping, ...
  - Data cleaning → “tidy data”
  - Model building & visualization
  - Reproducible report writing





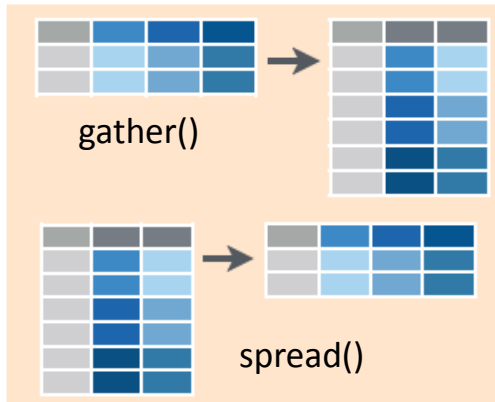
# The tidyverse of R packages



# Tidy tools: overview

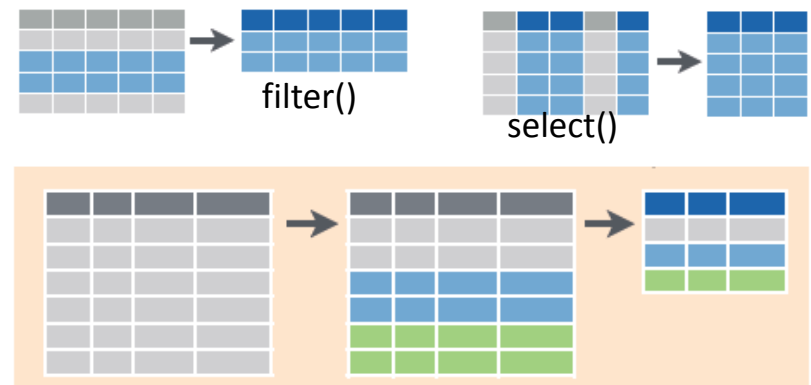


## tidyr



Reshape data to be tidy

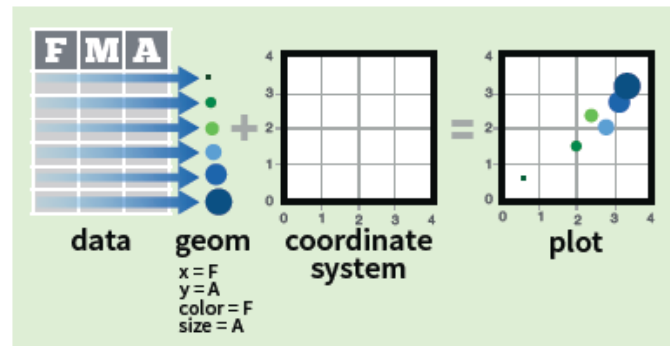
## dplyr



Manipulate & summarize tidy data



## ggplot2



Visualize me!

# Data wrangling with dplyr & tidyr

## What is Tidy Data?

A dataset is said to be tidy if:

- observations are in **rows**
- variables are in **columns**
- each value is in its own **cell**.

country	year	cases	population
Afghanistan	1999	18215	18000000
Afghanistan	2000	18666	20095360
Brazil	1999	31737	172006362
Brazil	2000	80488	174004898
China	1999	212258	1272015272
China	2000	212966	1280000583

variables

country	year	cases	population
Afghanistan	1999	18215	18000000
Afghanistan	2000	18666	20095360
Brazil	1999	31737	172006362
Brazil	2000	80488	174004898
China	1999	212258	1272015272
China	2000	212966	1280000583

observations

country	year	cases	population
Afghanistan	1999	18215	18000000
Afghanistan	2000	18666	20095360
Brazil	1999	31737	172006362
Brazil	2000	80488	174004898
China	1999	212258	1272015272
China	2000	212966	1280000583

values

A “messy” dataset: Survey of income by religion from Pew Research

- Values of **income** are in separate columns
- Column headers are **values**, not variable names
- Cell values are frequencies--- **implicit**, not explicit

religion	<\$10k	\$10-20k	\$20-30k	\$30-40k	\$40-50k	\$50-75k
Agnostic	27	34	60	81	76	137
Atheist	12	27	37	52	35	70
Buddhist	27	21	30	34	33	58
Catholic	418	617	732	670	638	1116

