

# Graphics

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## Agenda

- Basic graphics
- Custom graphics

## Functions for graphics

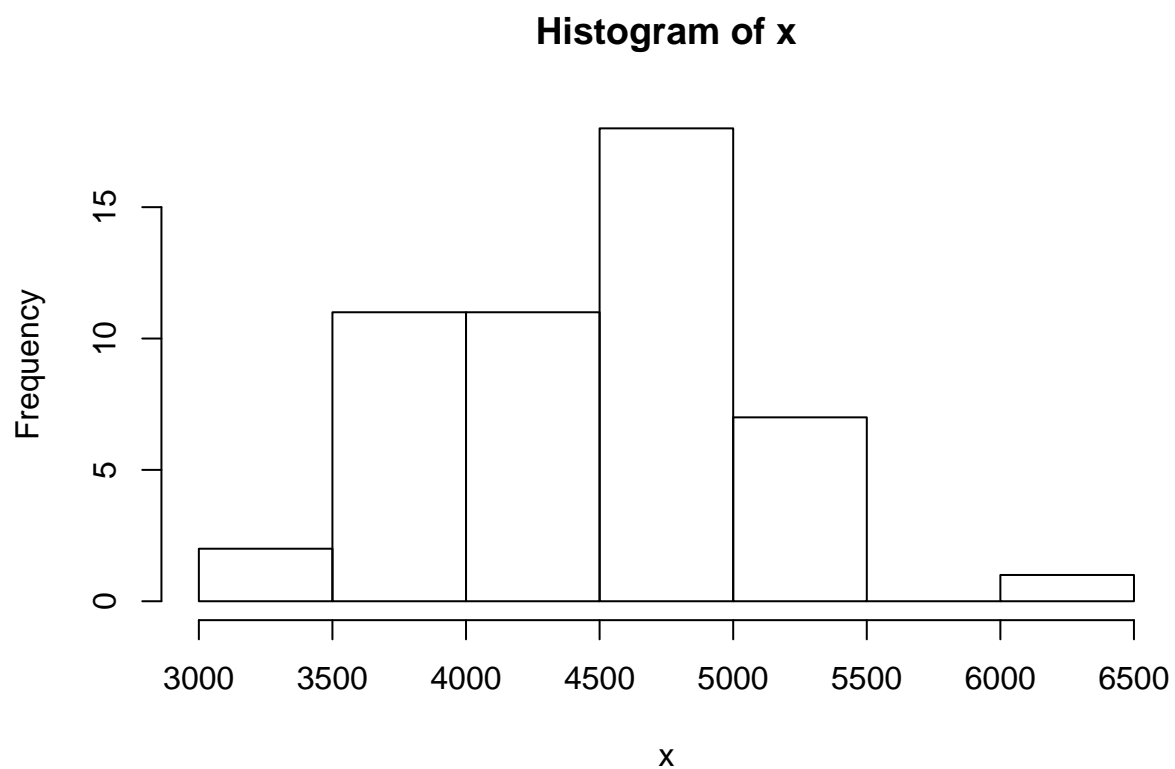
- The functions `hist()`, `boxplot()`, `plot()`, `points()`, `lines()`, `text()`, `mtext()`, `axis()`, etc. form a suite that plot graphs and add features to the graph
- Each of these functions have various options, to learn more about them, use the help
- `par()` can be used to set or query graphical parameters

## Basic graphics

- Univariate data:
  - continuous: density plots(histogram and kernel density plots), ecdf
  - Categorical: pie charts, bar charts
- Bivariate data:
  - Two continuous data: Scatterplots, qqplots
  - One continuous and one categorical data: boxplots
  - Two categorical data: stacked/grouped bar charts
- Trivariate data:
  - pairwise plots
  - 3D plots(image plots, contour plots)

