Last week we left off with a discussion on colors and how they can improve a chart or plot.

We found that we could use built-in palettes from which we could pick "premade" colors. This makes it easy for those of us who can't really pick matching color schemes so easily.

This week we extend this idea some and learn about "color ramps", annotation, as well as some other chart types.

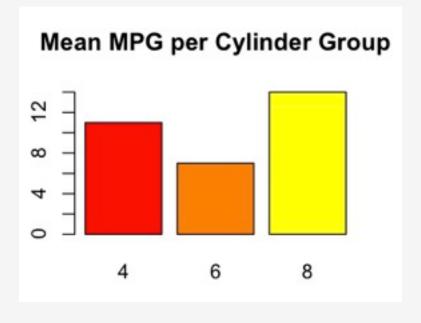
We also explore lattice graphics.

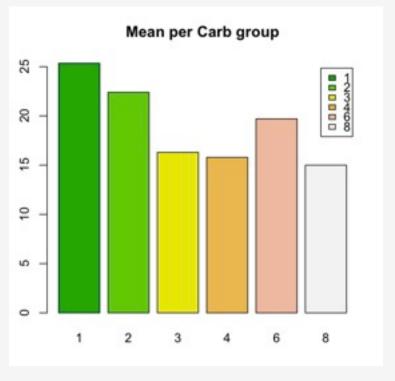
```
myt = tapply(mtcars$mpg, mtcars$cyl, mean)

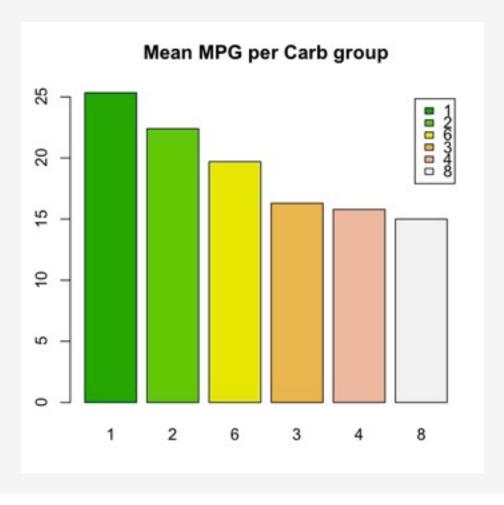
# tapply produces a table

    4     6     8
26.66364 19.74286 15.10000

barplot(myt, main = "Mean MPG per Cylinder Group", col=heat.colors(3))
```







```
barplot(rev(sort(myt)), main = "Mean MPG per Carb group",
             col= terrain.colors(length(myt)),legend=T)
sort(myt)
15.00000 15.79000 16.30000 19.70000 22.40000 25.34286
rev(sort(myt))
25.34286 22.40000 19.70000 16.30000 15.79000 15.00000
# I reverse the sort to make it so the legend doesn't overwrite one of the bars. Try
# plotting it without doing the rev to see what I mean.
barplot(sort(myt), main = "Mean MPG per Carb group",
             col= terrain.colors(length(myt)),legend=T)
```

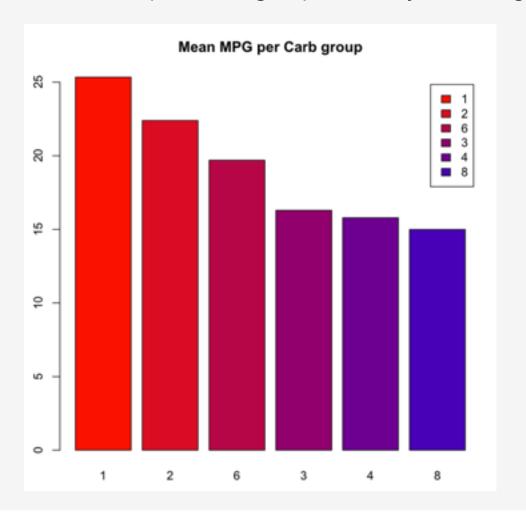
What about making the colors reflect the mean? If we have an increasing sequence of bars why not make the color for each group a different shade from a graduated color scale.

To do this we need a "color ramp" command to generate a palette. Its easier to understand this with an example.

```
mycols = colorRampPalette(c("red","blue"))( 6 )
[1] "#FF0000" "#DA0024" "#B60048" "#91006D" "#6D0091" "#4800B6"
```

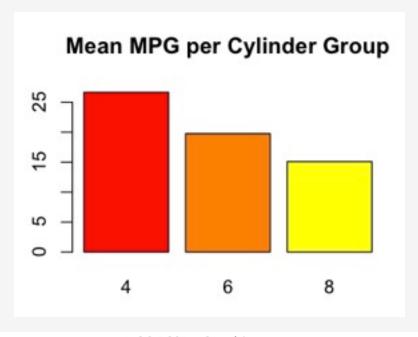
This creates a graduated color scheme between red and blue. Let's apply this to our barplot.

barplot(rev(sort(myt)),
 main = "Mean MPG per Carb group", col=mycols, legend=T)



```
myt = tapply(mtcars$mpg, mtcars$cyl, mean)
# tapply produces a table

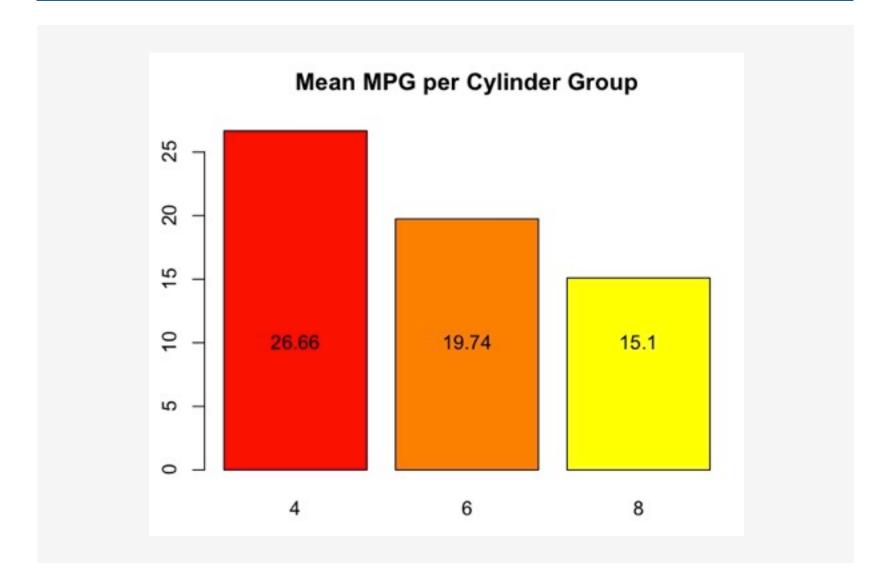
    4     6     8
26.66364 19.74286 15.10000
barplot(rev(sort(myt)), main = "Mean MPG per Cylinder Group", col=heat.colors(3))
```



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```
myt = tapply(mtcars$mpg, mtcars$cyl, mean)
# tapply produces a table
26.66364 19.74286 15.10000
temp = barplot(myt, main = "Mean MPG per Cylinder Group",
                    col=heat.colors(3))
            # These represent the X coordinates for each bar
temp
     \lceil , 1 \rceil
[1,] 0.7
[2,] 1.9
[3,] 3.1
#
     x y text
text(0.7, 10, 26.66) # We put up the means one by one
text(1.9, 10, 19.74)
text(3.1, 10, 15.10)
```



But there is an easier way. The text function arguments can accept vectors. So an easier way to do this:

```
#
              text
     x y
text(0.7, 10, 26.66) # We put up the means one by one
text(1.9, 10, 19.74)
text(3.1, 10, 15.10)
# is:
text(temp, 10, round(myt, 2))
temp
    [,1]
[1,] 0.7
[2,] 1.9
[3,] 3.1
```

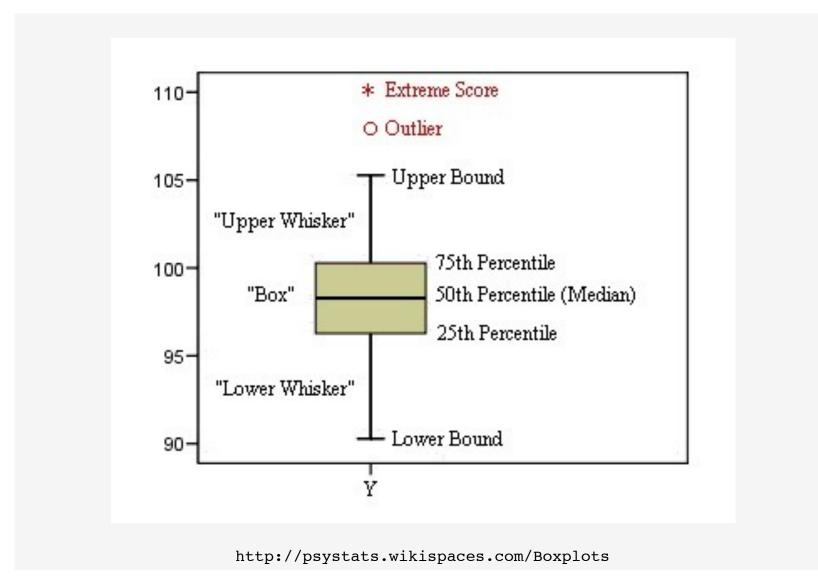
If you are doing a X/Y scatterplot and one of them is a categorical variable then the resulting plot will be a boxplot.

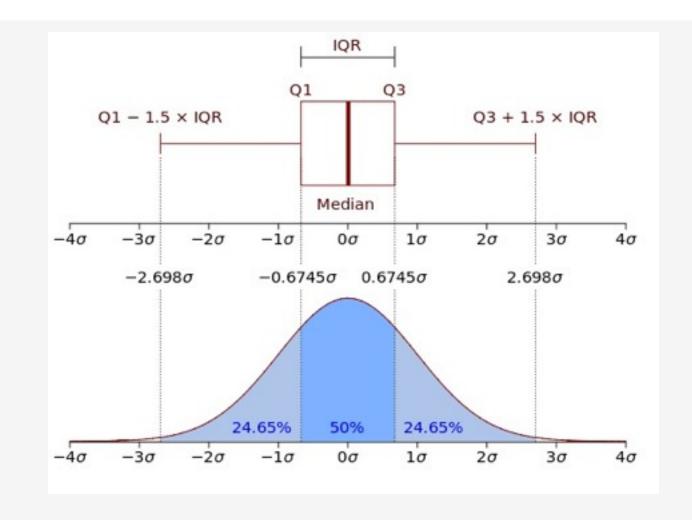
A box plot summarizes a lot of information clearly. It shows the location and spread of data as well as skewness.

The horizontal line shows the median. The bottom and top lines show the 25th and 75th percentiles respectively.

The vertical dashed lines are called "whiskers". The show the maximum of 1) the smaller of the data being plotted or 1.5 times the interquartile range.

1.5 times the interquartile is roughly two standard deviations. And the IQR is the difference in the response variable between the first and third quartile.



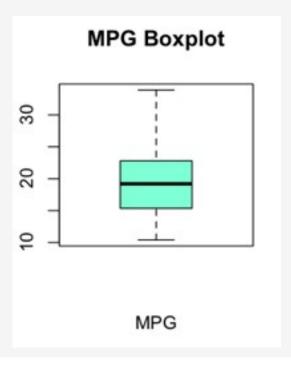


http://en.wikipedia.org/w/index.php?title=File:Boxplot_vs_PDF.svg

Note that any boxplot output will also match the output from the "fivenum" command:

```
fivenum(mtcars$mpg)
[1] 10.40 15.35 19.20 22.80 33.90
```

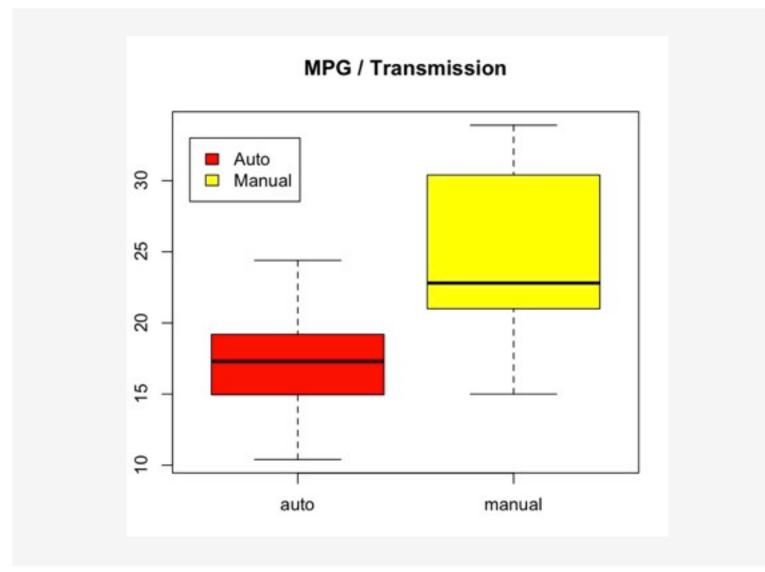
boxplot(mtcars\$mpg, main="MPG Boxplot",xlab="MPG",col="aquamarine")



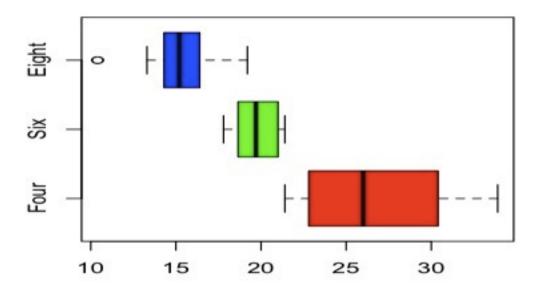
We can get more than one boxplot within a plot window. If you are doing a X/Y scatterplot and one of them is a categorical variable then the resulting plot will be a boxplot. Let's look at the mtcars dataset now.

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MPG by Cylinders



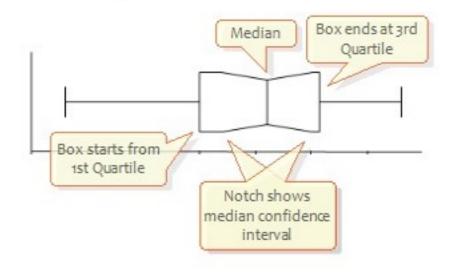
Boxplots are generally good at showing distribution of data around the median but not so good at showing the significance of differences between medians. Tukey introduced the idea of "notched" plots to address this problem.

Boxes for which the notches do not overlap are "likely" to have significantly different medians in terms of testing.

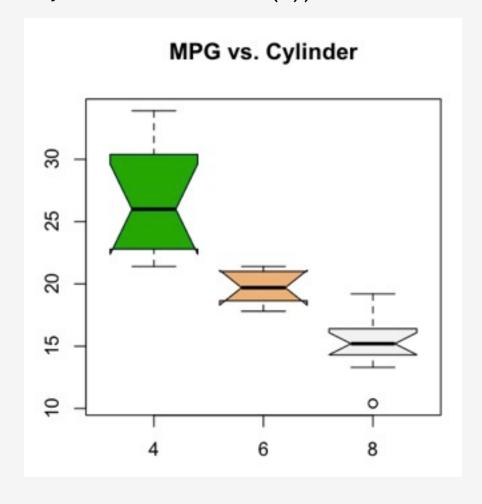
The width of the notches is proportional to the interquartile range of the sample and inversely proportional to the square root of the size of the sample.

http://analyse-it.com/docs/220/standard/summary_paired.htm

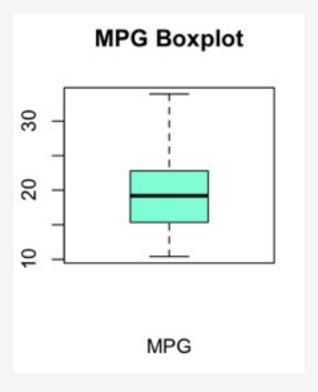
 Notched box plots show a basic box plot as above, with the addition of a notched (pinched or indented) section for the confidence interval around the median (see below).



http://analyse-it.com/docs/220/standard/summary_paired.htm



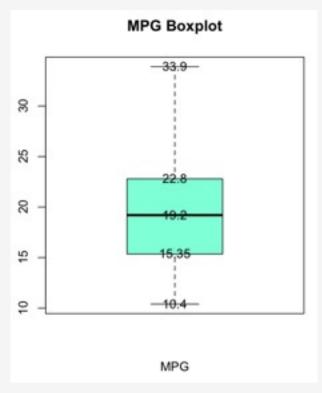
boxplot(mtcars\$mpg, main="MPG Boxplot",xlab="MPG",col="aquamarine")



```
hold = boxplot(mtcars$mpg, main="MPG Boxplot",
               xlab="MPG",col="aquamarine")
$stats
               # This contains the fivenum summary info
      [,1]
[1,] 10.40
[2,] 15.35
[3,] 19.20
[4,] 22.80
[5,] 33.90
$n
               # This contains the number of records in the boxplot
[1] 32
$conf
         [,1]
[1,] 17.11916
[2,] 21.28084
$out
numeric(0)
$group
numeric(0)
$names
[1] ""
```

There is only one boxplot so the X coordinate is "1".

```
text(1,10.4,10.4)
text(1,15.35,15.35)
text(1,19.20,19.20)
text(1,22.8,22.8)
text(1,33.9,33.9)
```



We could do this more simply by recognizing that the text function can handle vectors as well as single values (as in the previous example).

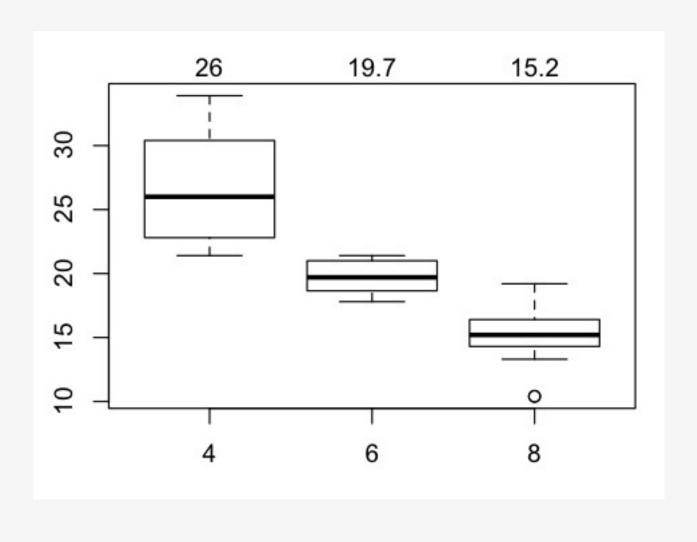
text(1,hold\$stats[,1],hold\$stats[,1])

As with the barchart command, the boxplot command returns some interesting information such as the fivenum value statistics and some other stuff:

```
tmpvar = boxplot(mpg~cyl,data=mtcars)
$stats
     [,1] [,2] [,3]
[1,] 21.4 17.80 13.3
[2,] 22.8 18.65 14.3
[3,] 26.0 19.70 15.2
                         # This row represents the median across the cylinder groups
[4,] 30.4 21.00 16.4
[5,] 33.9 21.40 19.2
$n
                    # This row represents the number of observations in each cyl group
[1] 11 7 14
$conf
         [,1]
                  [,2] [,3]
[1,] 22.37945 18.29662 14.31323
[2,] 29.62055 21.10338 16.08677
$out
[1] 10.4 10.4
$group
[1] 3 3
$names
[1] "4" "6" "8"
```

As with the barchart command, the boxplot command returns some interesting information such as the fivenum value statistics and some other stuff:

```
tmpvar = boxplot(mpg~cyl,data=mtcars)
tmpvar$stats
     [,1] [,2] [,3]
[1,] 21.4 17.80 13.3
[2,] 22.8 18.65 14.3
[3,] 26.0 19.70 15.2
[4,] 30.4 21.00 16.4
[5,] 33.9 21.40 19.2
tmpvar$stats[3,]
[1] 26.0 19.7 15.2
mtext(at=1:3, text=tmpvar$stats[3,])
# same as doing the following:
mtext(at=1, text=tmpvar$stats[3,1], side=3) # top margin
mtext(at=2, text=tmpvar$stats[3,2], side=3) # top margin
mtext(at=3, text=tmpvar$stats[3,3], side=3) # top margin
```

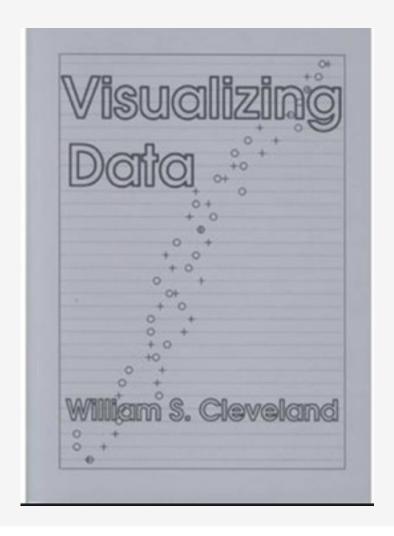


Graphics

LATTICE Graphics

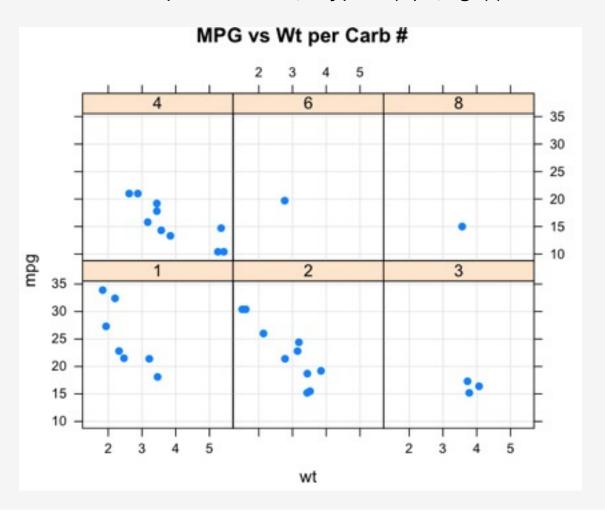
Note that much of the material in this section is attributable to the Lattice intro available at:

http://lattice.r-forge.r-project.org/Vignettes/src/lattice-intro/lattice-intro.pdf



Why "lattice" ?





Design goals

Visualization is an art, but it can benefit greatly from a systematic, scientific approach. In particular, ? has shown that it is possible to come up with general rules that can be applied to design more effective graphs.

One of the primary goals of Trellis graphics is to provide tools that make it easy to apply these rules, so that the burden of compliance is shifted from the user to the software to the extent possible. Some obvious examples of such rules are:

- Use as much of the available space as possible
- Force direct comparsion by superposition (grouping) when possible
- Encourage comparison when juxtaposing (conditioning): use common axes, add common reference objects such as grids.

The following display types are available in lattice.

Function	Default Display Histogram	
histogram()		
densityplot()	Kernel Density Plot	
qqmath()	Theoretical Quantile Plot	
qq()	Two-sample Quantile Plot	
stripplot()	Stripchart (Comparative 1-D Scatter Plots)	
bwplot()	Comparative Box-and-Whisker Plots	
dotplot()	Cleveland Dot Plot	
barchart()	Bar Plot	
xyplot()	Scatter Plot	
splom()	Scatter-Plot Matrix	
contourplot()	Contour Plot of Surfaces	
levelplot()	False Color Level Plot of Surfaces	
wireframe()	Three-dimensional Perspective Plot of Surface	
cloud()	Three-dimensional Scatter Plot	
parallel()	Parallel Coordinates Plot	

Lattice Function	Description	Traditional Analogue
barchart()	Barcharts	barplot()
bwplot()	Boxplots Box-and-whisker plots	boxplot()
densityplot()	Conditional kernel density plots Smoothed density estimate	none
dotplot()	Dotplots Continuous versus categorical	dotchart()
histogram()	Histograms	hist()
qqmath()	Quantile-quantile plots Data set versus theoretical distribution	qqnorm()
stripplot()	Stripplots One-dimensional scatterplot	stripchart()
qq()	Quantile-quantile plots Data set versus data set	qqplot()
xyplot()	Scatterplots	plot()
levelplot()	Level plots	image()
contourplot()	Contour plots	contour()
cloud()	3-dimensional scatterplot	none
wireframe()	3-dimensional surfaces	persp()
splom()	Scatterplot matrices	pairs()
parallel()	Parallel coordinate plots	none

Even though lattice graphics is part of R you have to first load the lattice library before using any of the commands:

library(lattice)

Trellis displays are defined by the type of graphic and the role different variables play in it. Each display type is associated with a corresponding high-level function (histogram, densityplot, etc.). Possible roles depend on the type of display, but typical ones are:

primary variables: those that define the primary display (e.g., usually some continuous variables)

conditioning variables: divides data into subgroups, each of which are presented in a different panel (e.g., a category or factor).

grouping variables: subgroups are contrasted within panels by superposing the corresponding displays (e.g., gender in the last example).

http://lattice.r-forge.r-project.org/Vignettes/src/lattice-intro/lattice-intro.pdf

Factor review:

```
Factors

Description:

The function 'factor' is used to encode a vector as a factor (the terms 'category' and 'enumerated type' are also used for factors).

If argument 'ordered' is 'TRUE', the factor levels are assumed to be ordered. For compatibility with S there is also a function 'ordered'.

'is.factor', 'is.ordered', 'as.factor' and 'as.ordered' are the membership and coercion functions for these classes.

Usage:

factor(x = character(), levels, labels = levels, exclude = NA, ordered = is.ordered(x), nmax = NA)
```

Factor review:

```
mtcars$am = factor(mtcars$am, labels=c("auto", "manual"))
mtcars$am
 [1] manual manual manual auto
                                                     auto
                                auto
                                       auto
                                              auto
                                                            auto
                                                                   auto
                                       auto
[11] auto
           auto
                  auto
                                              auto
                                                     manual manual manual
                         auto
                                auto
                                       manual manual manual manual
[21] auto
           auto
                  auto
                         auto
                                auto
[31] manual manual
Levels: auto manual
levels(mtcars$am)
[1] "auto"
            "manual"
```

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Lattice/Trellis graphics display a variable, or a relationship between variables, conditioned on one or more other variables. The typical format is:

graph_type(formula,data=some.data) where formula specifies the variable(s) to display and any conditioning variables.

 $\sim x/A$ means display numeric variable x for each level of factor A on separate panels.

bwplot(~mpg | factor(am), data=mtcars)

x ~ A means display numeric variable x for each level of A on One panel.
bwplot(mpg ~ factor(am), data=mtcars)

~x means display numeric variable x alone. hist(~mpg, data=mtcars)

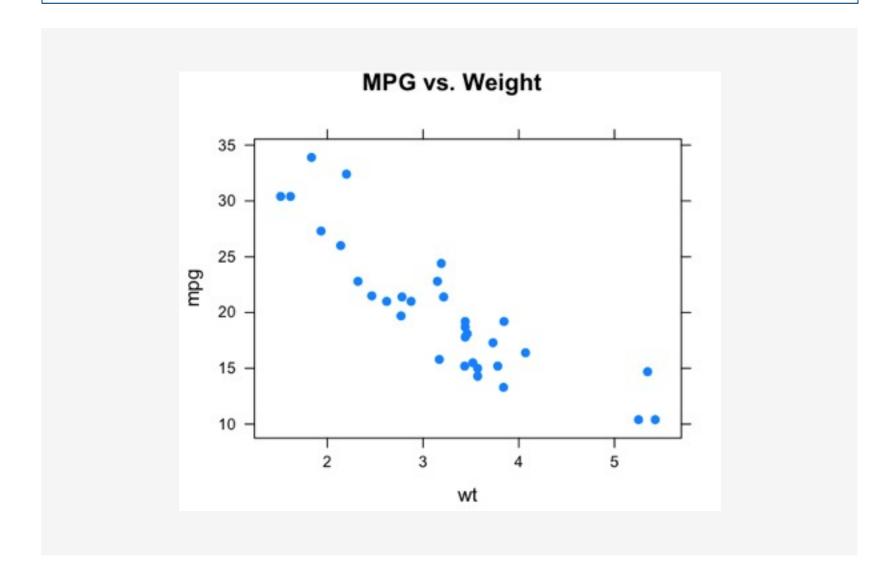
graph_type	description	formula examples
barchart	bar chart	x~A or A~x
bwplot	boxplot	x~A or A~x
cloud	3D scatterplot	z~x*y A
contourplot	3D contour plot	z~x*y
densityplot	kernal density plot	~x A*B
dotplot	dotplot	~x A
histogram	histogram	~x
levelplot	3D level plot	z~y*x
parallel	parallel coordinates plot	dataframe
splom	scatterplot matrix	dataframe
stripplot	strip plots	A~x or x~A
xyplot	scatterplot	y~x A

The typical Basic form when calling a lattice function such as xyplot looks like:

```
vertical.axis.variable ~ horizontal.axis.variable
```

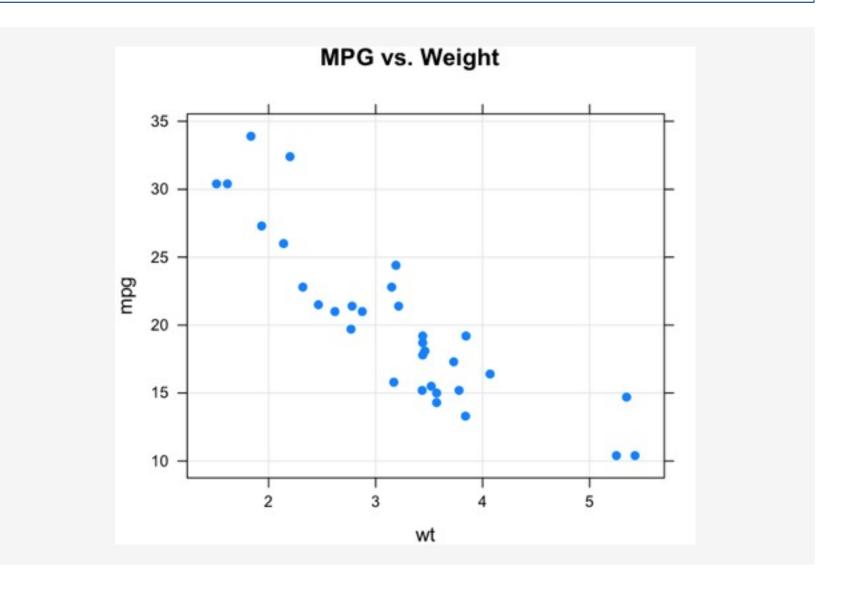
```
xyplot(mpg ~ wt, data = mtcars, pch = 19, main = "MPG vs. Weight")
```

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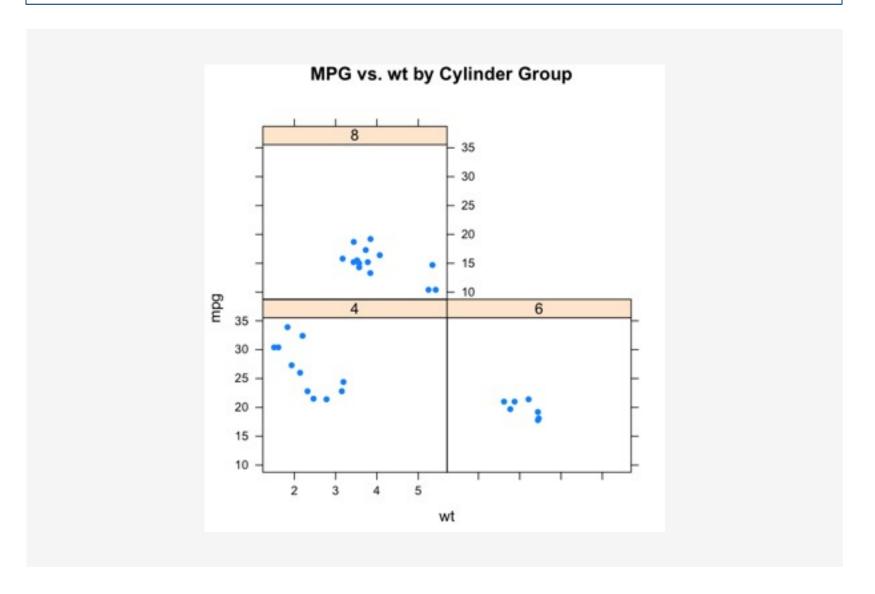
* Check out the "type" argument that let's us specify the type of graph we want (e.g. points, lines, grid)

```
xyplot(mpg ~ wt, data = mtcars, pch = 19,
    main = "MPG vs. Weight", type=c("p","g"))
```



* So check out this formula. "mpg" is the y variable and "wt" is the x variable.

- * "cylinder" is a conditioning variable (preceded by the | character). The conditioning variable divides the plot into separate panels
- * Conditioning variables are usually categorical.
- * Note that the "data" argument tells the function what data frame to use.

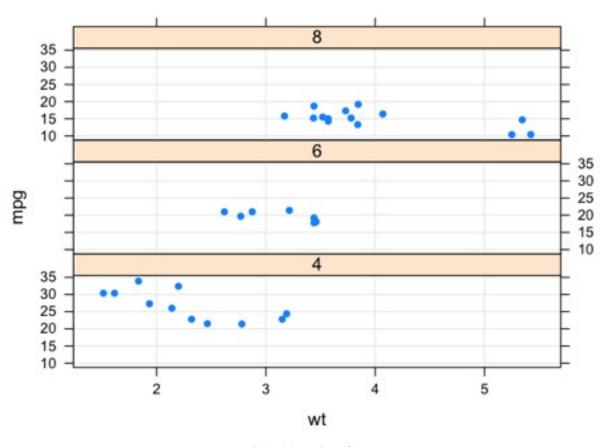


* So check out this formula. "mpg" is the y variable and "wt" is the x variable.

- * "cylinder" is a conditioning variable (preceded by the | character). The conditioning variable divides the plot into separate panels
- * Conditioning variables are usually categorical.
- * Note that the "data" argument tells the function what data frame to use.

In the Trellis terminology, this plot consists of three panels. Each panel in this case contains a scatterplot and above each panel there is a strip that presents the level of the conditioning variable.

MPG vs. wt by Cylinder Group



Remember our early X/Y plot with Base graphics where we wanted to plot MPG vs Weight for each cylinder category? To get a side-by-side panel plot we did something like this:

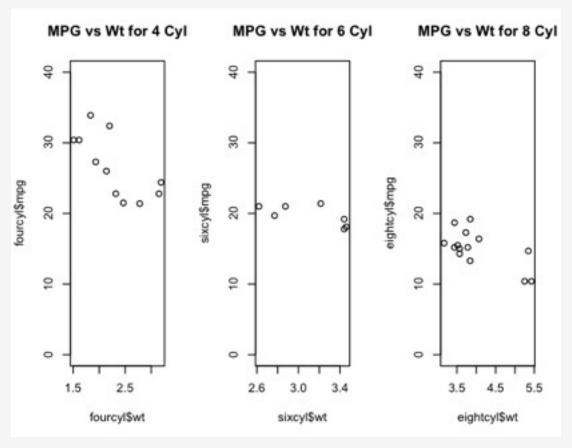
```
par(mfrow=c(1,3))

fourcyl = mtcars[mtcars$cyl == 4,]
sixcyl = mtcars[mtcars$cyl == 6,]
eightcyl = mtcars[mtcars$cyl == 8,]

plot(fourcyl$wt, fourcyl$mpg, main = "MPG vs Wt for 4 Cyl", ylim=c(0,40))
plot(sixcyl$wt, sixcyl$mpg, main = "MPG vs Wt for 6 Cyl", ylim=c(0,40))
plot(eightcyl$wt, eightcyl$mpg, main = "MPG vs Wt for 8 Cyl",
ylim=c(0,40))
```

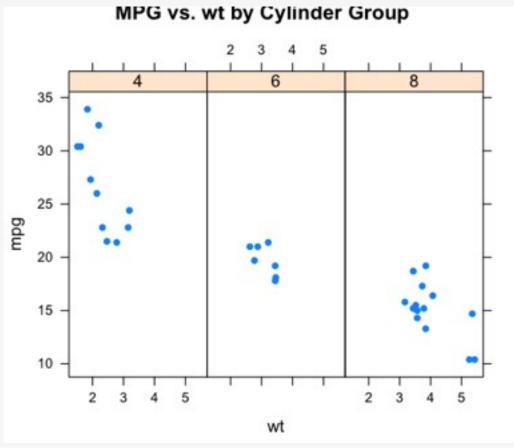
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Remember our early X/Y plot with Base graphics where we wanted to plot MPG vs Weight for each cylinder category? To get a side-by-side panel plot we did something like this:



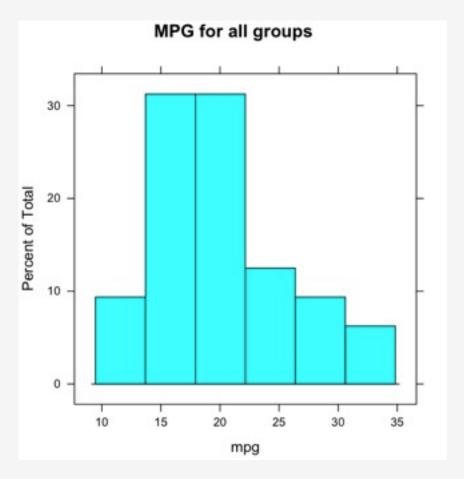
With lattice plots its much easier:

library(lattice)
xyplot(mpg~wt|factor(cyl),data = mtcars,layout=c(3,1))



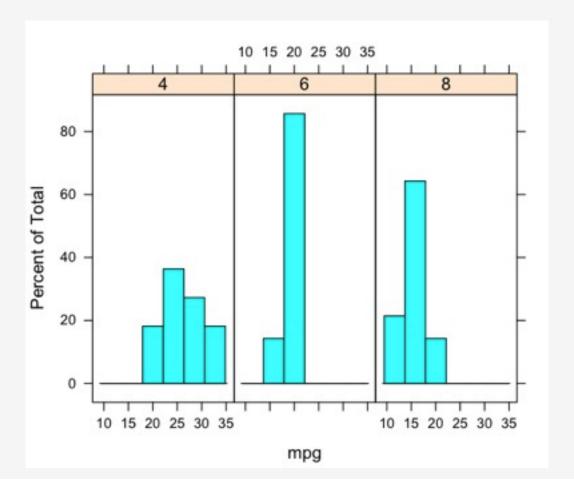
This works independently of the chart type:

histogram(~mpg, data = mtcars, main = "MPG for all groups")



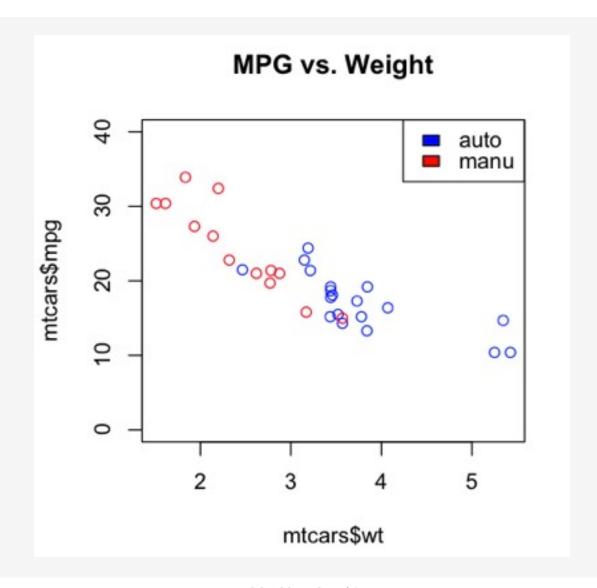
This works independently of the chart type:

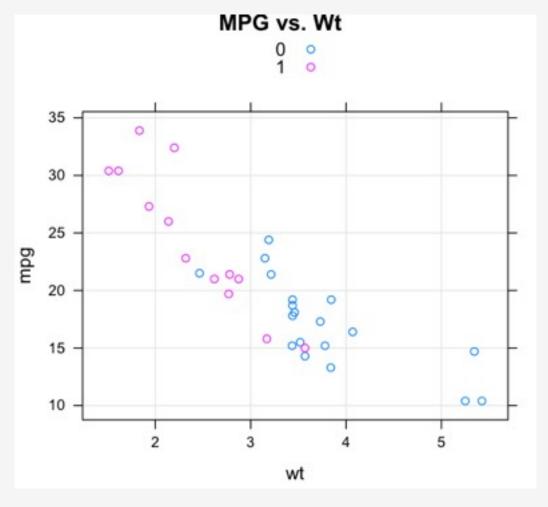
histogram(~mpg | factor(cyl), data = mtcars, layout = c(3,1))



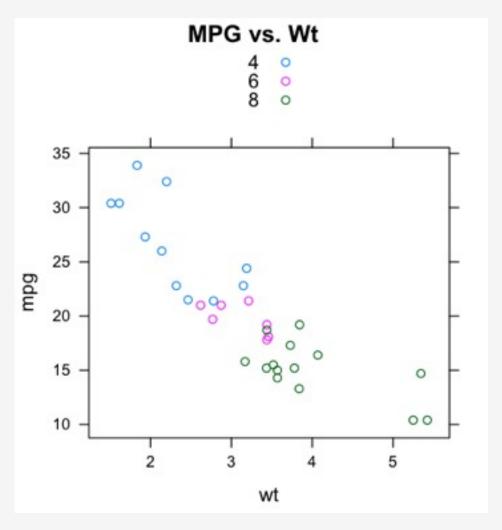
Remember our early X/Y plot with Base graphics where we wanted to plot MPG vs Weight for each transmission type and have it represented by a different color or plot character? The most basic solution looked like this (although we did improve it somewhat).