This organization is easy in Excel

But, this makes data analysis and graphing hard

# Tidying: reshaping wide to long

We can tidy the data by reshaping from wide to long format using `tidyr::gather()`

```
> pew <- read.delim(
  file = "http://stat405.had.co.nz/data/pew.txt",
  header = TRUE,
  stringsAsFactors = FALSE, check.names = FALSE)

> (pew1 <- pew[1:4, 1:6]) # small subset

  religion <$10k $10-20k $20-30k $30-40k $40-50k
1 Agnostic    27      34      60      81      76
2 Atheist     12      27      37      52      35
3 Buddhist    27      21      30      34      33
4 Catholic   418     617     732     670     638
```

Another solution, using `reshape2::melt()`

```
> library(reshape2)
> pew_tidy <- melt(
  data = pew1,
  id = "religion",
  variable.name = "income",
  value.name = "frequency"
)
```

key      value      columns

↓      ↓      ↓

```
> library(tidyr)
> gather(pew1, "income", "frequency", 2:6)

  religion income frequency
1 Agnostic <$10k      27
2 Atheist  <$10k      12
3 Buddhist <$10k      27
4 Catholic <$10k     418
5 Agnostic $10-20k     34
6 Atheist  $10-20k     27
7 Buddhist $10-20k     21
8 Catholic $10-20k    617
9 Agnostic $20-30k     60
10 Atheist  $20-30k     37
11 Buddhist $20-30k     30
12 Catholic $20-30k    732
13 Agnostic $30-40k     81
14 Atheist  $30-40k     52
15 Buddhist $30-40k     34
16 Catholic $30-40k    670
...      ...      ...
```

NB: income is a character variable; we might want to create an ordered factor or numeric version



# Using pipes: %>%

- R is a functional language

- This means that  $f(x)$  returns a value, as in  $y \leftarrow f(x)$
- That value can be passed to another function:  $g(f(x))$
- And so on:  $h(g(f(x)))$

```
> x <- c(0.109, 0.359, 0.63, 0.996, 0.515, 0.142)
> exp(diff(log(x)))
[1] 3.29 1.75 1.58 0.52 0.28
```

- This gets messy and hard to read, unless you break it down step by step

```
> # Compute the logarithm of `x`, calculate lagged differences,
> # return the exponential function of the result
> log(x)
[1] -2.216 -1.024 -0.462 -0.004 -0.664 -1.952
> diff(log(x))
[1] 1.19 0.56 0.46 -0.66 -1.29
> exp(diff(log(x)))
[1] 3.29 1.75 1.58 0.52 0.28
```



# Using pipes: %>%

- Pipes (%>%) change the syntax to make this easier

```
> # use pipes  
> x %>% log() %>% diff() %>% exp()  
[1] 3.29 1.75 1.58 0.52 0.28
```

- Basic rules

- $x \%>\% f()$  passes object on left hand side as **first** argument (or . argument) of function on right hand side
  - $x \%>\% f()$  is the same as  $f(x)$
  - $x \%>\% f(y)$  is the same as  $f(x, y)$
  - $y \%>\% f(x, ., z)$  is the same as  $f(x, y, z)$
- $x \%<>\% f()$  does the same, but assigns the result to x
  - Shortcut for  $x \leftarrow x \%>\% f()$

# dplyr: Subset observations (rows)

dplyr implements a variety of verbs to select a subset of observations from a dataset



In a pipe expression, omit the dataset name

**dplyr::filter(iris, Sepal.Length > 7)**

Extract rows that meet logical criteria.

**dplyr::distinct(iris)**

Remove duplicate rows.

