

3. Plotting functions and formulas

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In this session

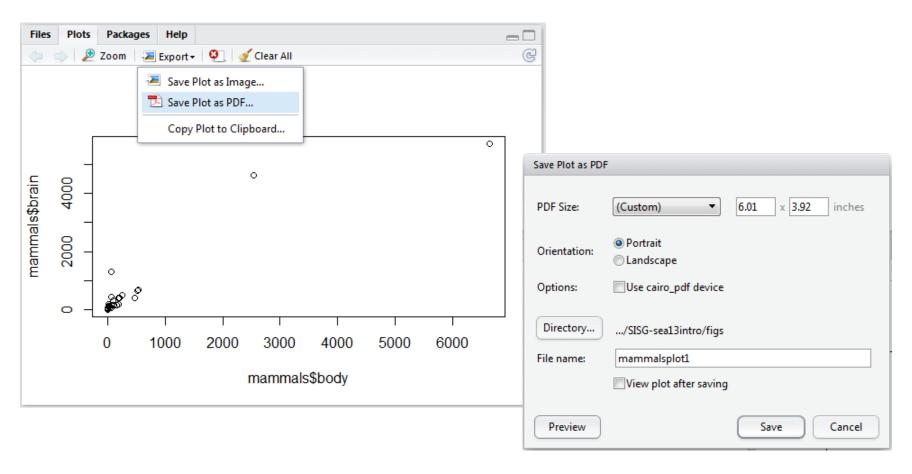
R is known for having good graphics — good for data exploration and summary, as well as illustrating analyses. Here, we wil see;

- Some generic plotting commands
- Making graphics files
- Fine-tuning your plots (and why not to do too much of this)
- The formula syntax

NB more graphics commands will follow, in the next session.

A first example, using the mammals dataset — and its output in the Plot window; (The preview button is recommended)

plot(x=mammals\$body, y=mammals\$brain)



Some other options for exporting;

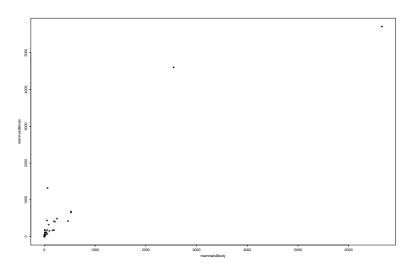
- Copy directly to clipboard as a bitmap or editable (Windows)
 metafile then paste into e.g. your Powerpoint slides
- With 'Save Plot as Image', PNG is a (good) bitmap format, suitable for line art, i.e. graphs. JPEG is good for photos, not so good for graphs
- For PNG/JPEG, previews disappear if they get too large!
- Many of the options (TIFF, EPS) are seldom used, today
- Handy hint; if too much re-sizing confuses your graphics device (i.e. the Plot window) enter dev.off() and just start over

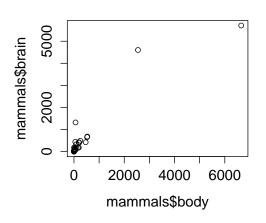
A golden rule for exporting;

Make the file the size it will be in the final document – because R is good at choosing font sizes

A 6:4 plot, saved at 24×16 inches

The same plot, saved at 4×2.67 inches





- Not the same plot 'blown up' note e.g. axes labels
- R likes to add white space around the edges good in documents, less good in slides, depending on your software

Better axes, better axis labels and a title would make the scatterplot better. But on looking up ?plot...

"For simple scatter plots, plot.default will be used. However, there are plot methods for many R objects, including functions, data.frameS, density objects, etc. Use methods(plot) and the documentation for these."

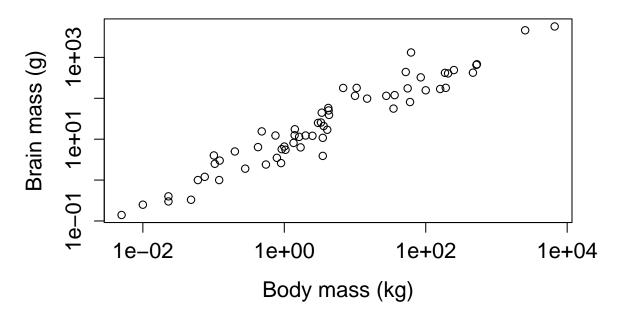
plot() is a *generic* function — it does different things given different input; see methods(plot) for a full list. For our plot of y vs x, the details we need are in ?plot.default...