```
plot(mtcars$wt, mtcars$mpg, main="MPG vs. Weight", type="n",
     ylim=c(0,40))
auto = mtcars[mtcars$am == 0,]
manu = mtcars[mtcars$am == 1,]
points(auto$wt, auto$mpg, col="blue")
points(manu$wt, manu$mpg, col="red")
legend("topright",c("auto","manu"),fill=c("blue","red"))
```





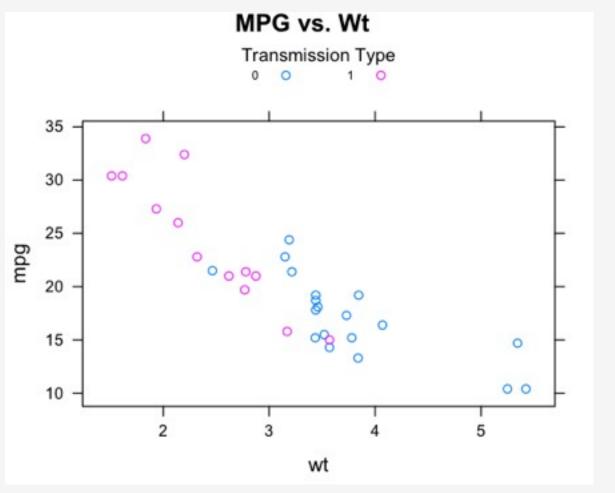
xyplot(mpg~wt, data=mtcars, groups=cyl, main="MPG vs. Wt", auto.key=TRUE)



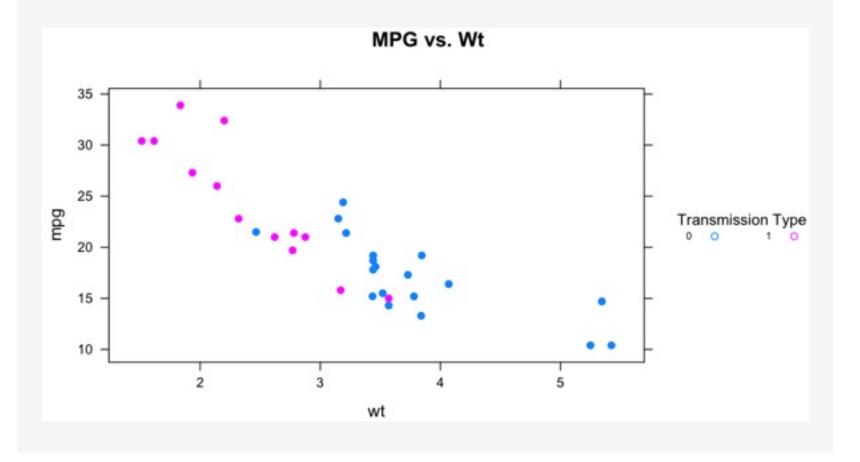
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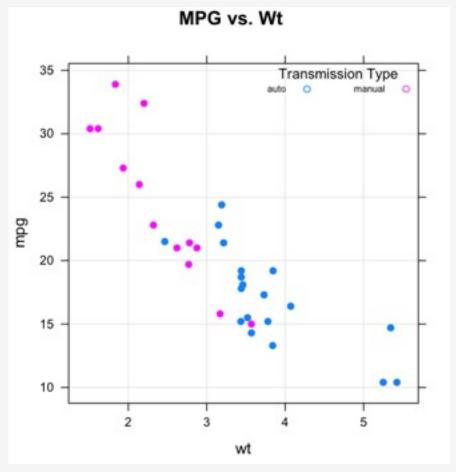
Graphics: lattice: legends



Graphics: lattice: legends

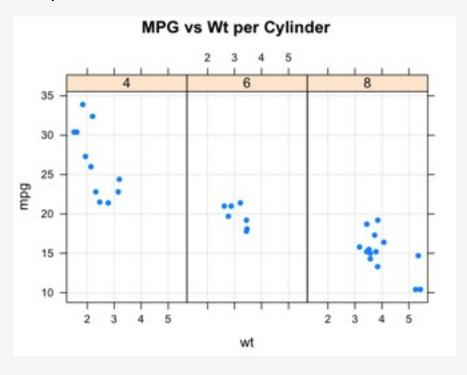


Graphics: lattice: legends



Graphics: lattice: grouping and conditions

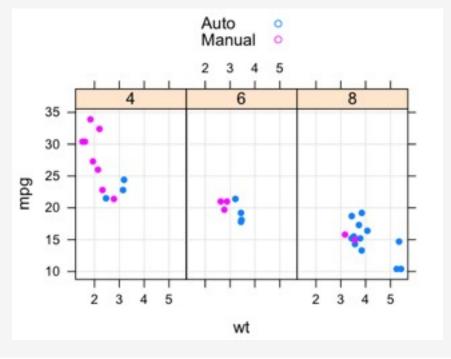
In this case we have a conditioning variable, which in this example is the variable cyl . So we have three panels:

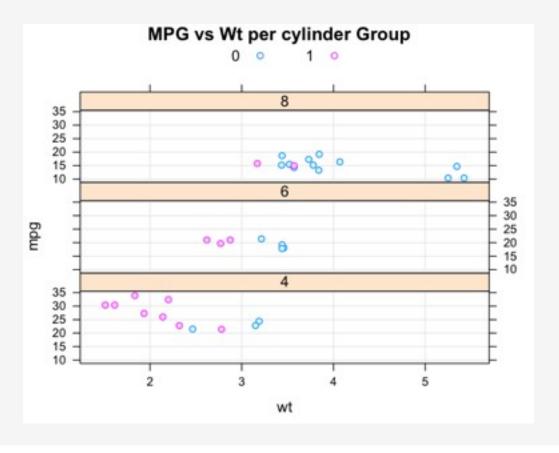


Graphics: lattice: grouping and conditions