**dplyr::sample\_frac(iris, 0.5, replace = TRUE)**

Randomly select fraction of rows.

**dplyr::sample\_n(iris, 10, replace = TRUE)**

Randomly select n rows.

**dplyr::slice(iris, 10:15)**

Select rows by position.

**dplyr::top\_n(storms, 2, date)**

Select and order top n entries (by group if grouped data).

```
iris %>% filter(Sepal.Length > 7)
iris %>% filter(Species == "setosa")
```

```
iris %>% sample_n(10)
iris %>% slice(1:50) # setosa
```

# dplyr: Subset variables (columns)



```
dplyr::select(iris, Sepal.Width, Petal.Length, Species)
```

Select columns by name or helper function.

Many helper functions in dplyr allow selection by a **function** of variable names:

```
select(iris, contains("."))
```

Select columns whose name contains a character string.

```
select(iris, ends_with("Length"))
```

Select columns whose name ends with a character string.

```
select(iris, everything())
```

Select every column.

```
select(iris, matches(".t."))
```

Select columns whose name matches a regular expression.

```
select(iris, num_range("x", 1:5))
```

Select columns named x1, x2, x3, x4, x5.

```
select(iris, one_of(c("Species", "Genus")))
```

Select columns whose names are in a group of names.

```
select(iris, starts_with("Sepal"))
```

Select columns whose name starts with a character string.

```
select(iris, Sepal.Length:Petal.Width)
```

Select all columns between Sepal.Length and Petal.Width (inclusive).

```
select(iris, -Species)
```

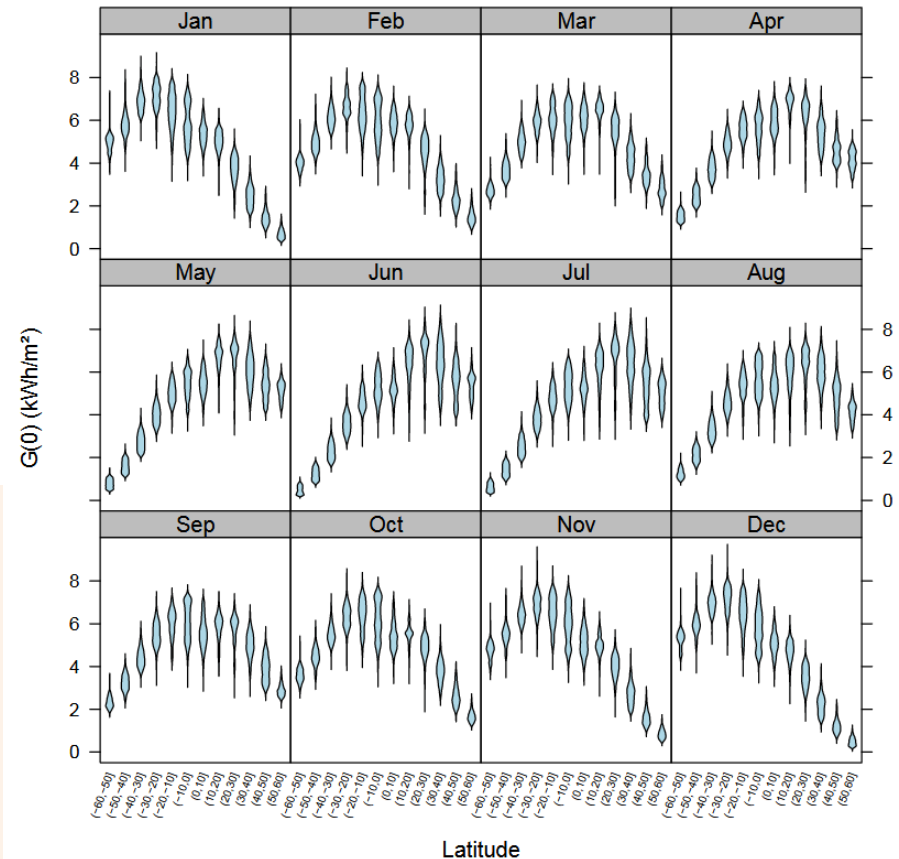
Select all columns except Species.

