

# Summer Institute in Statistical Genetics Module 6: Computing for Statistical Genetics

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2. Learning to Draw

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R (and S-PLUS) can produce graphics in many formats, including:

- on screen
- PDF files for LAT<sub>E</sub>X or emailing to people
- PNG or JPEG bitmap formats for web pages (or on non-Windows platforms to produce graphics for MS Office). PNG is also useful for graphs of large data sets.
- On Windows, metafiles for Word, Powerpoint, and similar programs

Graphs should usually be designed on the screen and then may be replotted on eg a PDF file (for Word/Powerpoint you can just copy and paste)

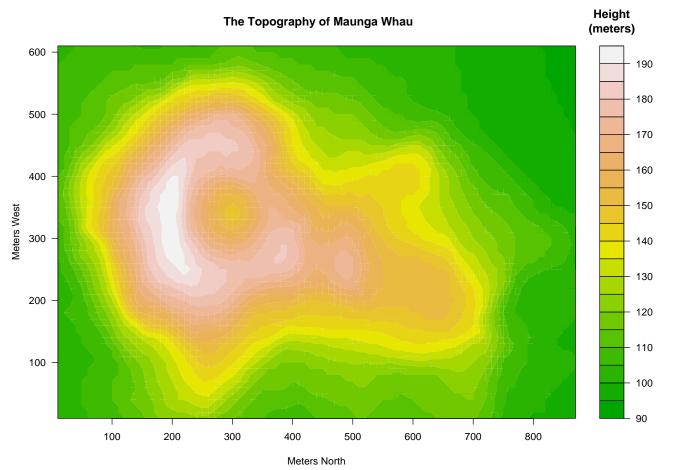
For printed graphs, you will get better results if you design the graph at the size it will end up, eg:

```
## on Windows
windows(height=4,width=6)
## on Unix
x11(height=4,width=6)
```

Word or  $IAT_EX$  can rescale the graph, but when the graph gets smaller, so do the axis labels...

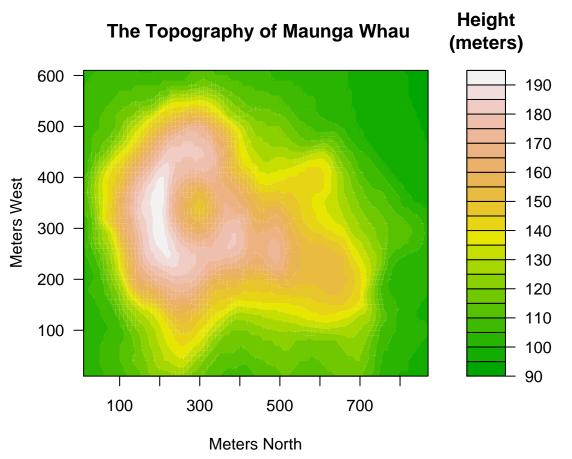
# Setting up a graph

#### Created at full-page size



filled.contour(.) from R version 2.5.1 (2007-06-27)

Created at 5x6 inches



filled.contour(.) from R version 2.5.1 (2007-06-27)

After you have the right commands to draw the graph you can produce it in another format: eg

## start a PDF file
pdf("picture.pdf",height=4,width=6)
## your drawing commands here
....
#### close the PDF file
dev.off()

# Drawing

Usually use plot() to create a graph and then lines(), points(), legend(), text(), and other commands to annotate it.

plot() is a generic function : it does appropriate things for different types of input

## scatterplot
plot(salary\$year, salary\$salary)
## boxplot
plot(salary\$rank, salary\$salary)
## stacked barplot
plot(salary\$field, salary\$rank)

and others for other types of input.

The plot() command can be written

plot(salary~rank, data=salary)

introducing the *formula* system that is also used for regression models. The variables in the formula are automatically looked up in the data= argument.

Two important aspects of designing a graph

- It should have something to say
- It should be legible

Having something to say is your problem; software can help with legibility.