```
xyplot(mpg ~ wt | factor(cyl), data=mtcars,
    pch=16,layout=c(3,1),
    groups = factor(am,labels=c("Auto","Manual")),
    type=c("p","g"),auto.key = TRUE)
```

In this case we have a grouping variable, which in this example is the variable am . The grouping variable segregates data into subgroups within each panel via color.





Graphics

Some useful arguments that you will see again and again:

data = (this specifies the data frame of interest)

xlab = and ylab = (this indicates the X/Y axis labels

aspect = (sets the aspect ration of the plot – not so common but useful)

xlim = and ylim = (sets the numerical limits of the X and Y vals)

main = (sets the title for the plot)

cex = (changes the size of an element like text or the plotting symbol

http://polisci.msu.edu/jacoby/icpsr/graphics/lattice/Lattice,%20ICPSR%202012%20Outline,%20Ver %201.pdf

Graphics

Some useful arguments that you will see again and again:

The **auto.key** = **T** argument gives you an automatic legend.

The **scales** = argument uses a "mini-language" to modify the characteristics of the display axes such as ticks, and tick labels. Its basically a "list" (in the R sense)

The **group** = argument lets you pass a categorical variable that can be used to differentiate subgroups in a single display.

http://polisci.msu.edu/jacoby/icpsr/graphics/lattice/Lattice,%20ICPSR%202012%20Outline,%20Ver %201.pdf

Summary

Strengths of Base Graphics:

Oldest and Most Well Known

Has Both High Level and Low Level Functions

Can easily assemble a plot in stages

Primitive functions (e.g. points, abline, lines) are simple to use

Weaknesses of Base Graphics:

Lots of independent functions to keep up with

Each high level plot command has its own specific arguments

You have to pick colors and plot characters using basic programming tricks

Summary

Strengths of Lattice Graphics

Developed according to a well-thought out philosophy

Can generate colors, legends, and axes as part of the high level commands

Can use "condition" variables

Can group variables

Weaknesses of Lattice Graphics:

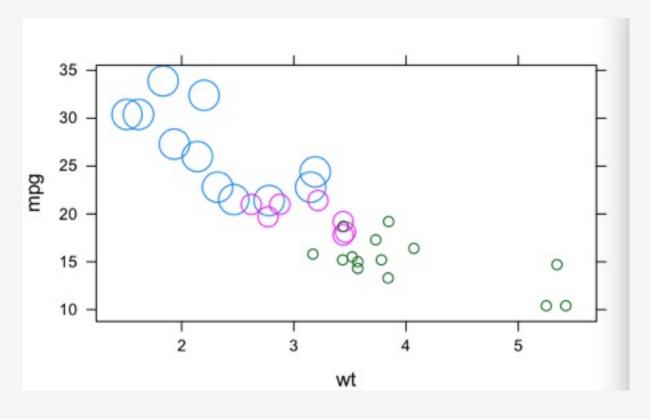
Commands can get long and involved

Commands can have many arguments

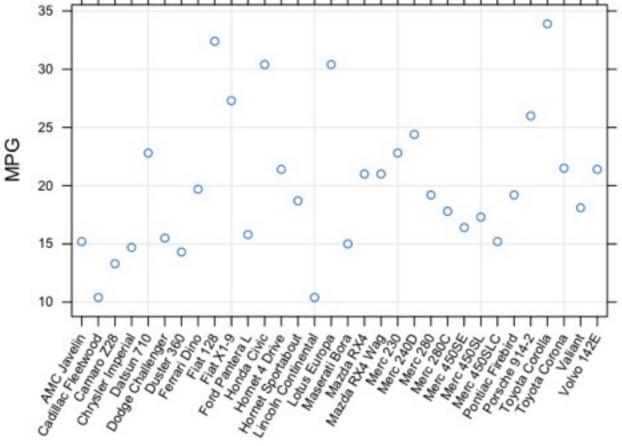
THE FOLLOWING SLIDES ARE SUPPLEMENTAL THOUGH ARE VERY USEFUL

Remember our early X/Y plot with Base graphics where we wanted to plot MPG vs Weight and have the point sizes to reflect the cylinder group (4,6, or 8)? It was quite a bit of work to do with Base graphics. Here we can do it with one line:

xyplot(mpg ~ wt, groups = factor(cyl), cex=3:1, auto.key=T)

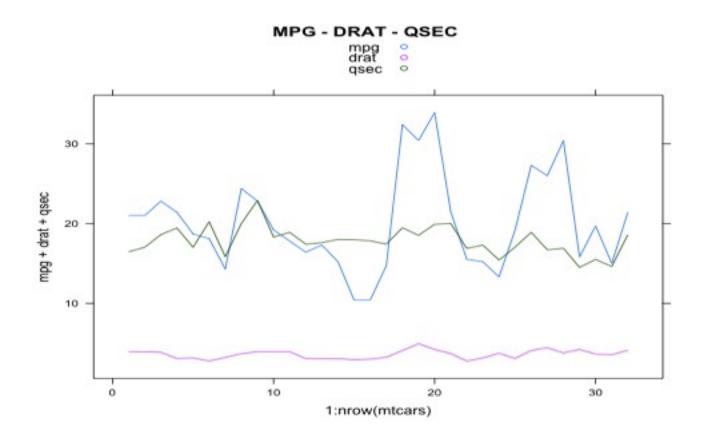


```
xyplot(mpg ~ factor(rownames(mtcars)),
    mtcars, xlab = "Car Names", ylab = "MPG",
    scales = list(cex = 0.7,x = list(rot = 60)),type=c("p","g"))
```



Car Names BIOS 560R - Graphics

```
xyplot(mpg + drat + qsec ~ 1:nrow(mtcars),
    data= mtcars, type="l",auto.key=T,
    main = "MPG - DRAT - QSEC")
```



Every lattice function has a default "panel" function. It is usually the name of the function prefixed by **panel**. As an example here is the basic xyplot command:

```
xyplot (y \sim x, data= dataset)
```

This is actually equivalent to:

```
xyplot (y ~ x, data= dataset, panel = panel.xyplot)
```

Yea, its a bit weird but if you have to stack up a bunch of options and arguments it becomes easier. You'll see eventually.

So a call like this:

```
xyplot(y ~ x, data=dataset, col="black")
```

will "silently" pass the "col='black' argument to panel.xyplot.

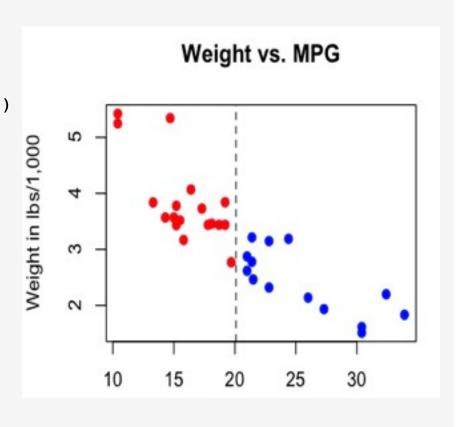
http://polisci.msu.edu/jacoby/icpsr/graphics/lattice/Lattice,%20ICPSR%202012%20Outline,%20Ver %201.pdf

We can look at group interaction within the panels

```
So a call like this:
xyplot(y ~ x, data=dataset, col="black")
will "silently" pass the "col='black' argument to panel.xyplot. Its basically the
equivalent of:
xyplot(y ~ x, data = dataset,
   panel = function (x, y) {
        panel.xyplot(x, y, col = "black") }
http://polisci.msu.edu/jacoby/icpsr/graphics/lattice/Lattice,%20ICPSR%202012%20Outline,%20Ver
%201.pdf
```

So remember that we did this in BASE graphics to get a plot with points less than the mean MPG to be one color and the points greater than the mean to be another.

```
attach(mtcars)
plot(mpg,wt,type="n",
      main="Weight vs. MPG",
      ylab="Weight in lbs/1,000")
points(mpg[mpg > mean(mpg)],
         wt[mpg > mean(mpg)],
         col="blue",bq="blue",
         pch=21)
points(mpg[mpg <= mean(mpg)],</pre>
         wt[mpg <= mean(mpg)],</pre>
         col="red",bq="red",
         pch=21)
abline(v=mean(mpg), lty=2)
```

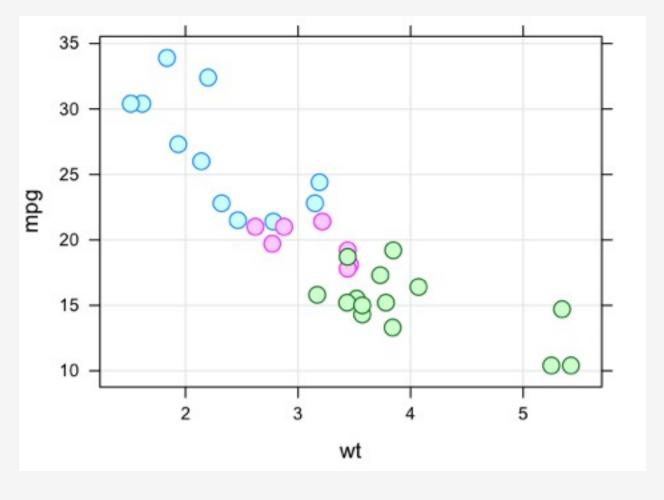


So the two commands are basically the same:

```
xyplot(mpg~wt, data=mtcars, groups=factor(cyl),
    panel = function(x,y,...) {
        panel.xyplot(x,y, cex=1.5, pch=21,...)
        panel.grid()
    }
)

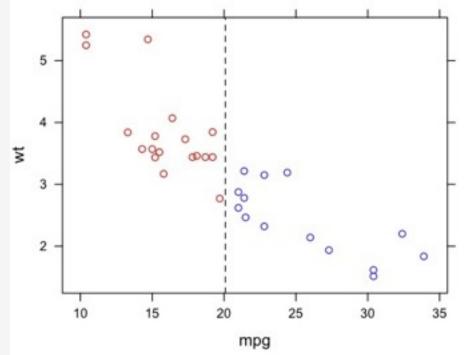
xyplot(mpg~wt, data=mtcars, groups=factor(cyl),
        cex=1.5, pch=21, type=c("p","g"))
```

So the two commands are basically the same:



Here is one way to do it with lattice graphics:

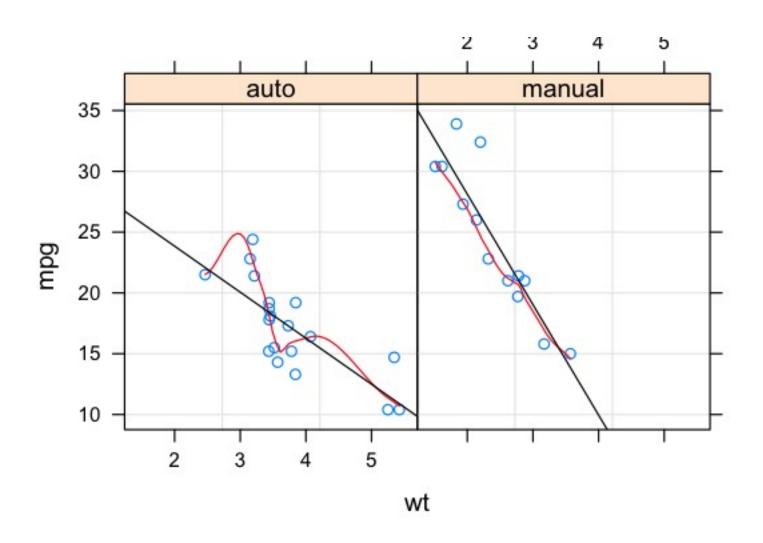
```
xyplot(mpg ~ wt, mtcars,
   panel = function(x,y) {
     panel.xyplot(x[x <= mean(x)], y[x <= mean(x)], col="red")
     panel.xyplot(x[x > mean(x)], y[x > mean(x)], col="blue")
     panel.abline(v=mean(x),lty="dashed")
   }
)
```

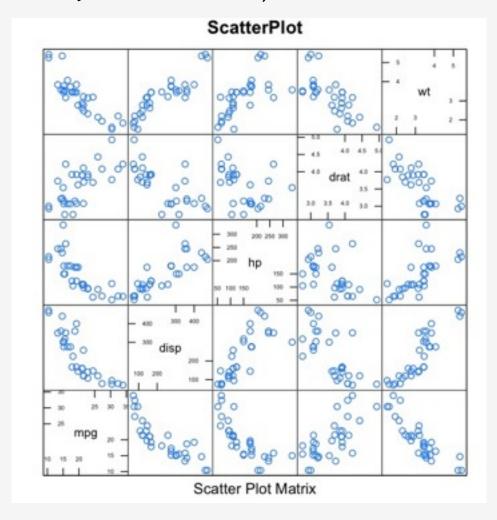


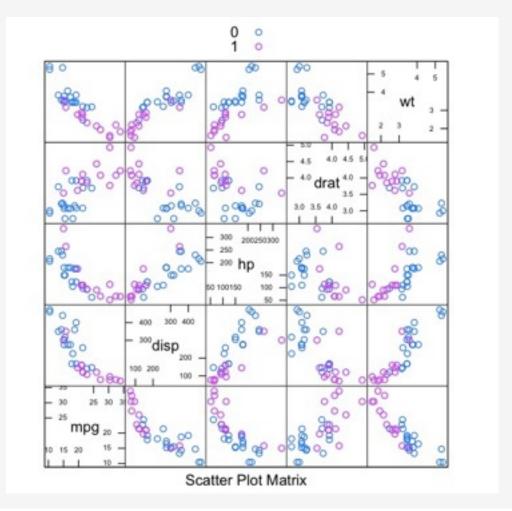
Or alternatively:

```
my.mean = ifelse(mtcars$mpg > mean(mtcars$mpg), "B", "R")
xyplot(wt ~ mpg, mtcars, groups = my.mean, col=c("blue","red"),
    panel = function(...) {
       panel.abline(v=mean(mtcars$mpg), lty="dashed")
       panel.xyplot(...)
                                                 0
                                       5
                                     ₹
                                       2
                                           10
                                                 15
                                                        20
                                                               25
                                                                      30
                                                          mpg
```

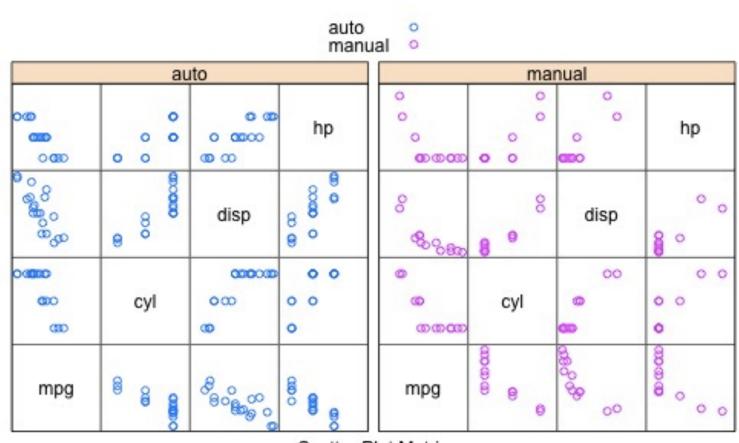
The default panel function for the xyplot() function is panel.xyplot(). This is a "busy" example but it drives home the point that you can do a lot of customization within the panel function. You don't always HAVE to but its a good option to have.



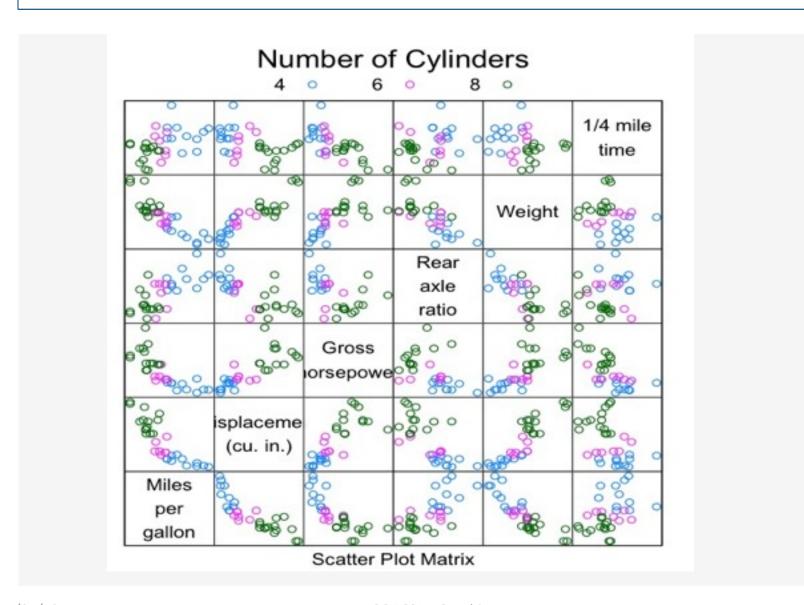




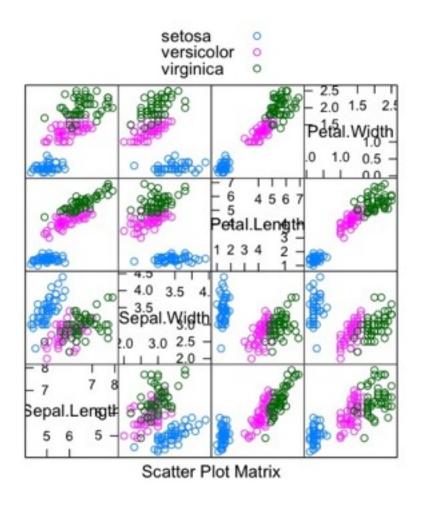
> splom(~mtcars[c(1:4)] | am,groups=am,mtcars,auto.key=TRUE,pscales=0)



Scatter Plot Matrix



splom(~iris[1:4], groups = Species, data = iris,auto.key=T)



Graphics: lattice

plot(x,y) where x and y are continuous:

X/Y, Scatterplot, Sunflower Plot

plot(x,[y]) where x and y are categorical. Note that y can be optional:

dotplot, barplot, stacked bar plot, pie chart

plot(x) where x is a single continuous variable:

dotplot, barplot, stripchart, boxplot, density, histogram, QQ Plot

plot(x,y) where one of x and y is continuous and the other is discrete

Side-by-Side DotPlot and BoxPlot, Notched BoxPlot

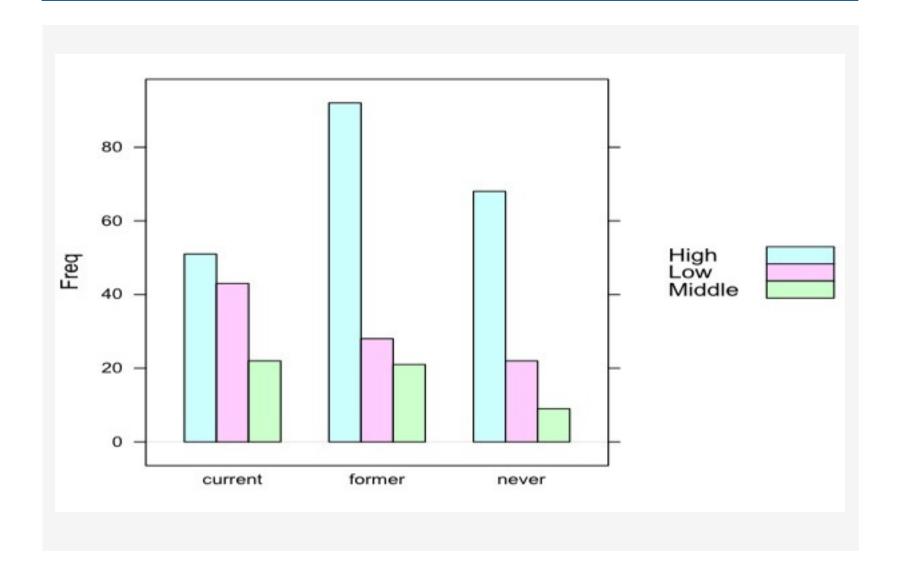
Graphics

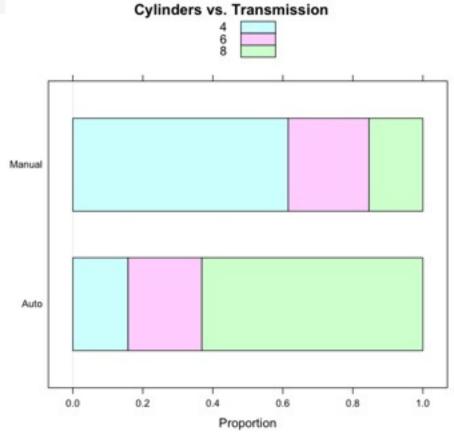
Let's look at a different data set that is smaller. For small samples, summarizing is often unnecessary, and simply plotting all the data reveals interesting features of the distribution.