# Faceting & tidy data

Here is a complex graph, showing distributions of solar radiation from NASA, by months of the year and latitude

This is complicated, because the data structure is **untidy**--- months were in separate variables (wide format)

```
> str(nasa)
'data.frame': 64800 obs. of 15 variables:
 $ Lat: int -90 -90 -90 -90 -90 -90 -90 -90 -90 -90 ...
 $ Lon: int -180 -179 -178 -177 -176 -175 -174 -173 -172 -171 ...
 $ Jan: num 9.63 9.63 9.63 9.63 9.63 9.63 9.63 9.63 9.63 9.63 ...
 $ Feb: num 5.28 5.28 5.28 5.28 5.28 5.28 5.28 5.28 5.28 5.28 ...
 $ Mar: num 0.75 0.75 0.75 0.75 0.75 0.75 0.75 0.75 0.75 0.75 ...
 $ Apr: num 0 0 0 0 0 0 0 0 0 0 ...
 $ May: num 0 0 0 0 0 0 0 0 0 0 ...
 $ Jun: num 0 0 0 0 0 0 0 0 0 0 ...
 $ Jul: num 0 0 0 0 0 0 0 0 0 0 ...
 $ Aug: num 0 0 0 0 0 0 0 0 0 0 ...
 $ Sep: num 0.1 0.1 0.1 0.1 0.1 0.1 0.1 0.1 0.1 0.1 ...
 $ Oct: num 3.24 3.24 3.24 3.24 3.24 3.24 3.24 3.24 3.24 3.24 ...
 $ Nov: num 8.28 8.28 8.28 8.28 8.28 8.28 8.28 8.28 8.28 8.28 ...
 $ Dec: num 11 11 11 11 11 ...
 $ Ann: num 3.19 3.19 3.19 3.19 3.19 3.19 3.19 3.19 3.19 3.19 ...
```



Each distribution is shown as a **violin plot**, a mirrored density plot

# tidying the data

In wide format, using lattice, I had to construct a plot formula to plot those columns

```
> x <- paste(names(nasa)[3:14], collapse='+')  
> (formula <- as.formula(paste(x, '~cut(Lat, pretty(Lat, 20))', sep=")))  
Jan + Feb + Mar + Apr + May + Jun + Jul + Aug + Sep + Oct + Nov +  
Dec ~ cut(Lat, pretty(Lat, 20))
```

Ugh!

It is much easier to reshape the data to long format, so solar is all in one column

```
library(tidyr)  
library(dplyr)  
library(ggplot2)
```

```
nasa_long <- nasa %>%  
  select(-Ann) %>%  
  gather(month, solar, Jan:Dec, factor_key=TRUE) %>%  
  filter( abs(Lat) < 60 ) %>%  
  mutate( Lat_f = cut(Lat, pretty(Lat, 12)))
```

%>% “pipes” data to the next stage

**select()** extracts or drops columns

**gather()** collapses columns into key-value pairs

**filter()** subsets observations

**mutate()** creates new variables

# tidying the data

```
> str(nasa_long)
'data.frame': 514080 obs. of 5 variables:
 $ Lat : int -59 -59 -59 -59 -59 -59 -59 -59 -59 -59 ...
 $ Lon : int -180 -179 -178 -177 -176 -175 -174 -173 -172 -171 ...
 $ month: Factor w/ 12 levels "Jan","Feb","Mar",...: 1 1 1 1 1 1 1 1 1 1 ...
 $ solar: num 5.19 5.19 5.25 5.25 5.17 5.17 5.15 5.15 5.15 5.15 ...
 $ Lat_f: Factor w/ 12 levels "(-60,-50]", "(-50,-40]",...: 1 1 1 1 1 1 1 1 1 1 ...
```

solar is now the single response variable

For ease of plotting, I created a factor version of Lat with 12 levels

```
> head(nasa_long)
  Lat Lon month solar  Lat_f
1 -59 -180  Jan  5.19 (-60,-50]
2 -59 -179  Jan  5.19 (-60,-50]
3 -59 -178  Jan  5.25 (-60,-50]
4 -59 -177  Jan  5.25 (-60,-50]
5 -59 -176  Jan  5.17 (-60,-50]
6 -59 -175  Jan  5.17 (-60,-50]
```

