

Seminar III: R/Bioconductor

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Advanced Plotting

intro

lattice

plotrix

ggplot2

car

R is strong in plots

- ▶ As you might recall, R is very strong for making **plots**, and it does so fast.
- ▶ We've seen how to make barplots, qqplots, mosaicplots, and many other ones.
- ▶ After all, plotting is very important for doing **exploratory data analysis**.
- ▶ However, all of them just make a small part.

Install some packages

To gain some time, please install these packages:

```
> install.packages("lattice")  
> install.packages("mlmRev")  
> install.packages("plotrix")  
> install.packages("ggplot2")  
> install.packages("car")  
> install.packages("DAAG")
```

Task Views

- ▶ First of all, remember the CRAN **Task Views**.
- ▶ <http://cran.r-project.org/web/views/Graphics.html>
- ▶ From there, go to the plotrix page.
- ▶ What two functions did they introduce on version 2.5-3?

tools

- ▶ You might decide to check the reference manual and test out the examples, but that's quite time consuming.
- ▶ I found out on the [R Journal](#) about a new function on the tools package.

```
> library(tools)  
> testInstalledPackage(pkgname)
```
- ▶ Its very **easy** to create pdf files with all the example plots of a given package!

Remember to check the help

- ▶ Remember to use:
 - > `help.start()`
 - > `help(package = pkgname)`
- ▶ What is the replacement of the **hist** function on the lattice package?

Intro

- ▶ It's an *implementation* of Trellis graphics and created by Deepayan Sarkar.
- ▶ <http://dsarkar.fhcrc.org/>
- ▶ Basically, its great for plotting multivariate data!
> `?` (*Lattice*)
- ▶ How are the lattice high level functions special?

Data

- ▶ We'll use a data set from the `mlmRev` package and in general we'll follow the BioC2008 lattice lab.

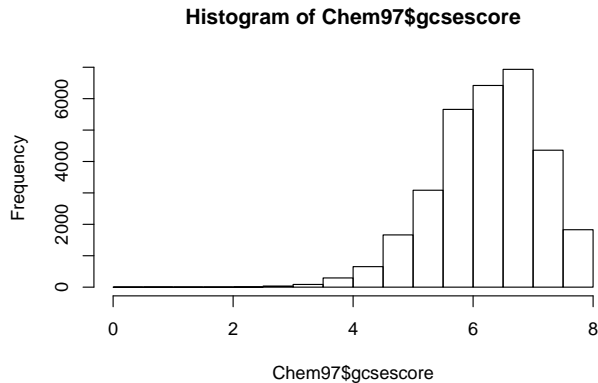
```
> library(lattice)  
> data(Chem97, package = "mlmRev")
```
- ▶ What is the class of `Chem97`?
- ▶ How many variables does it have? *You might want to use `length`*

Formula syntax

- ▶ We'll mostly use three variables: score, gcscscore and gender.
- ▶ Now, **lattice** uses the **formula** syntax.
- ▶ Basically its $y \mid g1$ where x is the variable with the numeric values and $g1$ is a factor.

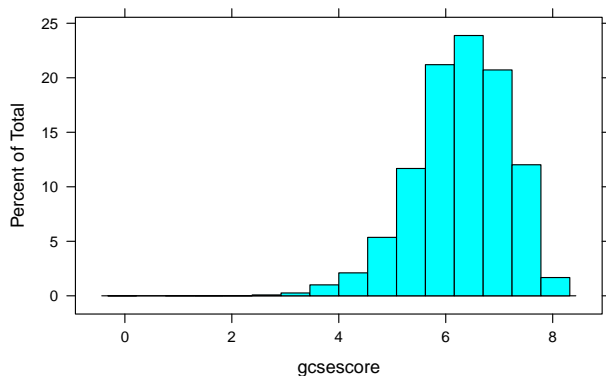
Comparing histograms

```
> hist(Chem97$gcscore)
```



Comparing histograms: part II

```
> print(histogram(~gcsescore, data = Chem97))
```



A grouping var

- ▶ The variable score **only** has values 0, 2, 4, 6, 8 and 10.

```
> head(Chem97$score)
```

```
[1]  4 10 10 10  8 10
```

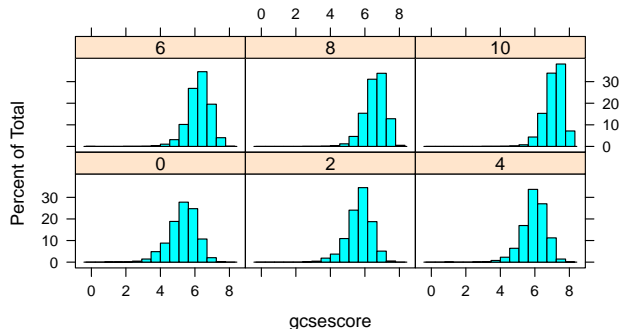
```
> class(Chem97$score)
```

```
[1] "numeric"
```

- ▶ We can use this variable as a factor!
- ▶ Lets make a more interesting plot :)

Multiple hist

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



And gender?