Important points

- Axes need labels (with units, large enough to read)
- Color can be very helpful (but not if the graph is going to be printed in black and white).
- Different line or point styles usually should be labelled.
- Points plotted on top of each other won't be seen

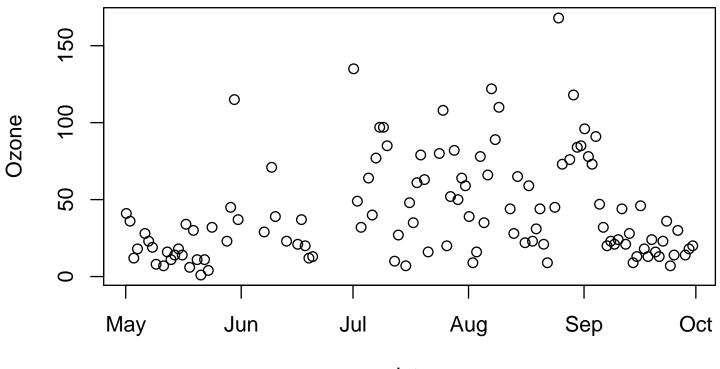
After these are satisfied, it can't hurt to have the graph look nice.

Set up a data set: daily ozone concentrations in New York, summer 1973

```
data(airquality)
names(airquality)
airquality$date<-with(airquality, ISOdate(1973,Month,Day))</pre>
```

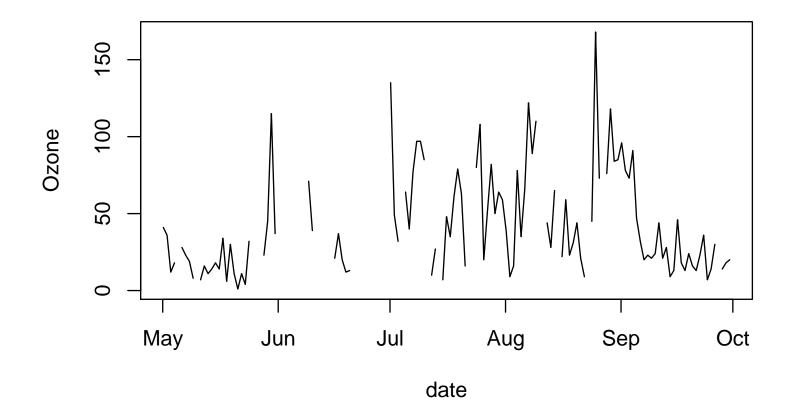
All these graphs were designed at  $4in\times 6in$  and stored as PDF files

plot(Ozone~date, data=airquality)

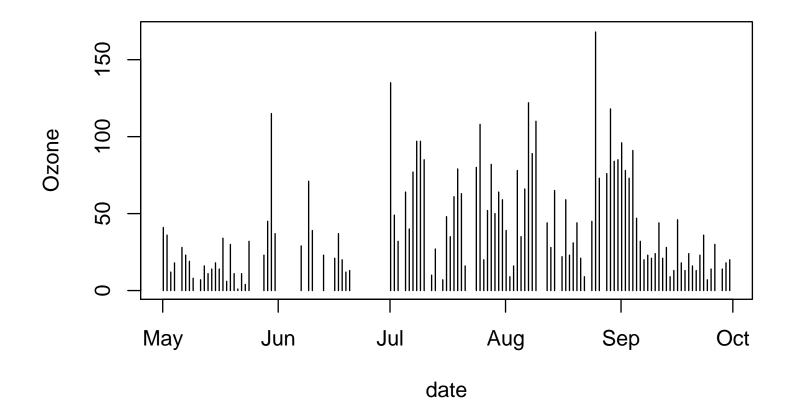


date

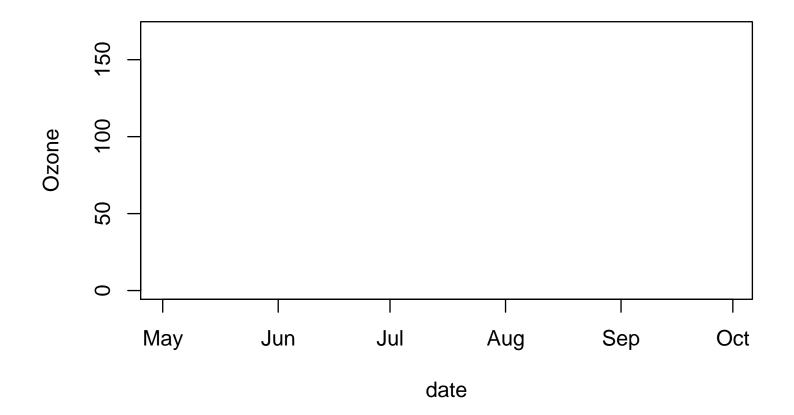
plot(Ozone~date, data=airquality,type="l")



plot(Ozone~date, data=airquality,type="h")