```
> head(quakes)
    lat long depth mag stations
1 -20.42 181.62
                562 4.8
                             41
2 -20.62 181.03 650 4.2
                             15
3 -26.00 184.10 42 5.4
                            43
4 -17.97 181.66 626 4.1
                             19
5 -20.42 181.96 649 4.0
                             11
6 -19.68 184.31 195 4.0
                             12
> sapply(quakes,class)
             long
     lat
                   depth
                                 maq
                                      stations
"numeric" "numeric" "integer" "numeric" "integer"
> sapply(quakes,range)
       lat long depth mag stations
[1,] -38.59 165.67 40 4.0
                                10
[2,] -10.72 188.13 680 6.4
                               132
```

We surveyed people to see if they smoked ever, heavily, or previously. We were then given some idea of their socioeconomic status ("high","middle","low"). Let's check it out:

```
smokers = read.table("http://www.cyclismo.org/tutorial/R/
smoker.csv",header=T,sep=",")
head(smokers)
     Smoke SES
1 former High
2 former High
( my.table = table(smokers$Smoke,smokers$SES) )
          High Low Middle
            51 43
                       22
  current
  former
            92 28
                       21
            68 22
  never
> barchart(my.table,auto.key=list(space="right"),stack=F,horizontal=F)
```





Graphics: lattice

plot(x,y) where x and y are continuous:

X/Y, Scatterplot, Sunflower Plot

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dotplot, barplot, stacked bar plot, pie chart

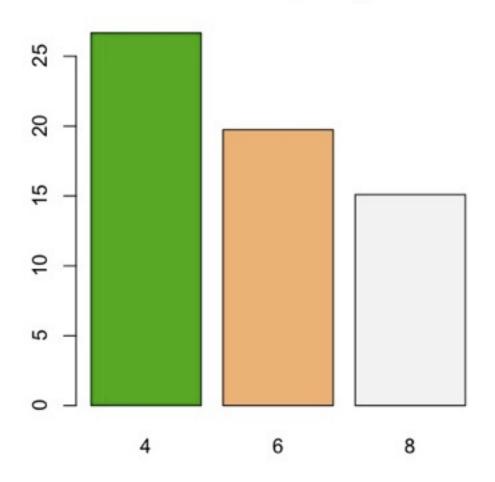
plot(x) where x is a single continuous variable:

dotplot, barplot, stripchart, boxplot, density, histogram, QQ Plot

plot(x,y) where one of x and y is continuous and the other is discrete

Side-by-Side DotPlot and BoxPlot, Notched BoxPlot

Means of MPG per Cylinder



A well-known graphical design that allows comparison between an arbitrary number of samples is the comparative box-and-whisker plot. They are related to the Q-Q plot: the values compared are five "special" quantiles, the median, the first and third quartiles, and the extremes.

More commonly, the extents of the "whiskers" are defined differently, and values outside plotted explicitly, so that heavier-than-normal tails tend to produce many points outside the extremes.

```
> bwplot(factor(score) ~ gcsescore | gender, Chem97,main="A-
level score vs. GCSES score \n by Gender")
```

The decreasing lengths of the boxes and whiskers suggest decreasing variance, and the large number of outliers on one side indicate heavier left tails (characteristic of a left-skewed distribution).

Next. Let's look at some other examples. In We'll be using the mtcars data set as well as the Chem97 dataset contained in the mlmRev package. If you don't have that package installed please install it using your GUI or at the command line by doing:

> install.packages("mlmRev")

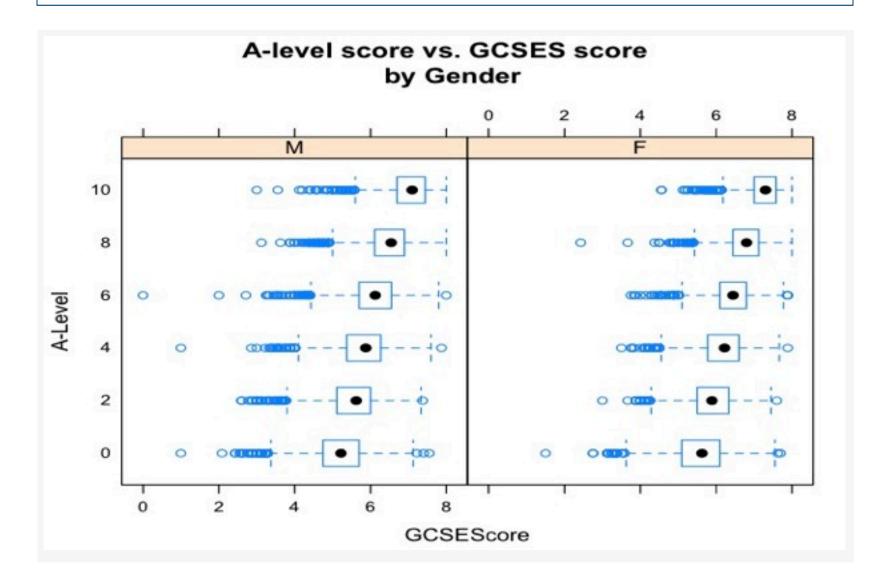
```
> data(Chem97, package="mlmRev")
> head(Chem97)
 lea school student score gender age gcsescore
                                        gcsecnt
1
   1
                          F 3
                                  6.625 0.3393157
                          F -3
2
   1
                   10
                                  7.625 1.3393157
3
   1 1
                   10 F -4
                                  7.250 0.9643157
                   10 F -2
   1
                                  7.500 1.2143157
5
   1
               5 8
                       F -1 6.444 0.1583157
                   10
                         F 4
                                  7.750 1.4643157
```

This dataset contains information in the 1997 A-level chemistry examination in Britain. Let's focus on the the following variables:

^{*} score: point score in the exam, with six possible values (0,2,4,6,8) (discrete)

^{*} gcsescore: average score in GCSE exams, This is a continuous score that can be used to predict the A-level scores. (continuous)

^{*} gender: Gender of the student (discrete/binary)

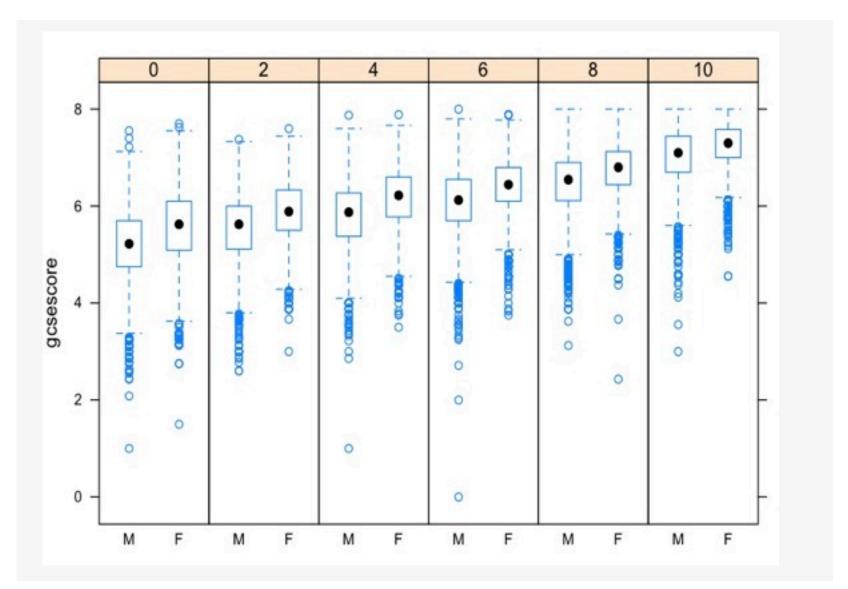


Note that we change the plot to get a different view. We do this by switching the order of the variables. The previous example looks like:

```
> bwplot(factor(score) ~ gcsescore | gender, Chem97,main="A-
level score vs. GCSES score \n by Gender")
Let's do this:
```

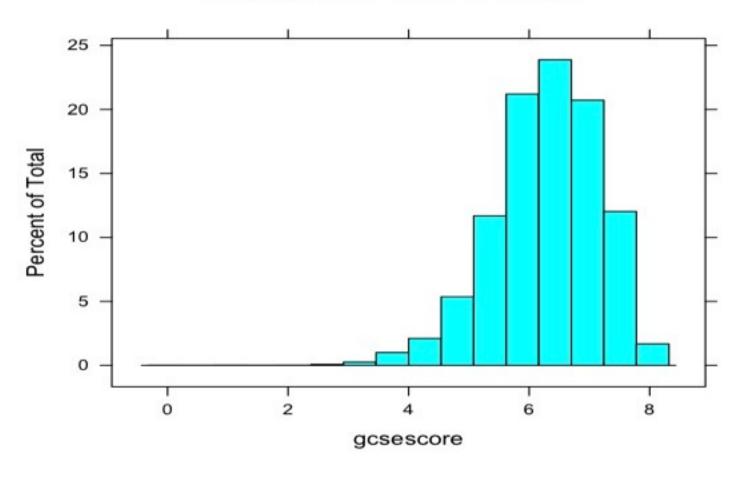
> bwplot(gcsescore ~ gender | factor(score), Chem97,
layout=c(6,1))

We get a different view of this information:



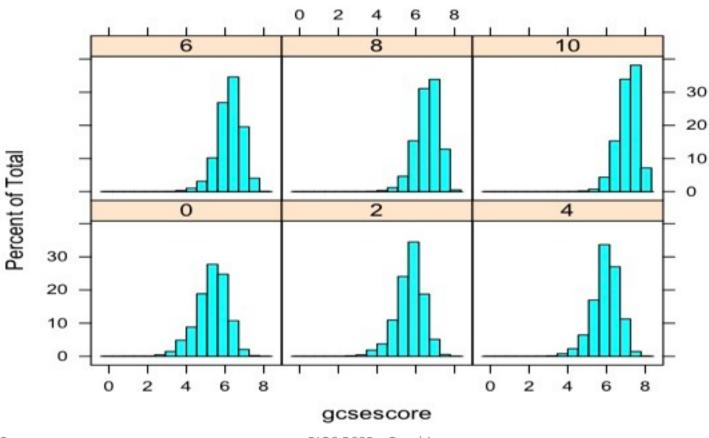
```
> histogram(~ gcsescore,data=Chem97,
+ main="Histogram of GCSES score")
```

Histogram of GCSES score



Okay, that's cool but it is kind of boring and uninformative. It might be better to look at a histogram across the scores, which could be considered as a factor grouping.

> histogram(~ gcsescore | factor(score), data = Chem97)



11/04/12

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