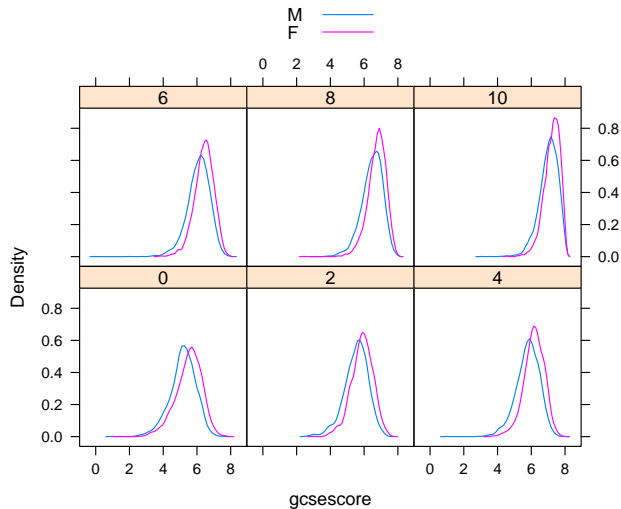
- ▶ But we want to use our third variable: gender

```
> class(Chem97$gender)
[1] "factor"
```
- ▶ Its **difficult** to plot two histograms on the same panel, but that's not the case with density lines!

Densities

```
> print(densityplot(~gcsescore |  
+   factor(score), Chem97, groups = gender,  
+   plot.points = FALSE, auto.key = TRUE))
```

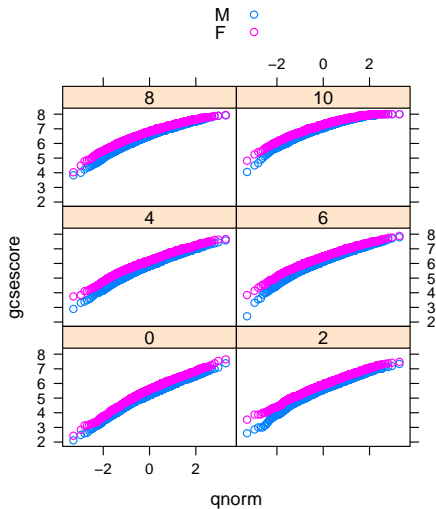

Densities



QQ norm too!

```
> print(qqmath(~gcsescore | factor(score),  
+           Chem97, groups = gender, auto.key = TRUE,  
+           aspect = "xy", f.value = ppoints(1000)))
```

QQ norm too!



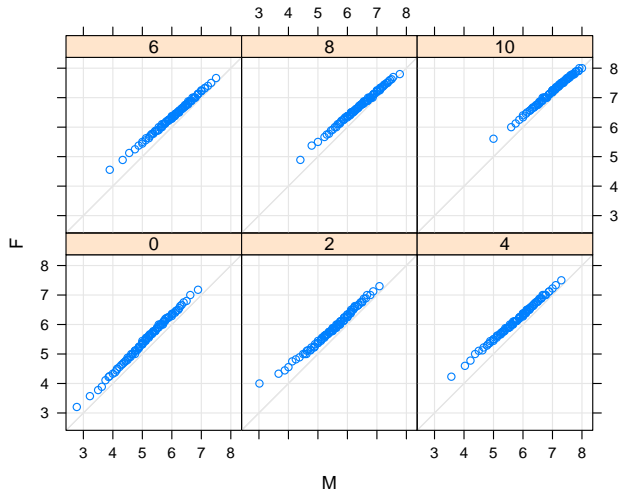
Compare QQ norm

- ▶ Re-do the above QQ norm plot with the following arguments:
 - > *f.value* = *ppoints*(100)
 - > *type* = *c*("p", "g")
- ▶ Which of the two QQ norm plots is clearer?