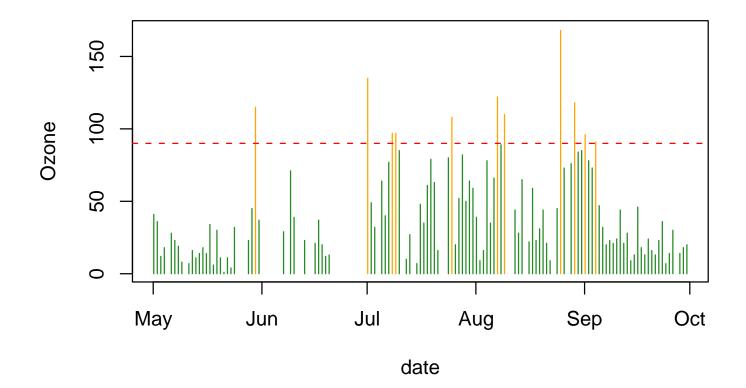


plot(Ozone~date, data=airquality,type="n")

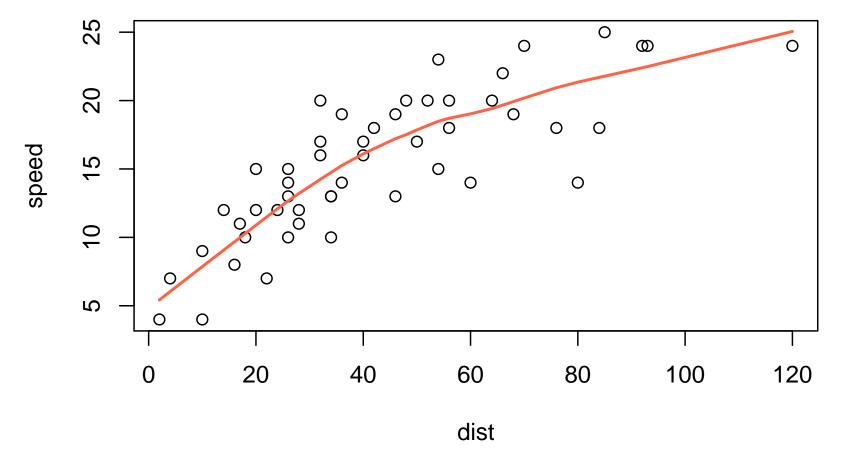


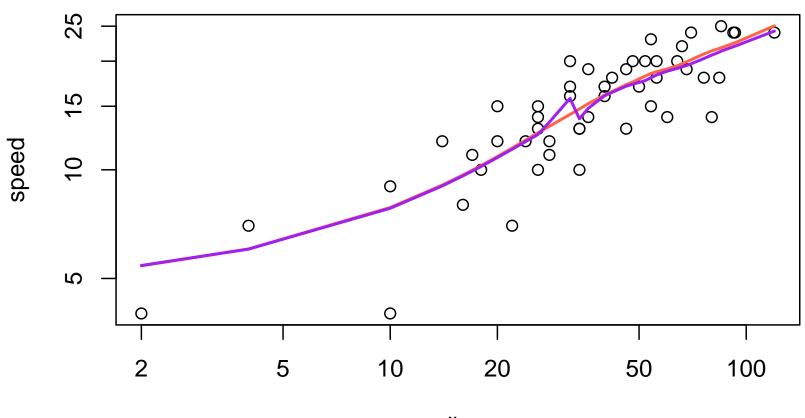
# **Options for** plot()

bad<-ifelse(airquality\$0zone>=90, "orange","forestgreen")
plot(0zone~date, data=airquality,type="h",col=bad)
abline(h=90,lty=2,col="red")