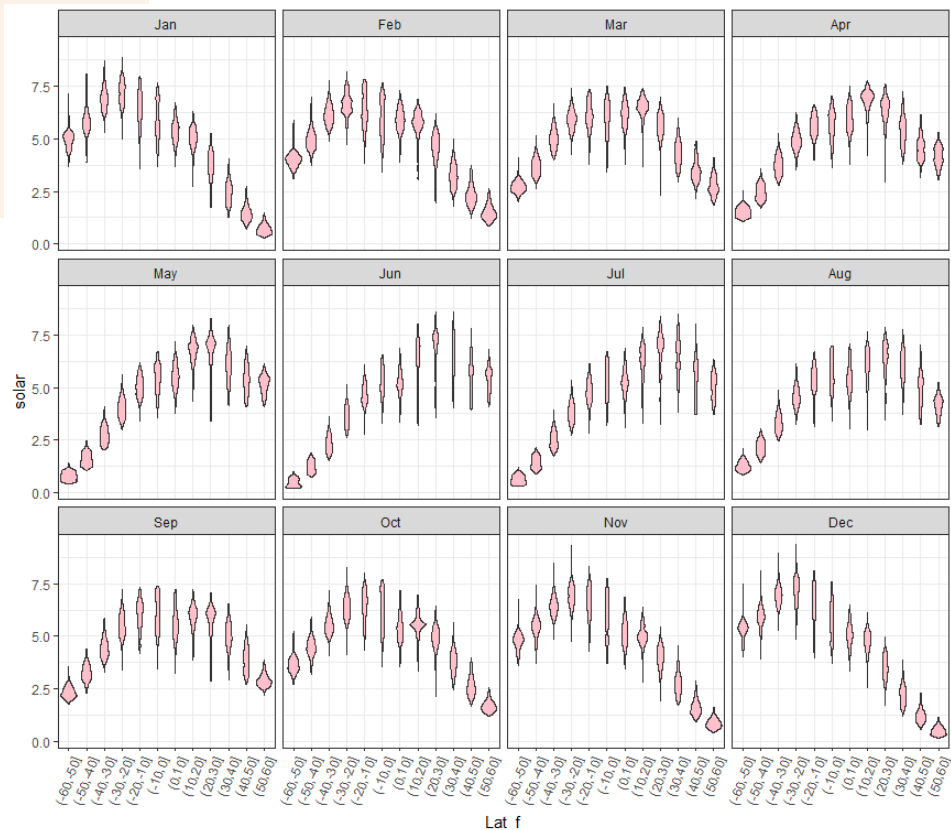
The data are now in a form where I can plot solar against Lat or Lat\_f and facet by month

# plotting the tidy data

Using `geom_violin()` shows the shapes of the distributions for levels of `Lat_f`

```
ggplot(nasa_long, aes(x=Lat_f, y=solar)) +  
  geom_violin(fill="pink") +  
  facet_wrap(~ month) +  
  theme_bw() +  
  theme(axis.text.x =  
    element_text(angle = 70,  
                  hjust = 1))
```