QQ plots

```
> print(qq(gender ~ gcscscore | factor(score),  
+         Chem97, f.value = ppoints(100),  
+         type = c("p", "g"), aspect = 1))
```

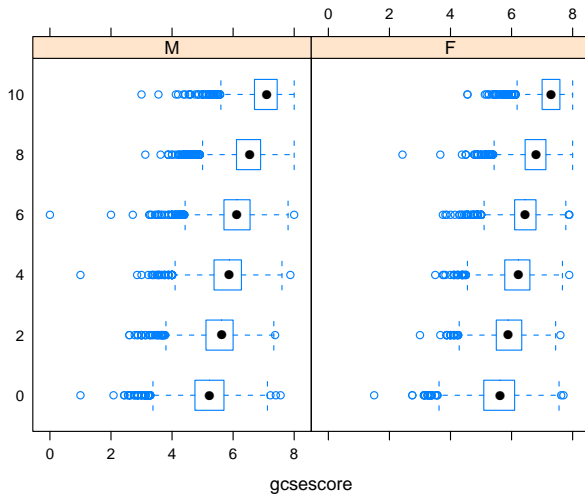
QQ plots



Boxplots

```
> print(bwplot(factor(score) ~ gcsescore |  
+           gender, Chem97))
```

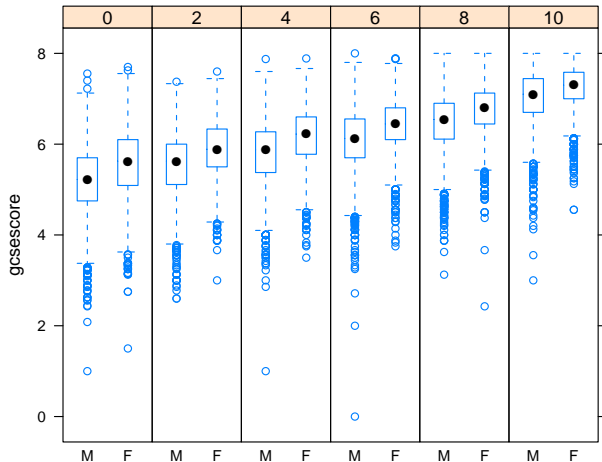
Boxplots



Boxplots II

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

Boxplots II

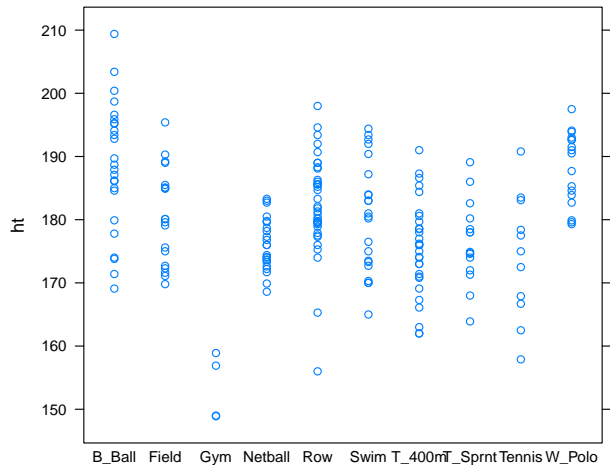


Stripplot

- Useful for small data sets :)

```
> library(DAAG)
> print(stripplot(ht ~ factor(sport),
+               data = ais))
```

Stripplot

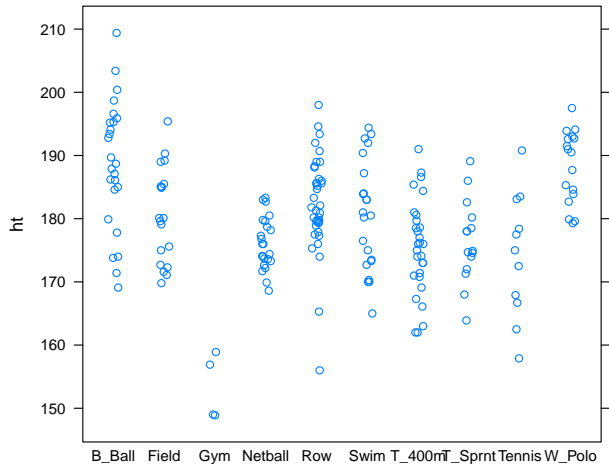


Stripplot II

- ▶ The **jitter** argument saves the day!
- ▶ Plus points in lattice are partially transparent

```
> print(stripplot(ht ~ factor(sport),  
+               data = ais, jitter = T))
```

Stripplot II

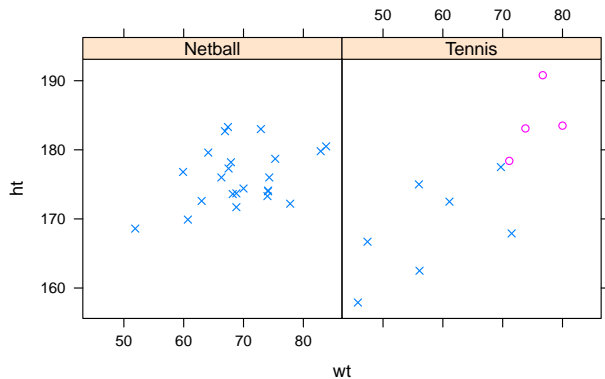


xyplot

- ▶ With lattice, we can also make something similar to **plot**
- ▶ But first, lets create a subset of the type of sports.

```
> subset <- ais$sport %in% c("Netball",  
+   "Tennis")  
> print(xyplot(ht ~ wt | sport, groups = sex,  
+   pch = c(4, 1), aspect = 1,  
+   subset = subset, data = ais))
```