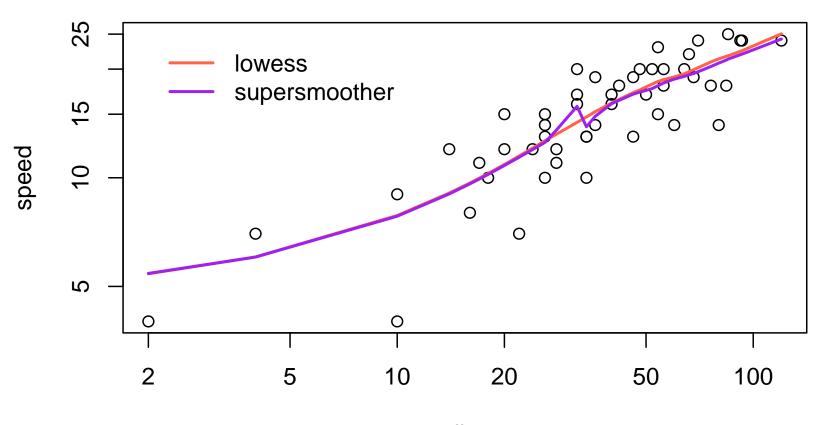


- type= controls how data are plotted. type="n" is not as useless as it looks: it can set up a plot for latter additions.
- Colors can be specified by name (the colors() function gives all the names), by red/green/blue values (#rrggbb with six base-sixteen digits) or by position in the standard palette of 8 colors.
- abline draws a single straight line on a plot
- ifelse() selects between two vectors based on a logical variable.
- lty specifies the line type: 1 is solid, 2 is dashed, 3 is dotted, then it gets more complicated.





dist

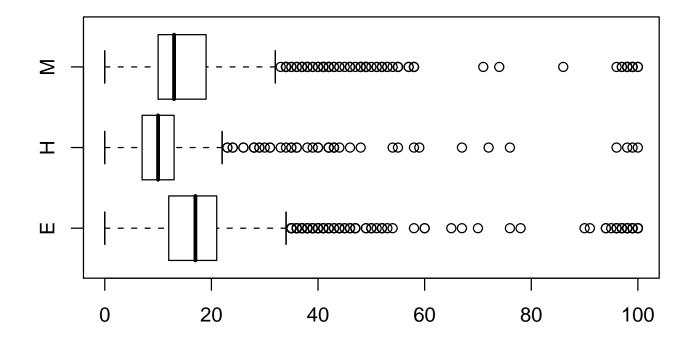


dist

- lines adds lines to an existing plot ( points() adds points).
- lowess() and supsmu() are scatterplot smoothers. They draw smooth curves that fit the relationship between y and x locally.
- log="xy" asks for both axes to be logarithm (log="x" would just be the x-axis)
- legend() adds a legend

# **Boxplots**

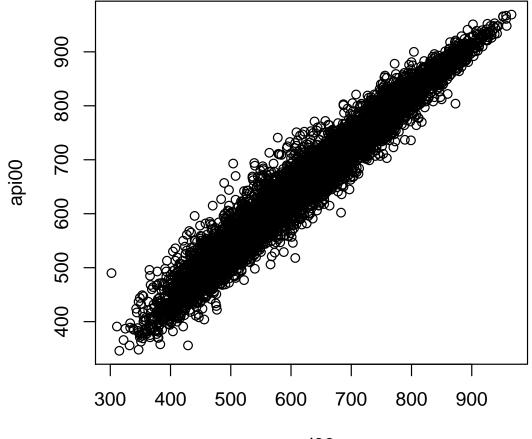
data(api, package="survey")
boxplot(mobility~stype,data=apipop, horizontal=TRUE)



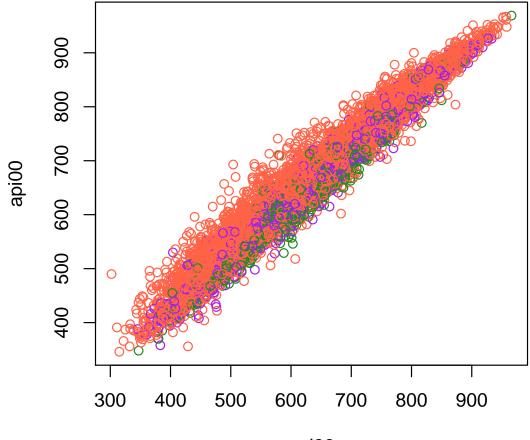
- boxplot computes and draws boxplots.
- horizontal=TRUE turns a boxplot sideways
- xlab and ylab are general options for x and y axis labels.