After checking the help page to see what these mean, we use;

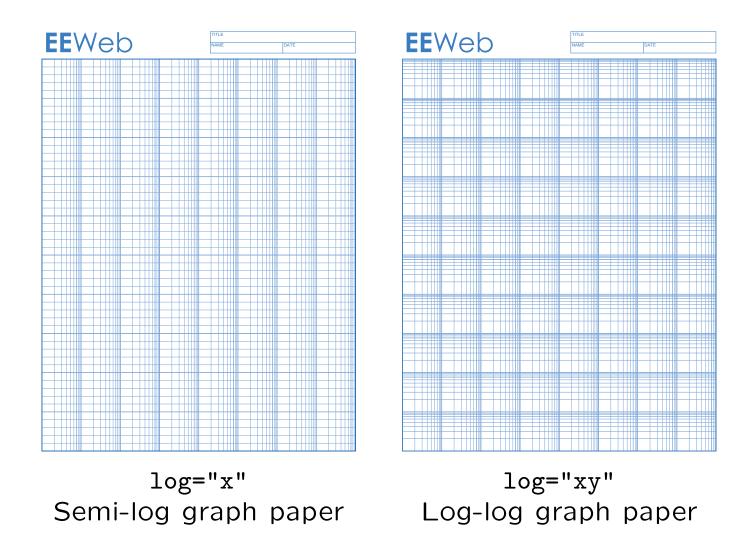
- xlab, ylab for the axis labels
- main for the main title
- log to log the axes − log="xy", to log them both

plot(x=mammals\$body, y=mammals\$brain, xlab="Body mass (kg)",
 ylab="Brain mass (g)", main="Brain and body mass, for 62 mammals",
 log="xy")

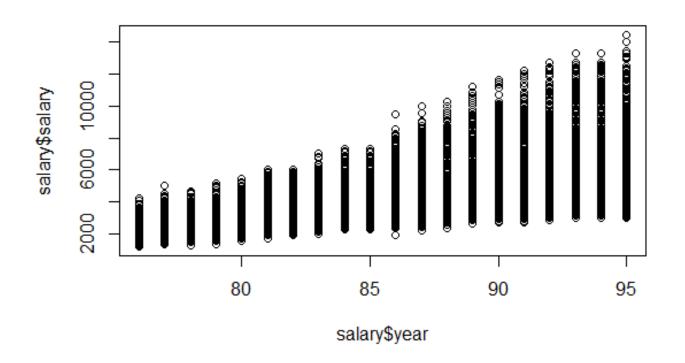
Brain and body mass, for 62 mammals



For those with historical interests (or long memories);

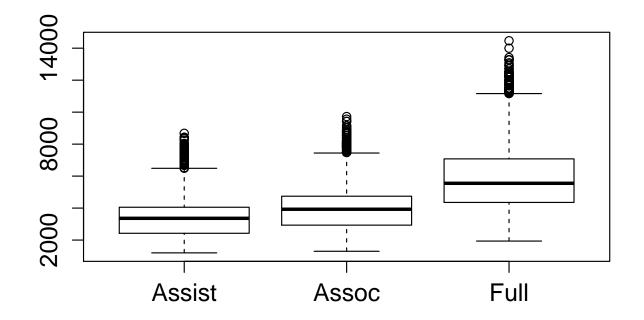


As the help file suggests, plot() gives different output for different types of input. First, another scatterplot; plot(x=salary\$year, y=salary\$salary)



Tip: export graphs of large datasets as PNG, not PDF or JPEG.

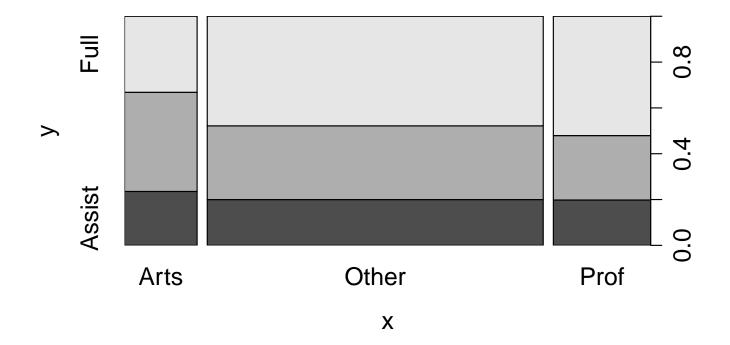
Plotting one numeric variable against a factor; plot(x=salary\$rank, y=salary\$salary)



There is also a boxplot() function.