xyplot

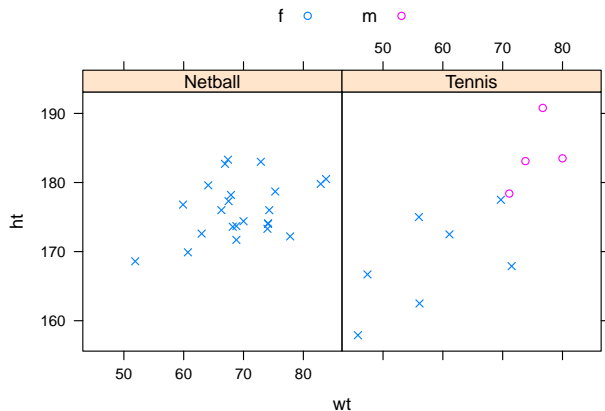


xyplot II

- ▶ What will happen if we say `auto.key=TRUE`?
- ▶ On this plot, we are visualizing data from how many variables?

```
> print(xyplot(ht ~ wt | sport, groups = sex,  
+           pch = c(4, 1), aspect = 1,  
+           auto.key = list(columns = 2),  
+           subset = subset, data = ais))
```

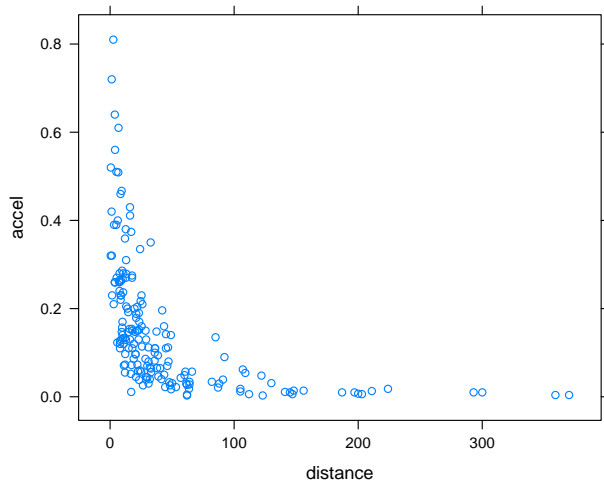
xyplot II



xyplot B

```
> data(Earthquake, package = "nlme")  
> print(xyplot(accel ~ distance,  
+           data = Earthquake))
```

xyplot B

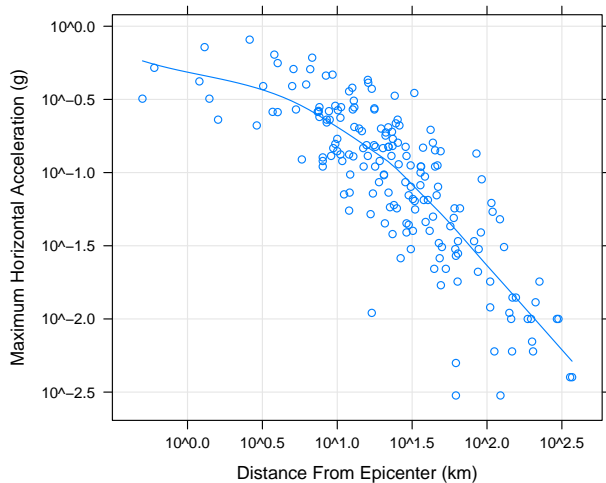


xyplot B II

- ▶ What does the `scales` argument control?
- ▶ What would happen if you delete `smooth` from the `type` argument?

```
> print(xyplot(accel ~ distance,  
+           data = Earthquake, scales = list(log = TRUE),  
+           type = c("p", "g", "smooth"),  
+           xlab = "Distance From Epicenter (km)",  
+           ylab = "Maximum Horizontal Acceleration (g)"))
```

xyplot B II

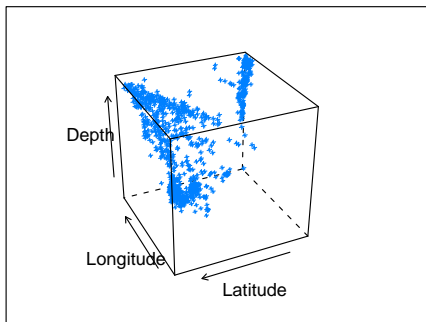


3D!