## Univariate data: Histogram

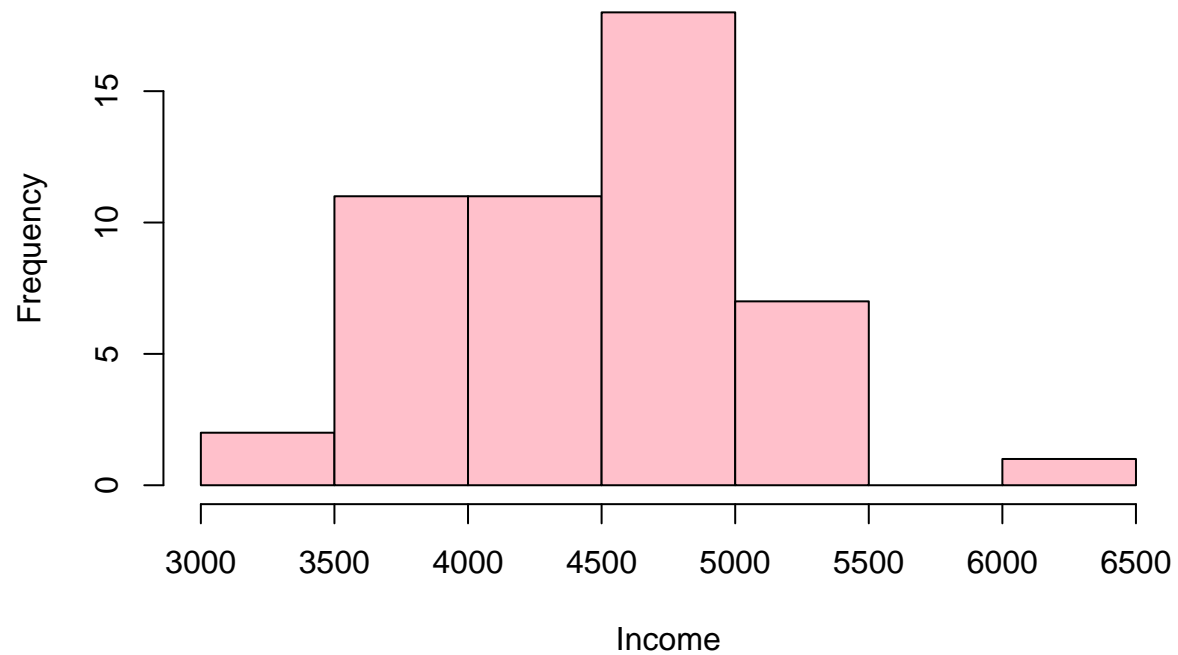
```
x <- state.x77[, 2]           # 50 average state incomes in 1977
hist(x)
```



Univariate data: Histogram

```
hist(x, breaks = 8, col="pink", xlab="Income", main="Histogram of State Income in 1977")
```

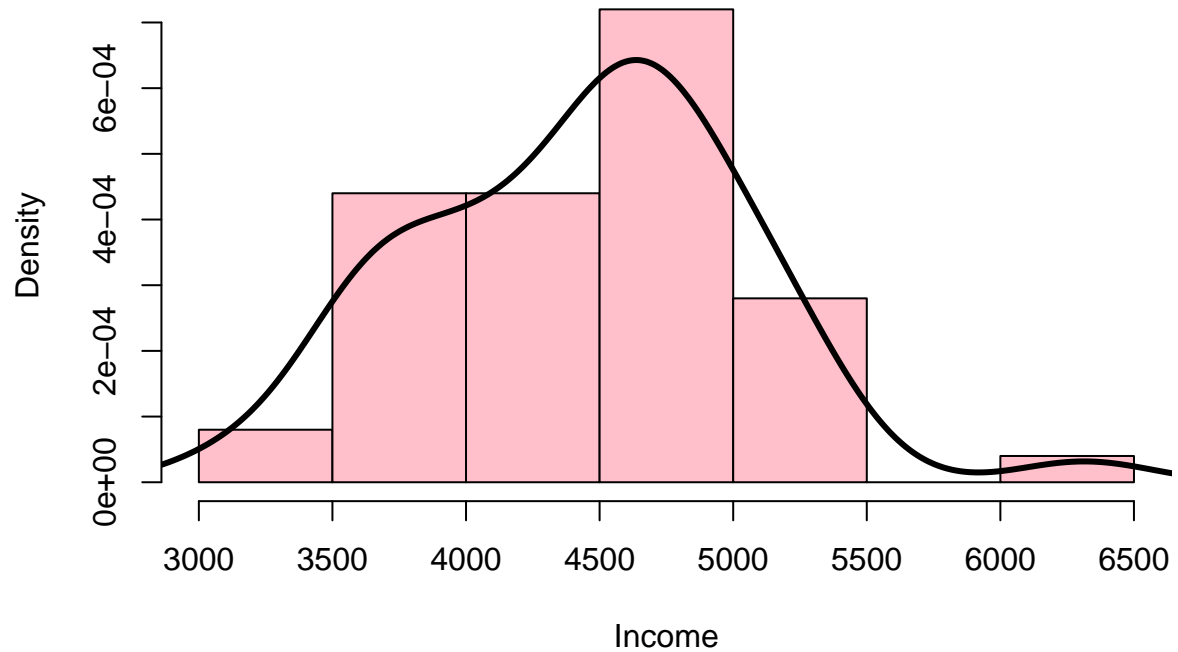
## Histogram of State Income in 1977



Univariate data: Histogram with a density plot

```
hist(x, breaks = 8, col="pink", freq = FALSE, xlab="Income", main="Histogram of State Income in 1977")  
lines(density(x), lwd=3)
```