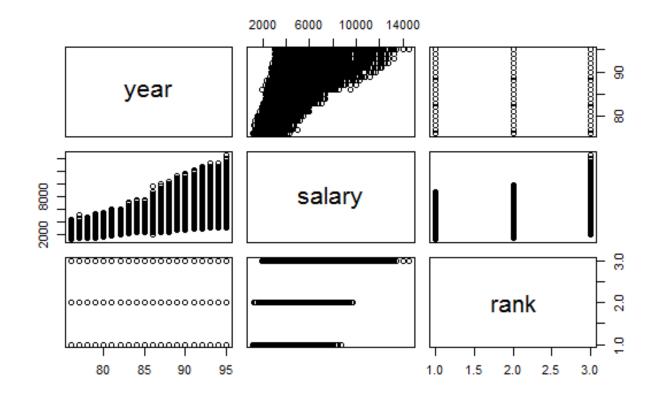
Plotting one factor variable against another;

plot(x=salary\$field, y=salary\$rank)



This is a *stacked barplot* — see also the barplot() function

Plotting an entire data frame (not too many columns) smallsalary <- salary[,c("year","salary","rank")] plot(smallsalary)

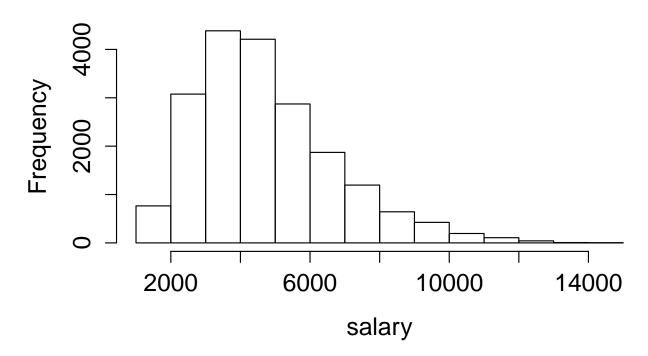


Not so clever! But quick, & okay if all numeric — see also pairs(). NB Plotting functions for large datasets are in later sessions.

For histograms, use hist();

hist(salary\$salary, main="Monthly salary", xlab="salary")

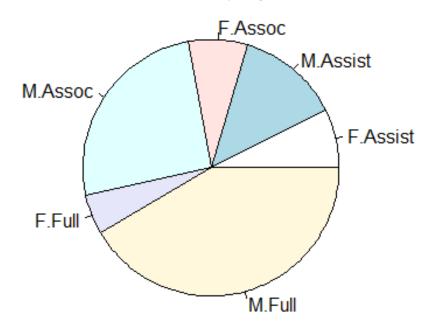
Monthly salary



For more control, set argument breaks to either a number, or a vector of the breakpoints.

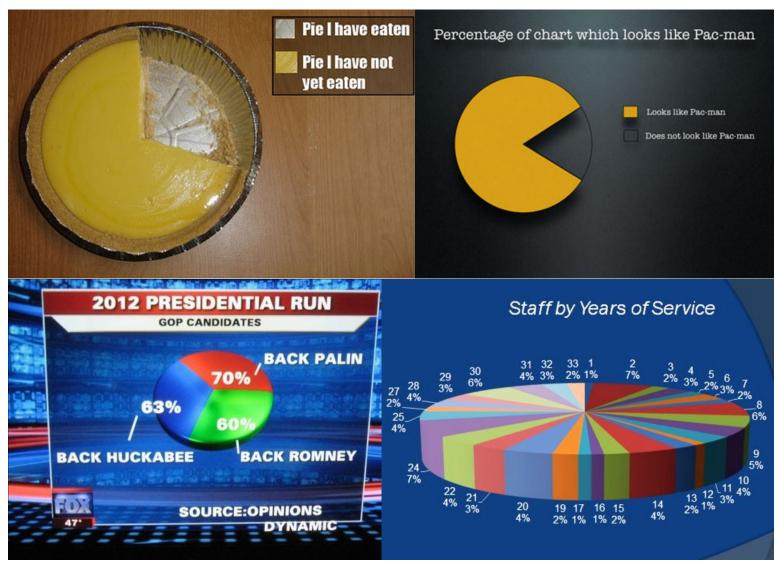
Please tell no-one I told you this one;

```
> table( interaction(salary$gender, salary$rank) )
F.Assist M.Assist F.Assoc M.Assoc F.Full M.Full
    1460    2588    1465    5064    1001    8210
> pie( table( interaction(salary$gender, salary$rank) ) )
```