- ▶ With the **cloud** function its possible to create 3D plots.
- ▶ To rotate it, you need to re-make it with different values for the x, y and z.

```
> print(cloud(depth ~ lat * long,  
+           data = quakes, zlim = rev(range(quakes$depth)),  
+           screen = list(z = 115, x = -60),  
+           panel.aspect = 0.75, xlab = "Longitude",  
+           ylab = "Latitude", zlab = "Depth"))
```

3D!



That's it for lattice

- ▶ **Lattice** has more plot functions such as **barchart** and **dotplot** which we won't cover today, but feel free to check them.
- ▶ There is also a book available on `lattice`:
`http://lmdvr.r-forge.r-project.org/`
- ▶ As I said at the beginning, use the **tools** package to explore `lattice` and `latticeExtra`.

Intro

- ▶ It contains loads of enhanced R functions.
- ▶ The reference manual has 139 pages!!!
- ▶ Functions such as adding a table, standard deviation bars, cutting axes, etc.

Barplot with table

- ▶ First, we'll create a data.frame with some data
- ▶ Then we'll use the **barp** function to create a barplot
- ▶ Finally, we'll add the table to our plot

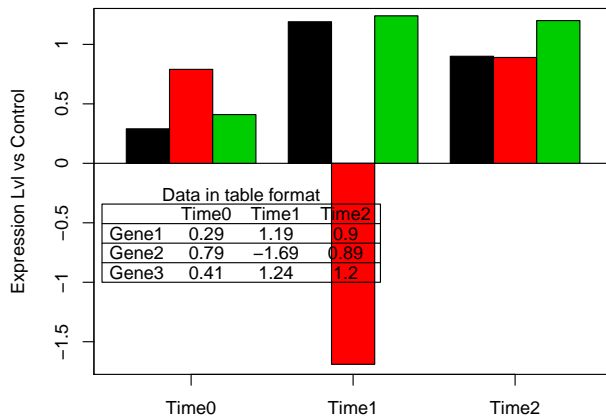
```
> set.seed(123)
> df <- data.frame(Time0 = runif(3),
+   Time1 = rnorm(3), Time2 = rlnorm(3))
> df <- round(df, digits = 2)
> rownames(df) <- c("Gene1", "Gene2",
+   "Gene3")
> df
```

Barplot with table

	Time0	Time1	Time2
Gene1	0.29	1.19	0.90
Gene2	0.79	-1.69	0.89
Gene3	0.41	1.24	1.20

```
> library(plotrix)
> barp(df, ylab = "Expression Lvl vs Control",
+       names.arg = colnames(df), col = 1:3)
> addtable2plot(0.45, -1, df, bty = "o",
+               display.rownames = TRUE, hlines = TRUE,
+               title = "Data in table format")
```

Barplot with table

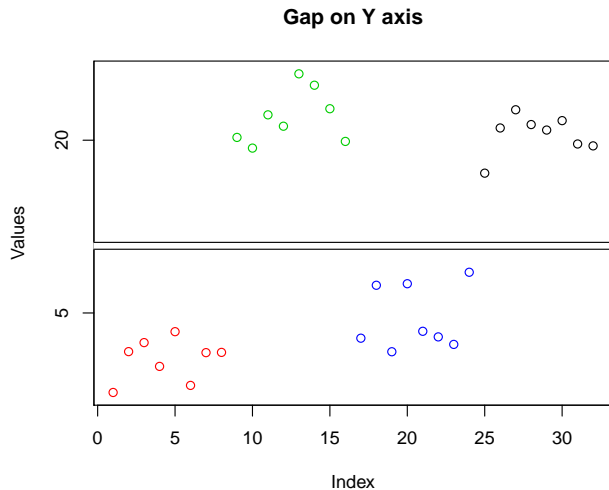


Plot with gaps

- ▶ With Plotrix we can make plots that have a gap on one axis.
- ▶ For example, a normal plot with a gap on the Y axis.

```
> data <- c(rnorm(8) + 3, rnorm(8) +  
+         21, rnorm(8) + 4.5, rnorm(8) +  
+         20)  
> color <- c(rep(2, 8), rep(3, 8),  
+           rep(4, 8), rep(1, 8))  
> gap.plot(data, gap = c(8, 16),  
+         xlab = "Index", ylab = "Values",  
+         main = "Gap on Y axis", col = color)
```