`facet_wrap(~month)` does the right thing



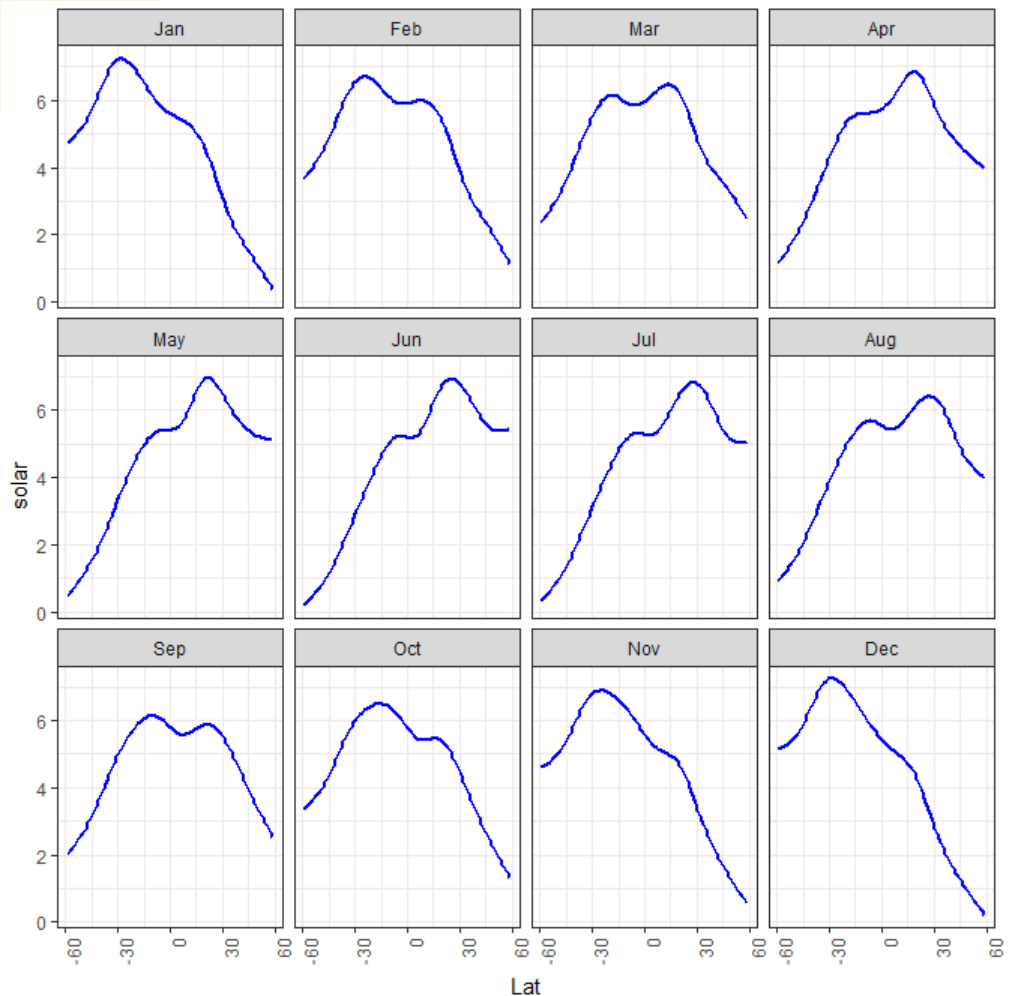
I had to adjust the x-axis labels for `Lat_f` to avoid overplotting

# plotting the tidy data: smoothing

```
ggplot(nasa_long, aes(x=Lat, y=solar)) +  
  geom_smooth(color="blue" ) +  
  facet_wrap(~ month) +  
  theme_bw()
```

Here I treat Lat as quantitative  
`geom_smooth()` uses method =  
“gam” here because of large  $n$

The variation in the smoothed  
trends over the year suggest  
quite lawful behavior





# build a model

What we saw in the plot suggests a generalized additive model, with a smooth,  $s(\text{Lat})$

```
library(mgcv)
nasa.gam <- gam(solar ~ Lon + month + s(Lat), data=nasa_long)
summary(nasa.gam)
```

Family: gaussian  
Link function: identity

Formula:  
solar ~ Lon + month + s(Lat)