Scatterplots quickly get crowded. For example, the California Academic Performance Index is reported on 6194 schools

- > plot(api00~api99,data=apipop)
- > colors<-c("tomato","forestgreen","purple")[apipop\$stype]</pre>
- > plot(api00~api99,data=apipop,col=colors)



api99



api99

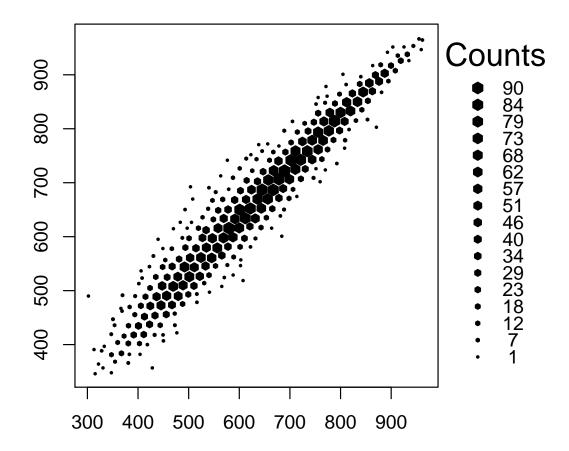
For a single large scatterplot some form of aggregation is useful

library(hexbin)
with(apipop, plot(hexbin(api99,api00), style="centroids"))

hexbin is in the hexbin package from the Bioconductor project. It computes the number of points in each hexagonal bin.

The style=centroids plot puts a filled hexagon with size depending on the number of points at the centroid of the points in the bin.