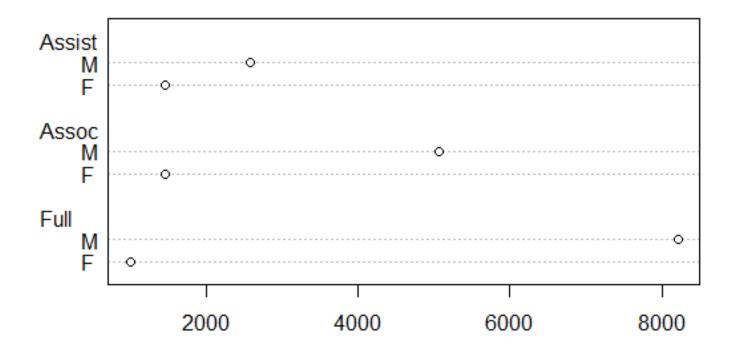
Why do statisticians hate pie charts with such passion?

... they really do!



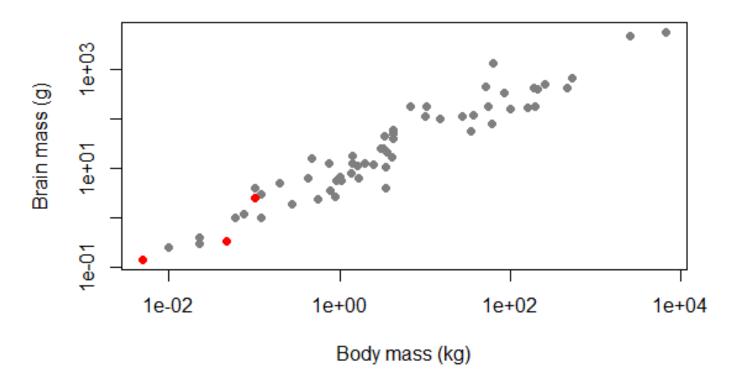
Because pie charts are a terrible way to present data. Dotcharts are *much* better – also easy to code;

dotchart(table(salary\$gender, salary\$rank))



See also stripchart(); with multiple symbols per line, these are a good alternative to boxplots, for small samples.

Suppose you want to highlight certain points on a scatterplot; other options to the plot() command change point style & color; > grep("shrew", row.names(mammals)) # or just look in Data viewer [1] 14 55 61 > is.shrew <- 1:62 %in% c(14,55,61) # 3 TRUEs and 59 FALSEs > plot(x=mammals\$body, y=mammals\$brain, xlab="Body mass (kg)", ylab="Brain mass (g)",log="xy", col=ifelse(is.shrew, "red", "gray50"), pch=19)



We used col=ifelse(is.shrew, "red", "gray50") — a vector of 3 reds and 59 gray50s.

- If we supply fewer colors than datapoints, what we supplied is recycled
- You could probably guess "red", "green", "purple" etc, but not "gray50". To find out the names of the (many) available R colors, use the colors() command – no arguments needed
- Can also specify colors by numbers; 1=black, 2=red,
 3=green up to 8, then it repeats
- Or consult this online chart or many others like it
- Can also supply colors by hexadecimal coding; #RRGGBB for red/green/blue – with #RRGGBBTT for transparency

NB legends will follow, in the next session.

We also used pch=19 - to obtain the same non-default plotting symbol, a filled circle.