Parametric coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	4.691e+00	6.833e-03	686.409	< 2e-16 ***
Lon	-1.713e-04	1.898e-05	-9.022	< 2e-16 ***
monthFeb	1.195e-01	9.664e-03	12.364	< 2e-16 ***

monthDec	-8.046e-02	9.664e-03	-8.326	< 2e-16 ***
----------	------------	-----------	--------	-------------

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	F	p-value
s(Lat)	8.997	9	37285	<2e-16 ***

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.398 Deviance explained = 39.8%

GCV = 2.0006 Scale est. = 2.0005 n = 514080

The violin plots suggest that variance is not constant. I'm ignoring this here by using the default gaussian model. (Good first start)

Model terms:

- Lon wasn't included before
- month is a factor, for the plots
- $s(\text{Lat})$  fits a smoothed term in latitude, averaged over other factors

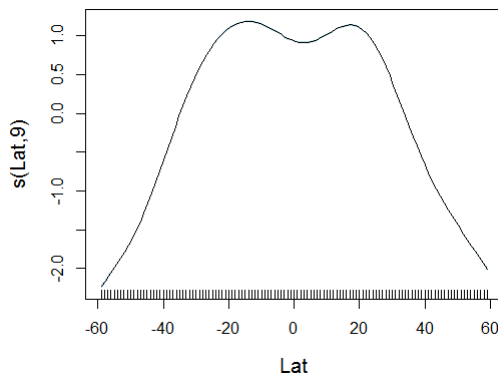
There are other model choices, but it is useful to visualize what we have done so far

# visualize the model

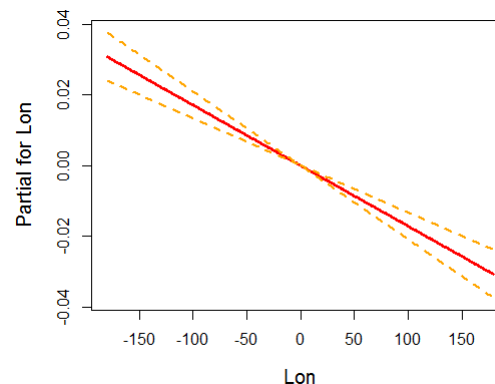
Effect plots show the fitted relationship between the response and model terms, averaged over other predictors.

The mgcv package has its own versions of these.

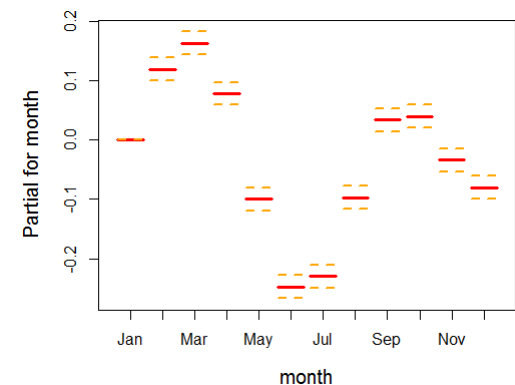
```
plot(nasa.gam, cex.lab=1.25)
termplot(nasa.gam, terms="month", se=TRUE, lwd.term=3, lwd.se=2, cex.lab=1.25)
termplot(nasa.gam, terms="Lon", se=TRUE, lwd.term=3, lwd.se=2, cex.lab=1.25)
```



why the dip at the equator?



effect of longitude is very small, but maybe interpretable



month should be modeled as a time variable

# Summary

- ggplot2 provides a new way of thinking about graphs
  - aes() – mapping data variables to visual properties
  - geom\_() – drawing geometric objects (points, lines, ...)
  - coord\_() – transform coordinate systems
  - layers – add stuff to an existing plot with '+'
  - themes – change the entire look of a graph
- tidyr & dplyr provide a new way of thinking about data analysis
- R Studio tools provide a way to organize your work, do analysis, and publish --- reproducible!