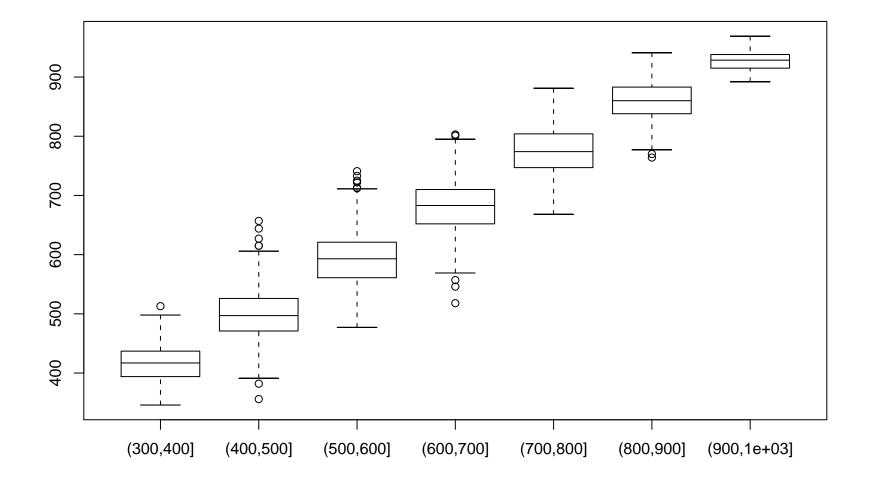
### **Density plots**

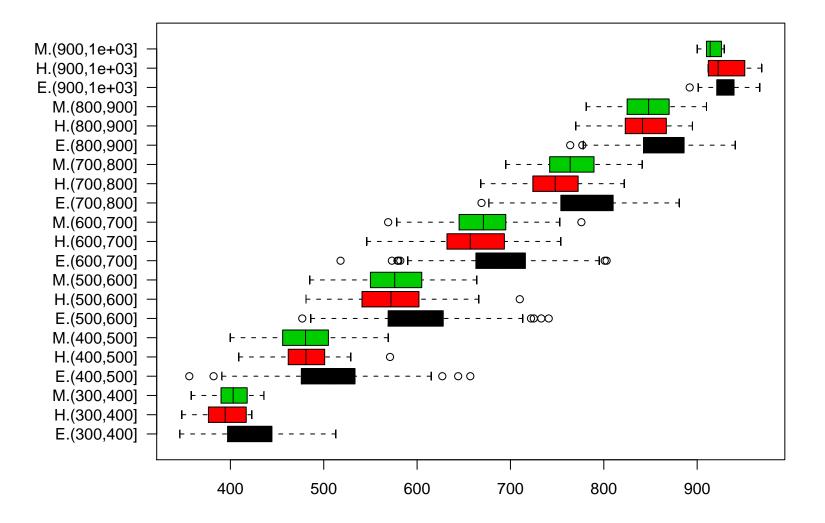


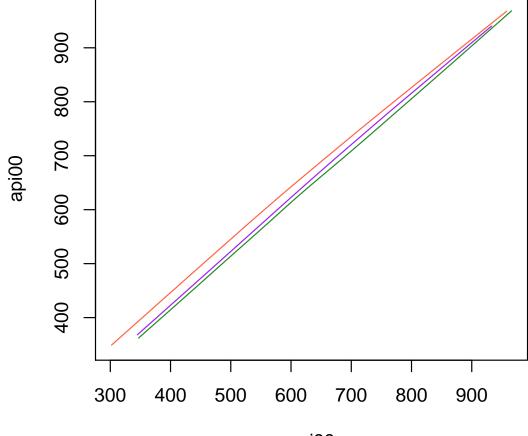
For showing multiple groups a scatterplot smoother or perhaps boxplots would be better.

- > boxplot(api00~cut(api99,(3:10)\*100), data=apipop)
- > par(las=1)
- > par(mar=c(5.1,10.1,2.1,2.1))
- > boxplot(api00~interaction(stype,

```
cut(api99,(3:10)*100)),
data=apipop, horizontal=TRUE,col=1:3)
plot(api00~api99,data=apipop,type="n")
with(subset(apipop, stype=="E"),
lines(lowess(api99, api00), col="tomato"))
with(subset(apipop, stype=="H"),
lines(lowess(api99, api00), col="forestgreen"))
with(subset(apipop, stype=="M"),
lines(lowess(api99, api00), col="purple"))
```







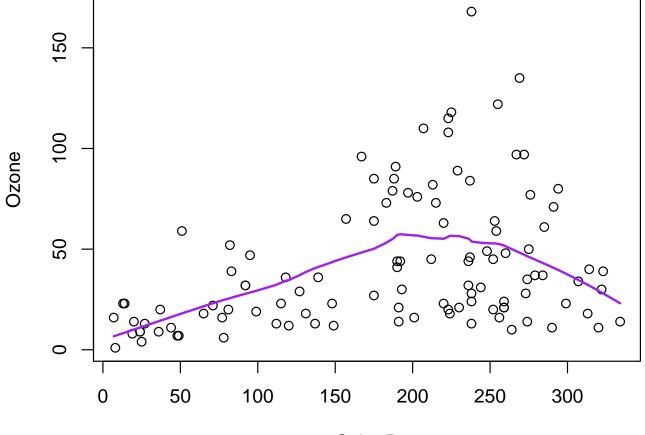
api99

- cut turns a variable into a factor by cutting it at the specified points.
- Note the use of type="n"
- par(mar=) sets the margins around the plot. We need a large left margin for the labels.
- subset takes a subset of a data frame.

Ozone is a *secondary pollutant*, it is produced from organic compounds and atmostpheric oxygen in reactions catalyzed by nitrogen oxides and powered by su nlight.