Plot with gaps

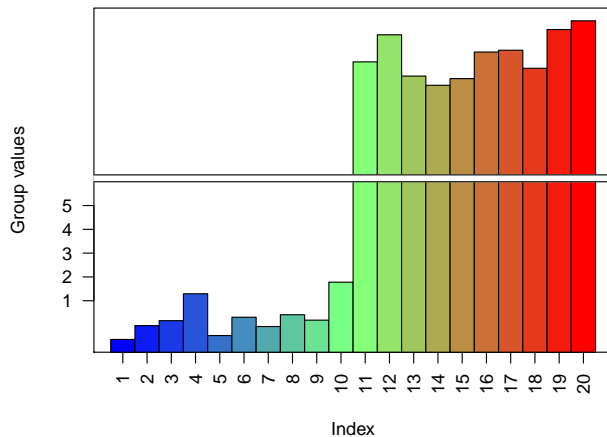


Gap on a barplot

- ▶ Or a barplot with a gap.
- ▶ Very helpful to visualize all your data.
- ▶ However, there is an issue with the labels on the Y axis T_T so be careful when using this kind of plot.

```
> data <- c(rnorm(10), rnorm(10) +  
+          30)  
> gap.barplot(data, gap = c(6, 25),  
+             xlab = "Index", ytics = c(1:30),  
+             ylab = "Group values", las = 2)
```


Gap on a barplot



Error bars

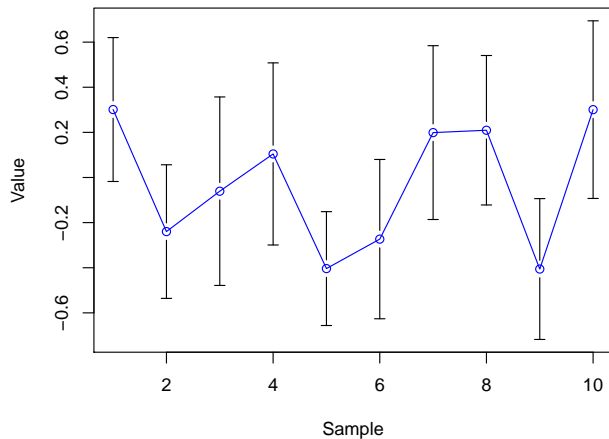
- ▶ Nowadays you get to see lots of graphs with the error bars.
- ▶ Experimental papers generally have 3 to 5 repeats of the same experiment.
- ▶ The **dispersion** function will be helpful to make this kind of plot.

```
> data <- matrix(rnorm(100), 10,  
+               10)  
> a <- colMeans(data)  
> b <- std.error(data)  
> plot(a, ylim = c(min(a - b), max(a +  
+                 b)), xlab = "Sample", ylab = "Value",  
+       col = 4, type = "o")
```

Error bars

```
> dispersion(1:10, colMeans(data),  
+           b)
```

Error bars



Some real data

- ▶ For the next plots, we'll use data from [this article](#) where they sequenced a Korean individual.
- ▶ I already saved as csv files two tables for easy import. We'll load them into R with the `read.csv` function.

```
> t1 <- read.csv("http://www.lcg.unam.mx/~lcollado/B/data/SuppTable01_kogen  
+   header = T)  
> t2 <- read.csv("http://www.lcg.unam.mx/~lcollado/B/data/SuppTable06_nsSnpr  
+   header = T)
```

- ▶ Use `head`, `dim`, `class` to find out more about the data.

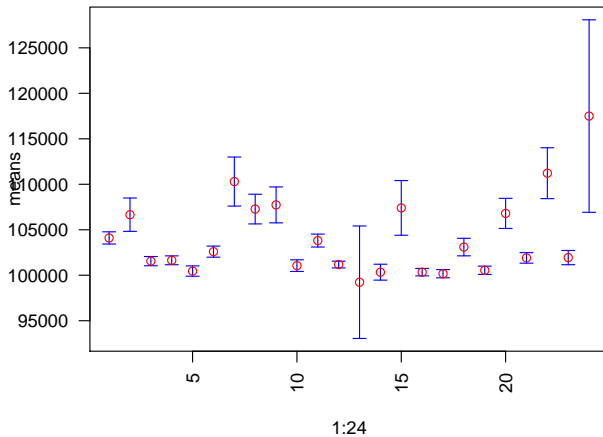
plotCI

- ▶ Plotrix has another function that plots error bars.
- ▶ We'll use our first table and get the data we need using `tapply`.

```
> means <- tapply(t1$bac_size, t1$chrNo,  
+               mean)  
> err <- tapply(t1$bac_size, t1$chrNo,  
+              std.error)  
> plotCI(1:24, means, err, col = "red",  
+        scol = "blue", las = 2, main = "bac_size per chrNo")
```