The full range;

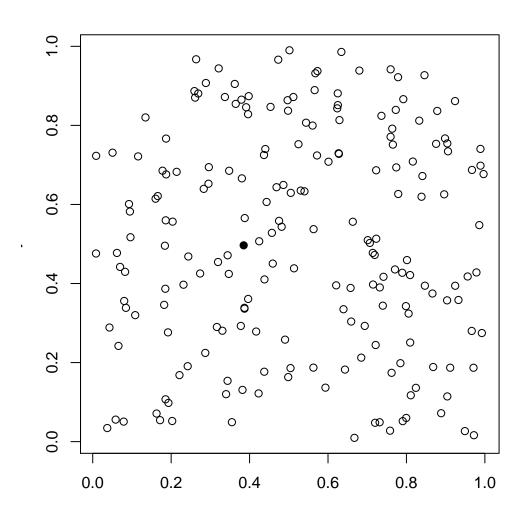
```
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 \bigcirc \triangle + \times \bigcirc \bigcirc \boxtimes \oplus \bigoplus \bigoplus \boxtimes \boxtimes \boxtimes \boxtimes \bigoplus \bigoplus \bigoplus \bigoplus \bigoplus \bigoplus \bigoplus \bigoplus
```

- Set the fill color for 21:25 with the bg argument
- The open circle (pch=1) is the default because it makes it easiest to see points that nearly overlap. Change it only if you have a good reason to
- Filled symbols 15:20 work well with transparent colors, e.g. col="#FF000033" for translucent pink

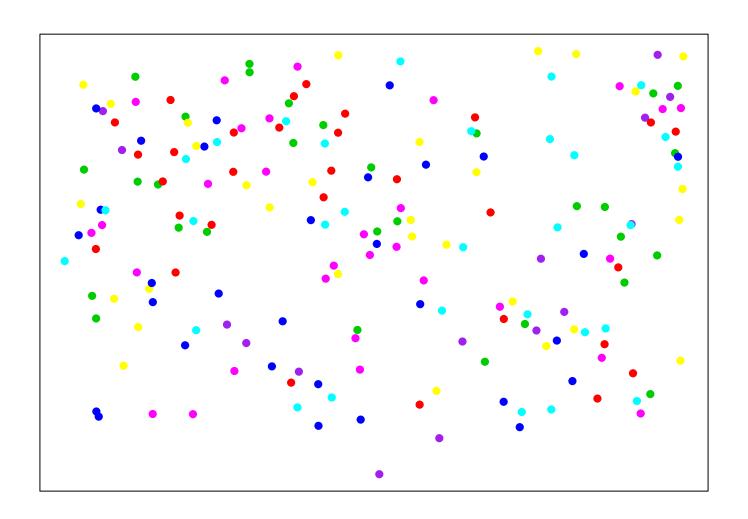
For different size symbols, there is a cex option; cex=1 is standard size, cex=1.5 is 50% bigger, etc.

But beware! These options should be used sparingly...

One of these points is not like the others...

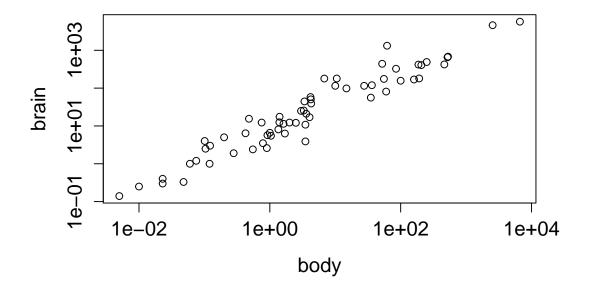


Too many colors (> 4, say) requires too much attention; what pattern is illustrated here?



To make plots, we've used arguments \mathbf{x} (on the X-axis) and \mathbf{y} (on the Y-axis). But another method makes a stronger connection to why we're making the plot;

plot(brain~body, data=mammals, log="xy")



"Plot how brain depends on body, using the mammals dataset, with logarithmic x and y axes"

A few more examples, using the salary dataset;

```
plot(salary~year, data=salary) # scatterplot
plot(salary~rank, data=salary) # boxplot
plot(rank~field, data=salary) # stacked barplot
```

In all of these, $Y \sim X$ can be interpreted as Y depends on X – the 'tilde' symbol is R's shorthand for 'depends on'.

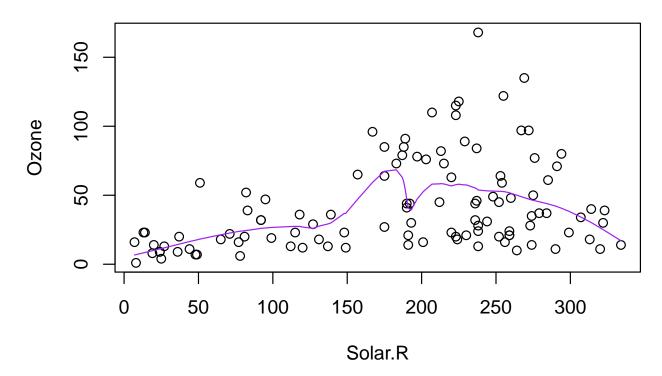
Statisticians (and scientists) like to think this way;

- How does some outcome (Y) depend on a covariate (X)? (a.k.a. a predictor)
- How does a dependent variable (Y) depend on an independent variable (X)?