However, looking at ozone concentrations in NY in summer we see a non-monotone relationship with sunlight

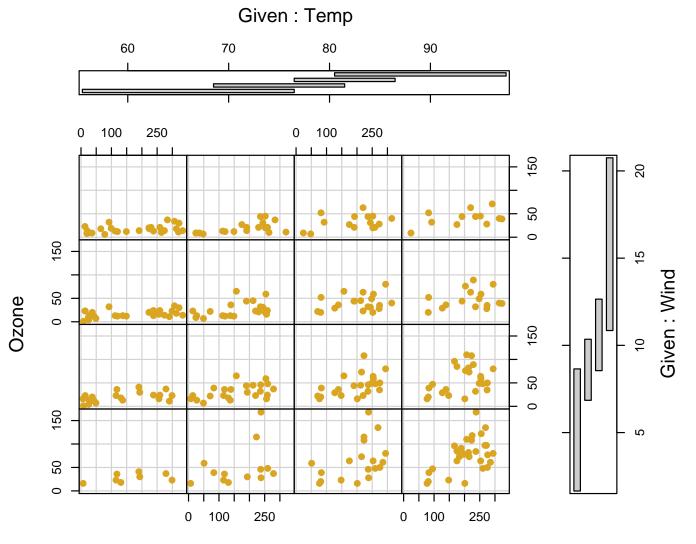
# **Conditioning plots**



Solar.R

Here we draw a scatterplot of Ozone vs Solar.R for various subranges of Temp and Wind . A simple version of what is possible with the Trellis system.

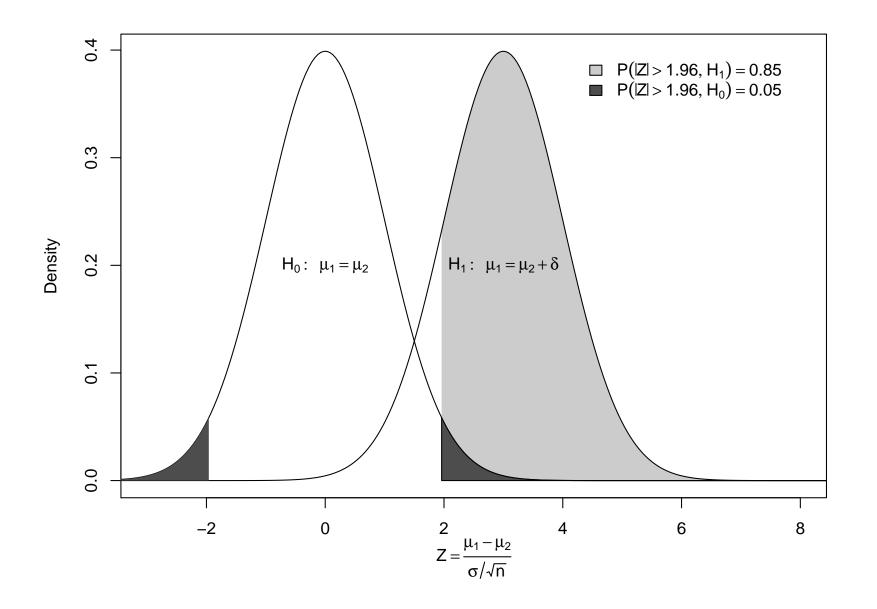
# Conditioning plots



Solar.R

```
An expression can be specified in R for any text in a graph (
help(plotmath) for details). Here we annotate a graph drawn
with polygon.
x<-seq(-10,10,length=400)
y1 < -dnorm(x)
y_2 < -dnorm(x, m=3)
par(mar=c(5,4,2,1))
plot(x, y2, xlim=c(-3, 8), type="n",
   xlab=quote(Z==frac(mu[1]-mu[2],sigma/sqrt(n))),
   ylab="Density")
polygon(c(1.96,1.96,x[240:400],10),
       c(0,dnorm(1.96,m=3),y2[240:400],0),
       col="grey80",lty=0)
lines(x,y2)
lines(x,y1)
polygon(c(-1.96,-1.96,x[161:1],-10),
       c(0,dnorm(-1.96,m=0),y1[161:1],0),
       col="grey30",lty=0)
polygon(c(1.96,1.96,x[240:400],10),
        c(0,dnorm(1.96,m=0),y1[240:400],0),
       col="grey30")
```

```
legend(4.2,.4,fill=c("grey80","grey30"),
    legend=expression(P(abs(Z)>1.96,H[1])==0.85,
        P(abs(Z)>1.96,H[0])==0.05),bty="n")
text(0,.2,quote(H[0]:~~mu[1]==mu[2]))
text(3,.2,quote(H[1]:~~mu[1]==mu[2]+delta))
```



#### **Proportion Hispanic**