plotCI

bac_size per chrNo

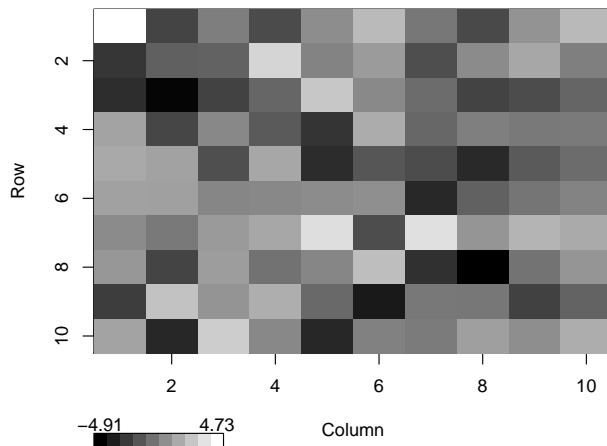


One similar to image

- ▶ With `color2D.matplot` we can make plots very similar to `image`
- ▶ What differences do you notice vs `image`?

```
> mat <- matrix(rnorm(100, 0, 2),  
+             10, 10)  
> color2D.matplot(mat, show.legend = T)
```


One similar to image



Hierobarp

- ▶ We'll use the default example for this powerful plot.

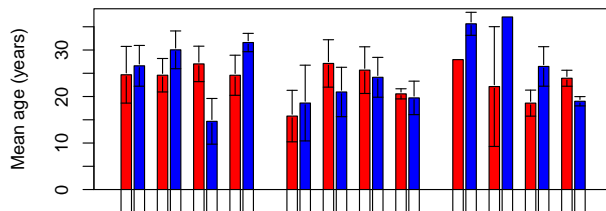
```
> test.df <- data.frame(Age = rnorm(100,  
+ 25, 10), Sex = sample(c("M",  
+ "F"), 100, TRUE), Marital = sample(c("D",  
+ "M", "S", "W"), 100, TRUE),  
+ Employ = sample(c("Full Time",  
+ "Part Time", "Unemployed"),  
+ 100, TRUE))  
> test.col <- list(Overall = "green",  
+ Employ = c("purple", "orange",  
+ "brown"), Marital = c("#1affd8",  
+ "#caeec", "#f7b3cc", "#94ebff"),  
+ Sex = c(2, 4))
```

Hierobarp

```
> hierobarp(formula = Age ~ Sex +  
+           Marital + Employ, data = test.df,  
+           ylab = "Mean age (years)",  
+           main = "Show only the final breakdown",  
+           errbars = TRUE, col = test.col$Sex)
```

Hierobarb

Show only the final breakdown



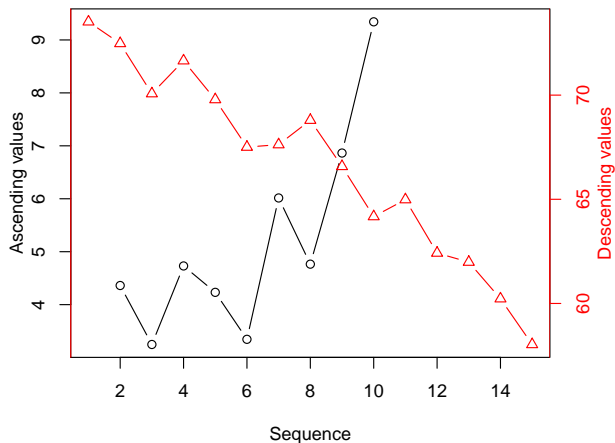
Two scales

- ▶ Sometimes you want two lines with different scales on the same plot.
- ▶ `twoord.plot` is the solution :)

```
> twoord.plot(2:10, seq(3, 7, by = 0.5) +  
+           rnorm(9), 1:15, rev(60:74) +  
+           rnorm(15), xlab = "Sequence",  
+           ylab = "Ascending values",  
+           rylab = "Descending values",  
+           main = "Plot with two ordinates - points and lines")
```

Two scales

Plot with two ordinates – points and lines



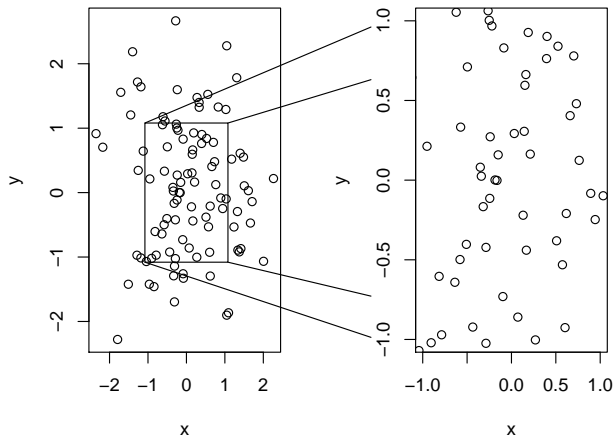
Zoom

- The final plot I'll show you from plotrix enables us to zoom into a section of the plot.

```
> zoomInPlot(rnorm(100), rnorm(100),  
+           rxlim = c(-1, 1), ryylim = c(-1,  
+           1), zoomtitle = "Zoom In Plot")
```

Zoom

Zoom In Plot



Intro

- ▶ **ggplot2** is a much more sophisticated plotting package.
- ▶ **199** pages long ref manual!!!
- ▶ Lets take a look at some examples.