And how does Y depend on X in observations with the same Z?

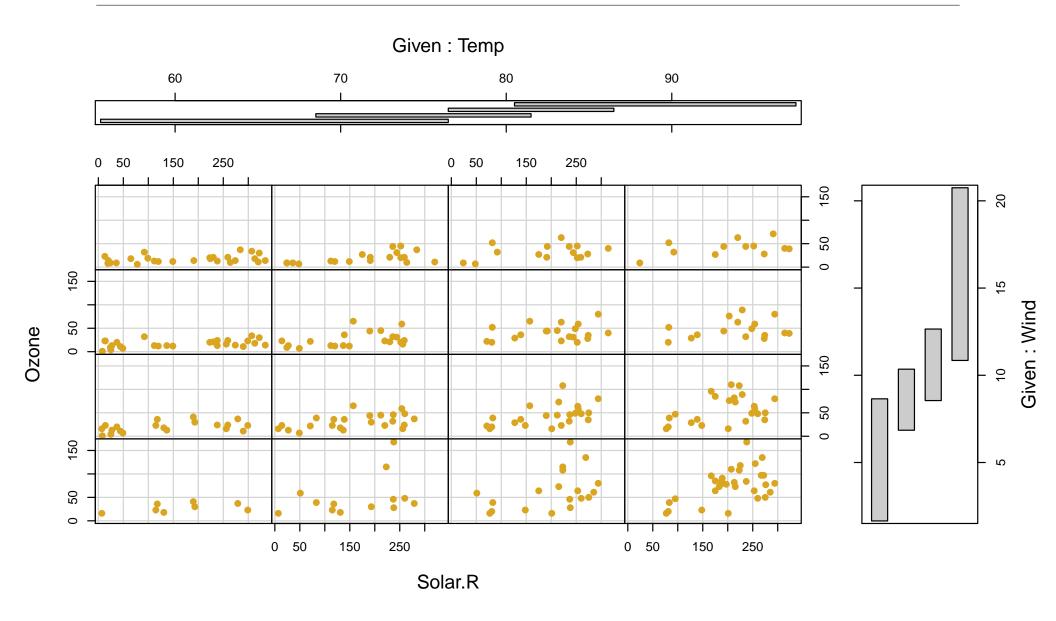
To help us illustrate how scientists think, a bit of science;

Ozone is a secondary pollutant, produced from organic compounds and atmostpheric oxygen, in reactions catalyzed by nitrogen oxides and powered by sunlight. But for ozone concentrations in NY in summer (Y) a smoother (code later) shows a non-monotone relationship with sunlight (X) ...



Now draw a scatterplot of Ozone vs Solar.R for various subranges of Temp and Wind.

- The vertical dash ("|") means 'given particular values of',
 i.e. 'conditional on'
- Here, "+" means 'and', not 'plus' see ?formula, and later sessions
- How does Ozone depend on Solar Radiation, on days with (roughly) the same Temperature and Wind Speed?
- ullet ...using the airquality data, with a 4 \times 4 layout, with solid dark yellow circular symbols



What does this show?

- A 4-D relationship is illustrated; the Ozone/sunlight relationship changes in strength depending on both the Temperature and Wind
- The horizontal/vertical 'shingles' tell you which data appear in which plot. The overlap can be set to zero, if preferred
- coplot()'s default layout is a bit odd; try setting rows, columns to different values
- Almost any form of plot can be 'conditioned' in this way –
 but the commands are in the non-default lattice package

NB it is possible to produce 'fake 3D' plots in R – but (on 2D paper) conditioning plots work better!

Summary

- R makes publication-quality graphics, as well as graphics for data exploration and summary
- plot() is generic, and adapts to what you give it. There are (necessarily) lots of arguments to consider; colors, plotting symbols, labels, etc
- hist(), boxplot(), dotplot() and coplot() offer more functionality
- The formula syntax is a (more) natural way from translate scientific aims to choice of what to plot
- Much more to come! In the next section we'll build up more complex plots