Plotmatrix

- ▶ We'll use the **iris** data set which is used quite frequently to exemplify scatterplots.
- ▶ Meaning that you are using 3 or more variables.
- ▶ Explore `iris` with `head` and other similar functions.

```
> plotmatrix(iris[, 1:4])
```

Plotmatrix II

- ▶ If we combine `plotmatrix` with `geom_smooth` we can get a much better graph.

```
> plotmatrix(iris[, 1:4]) + geom_smooth(method = "lm")
```

We'll be back

- ▶ On the class where we'll learn about linear regressions, we'll be back and make plots like this one:

```
> mod <- lm(mpg ~ wt, data = mtcars)
> qqplot(.fitted, .resid, data = mod) +
+       geom_hline() + geom_smooth(se = FALSE)
```

Intro

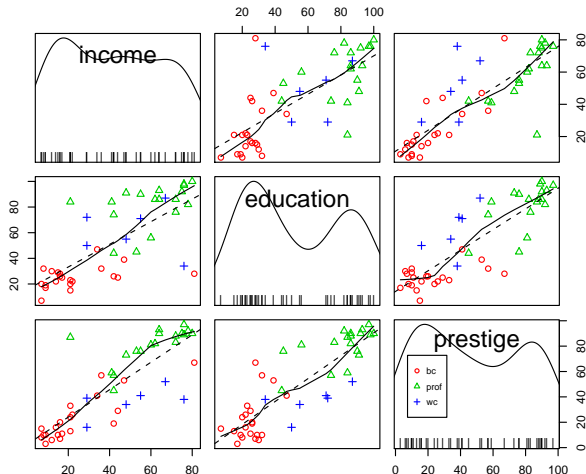
- ▶ While this package has quite a lot of functions too (105 page ref man), one special plot caught my eye.
- ▶ Feel free to check all the examples later if you want :D

scatterplot.matrix

- ▶ Quite similar plot to some we made before with automatic colors

```
> library(car)
> scatterplot.matrix(~income + education +
+   prestige | type, data = Duncan)
```

scatterplot.matrix

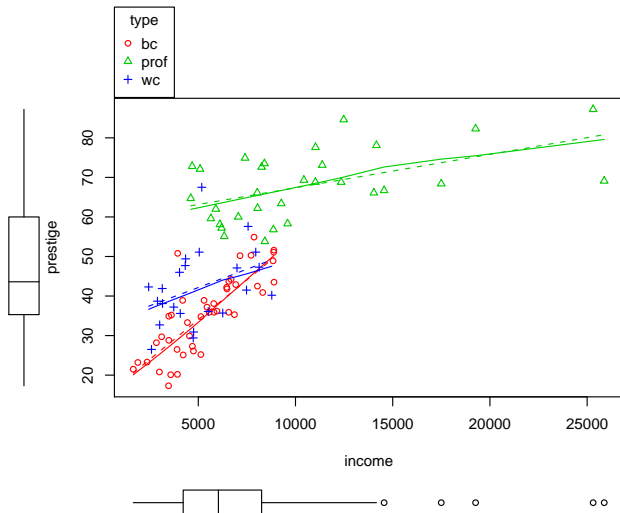


scatterplot

- ▶ With **scatterplot** we can create boxplots on our axis!!

```
> scatterplot+      type, data = Prestige, span = 1)
```


scatterplot



Session Info

```
> sessionInfo()
```

```
R version 2.10.0 Under development (unstable) (2009-07-21 r48968)  
i386-pc-mingw32
```

```
locale:
```

```
[1] LC_COLLATE=English_United States.1252  
[2] LC_CTYPE=English_United States.1252  
[3] LC_MONETARY=English_United States.1252  
[4] LC_NUMERIC=C  
[5] LC_TIME=English_United States.1252
```

```
attached base packages:
```

```
[1] stats      graphics  grDevices  
[4] utils      datasets  methods  
[7] base
```

```
other attached packages:
```

```
[1] car_1.2-15  
[2] plotrix_2.6-4
```

Session Info

```
[3] DAAG_1.00  
[4] randomForest_4.5-30  
[5] rpart_3.1-44  
[6] MASS_7.3-0  
[7] lattice_0.17-25
```

loaded via a namespace (and not attached):

```
[1] grid_2.10.0  tools_2.10.0
```