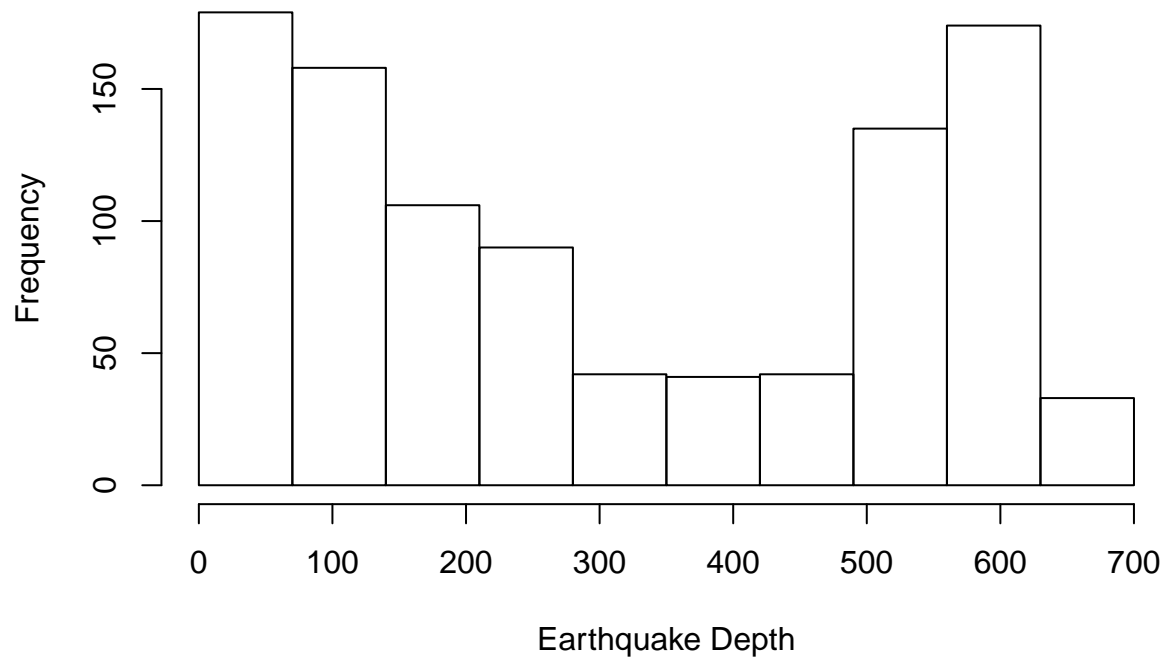
## Histogram of State Income in 1977



## Univariate data: Histogram

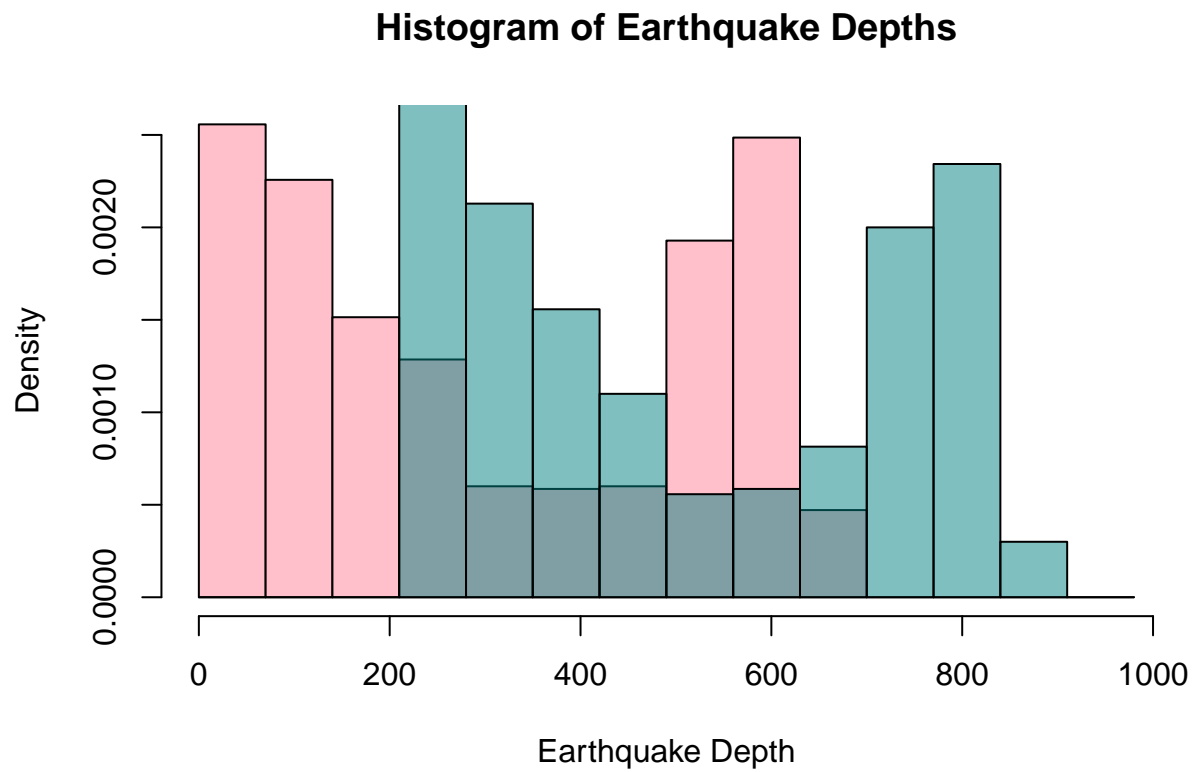
```
y <- quakes$depth                                     # 1000 earthquake depths
hist(y, seq(0, 700, by = 70), xlab="Earthquake Depth", main="Histogram of Earthquake Depths")
```

## Histogram of Earthquake Depths



### Univariate data: Histogram

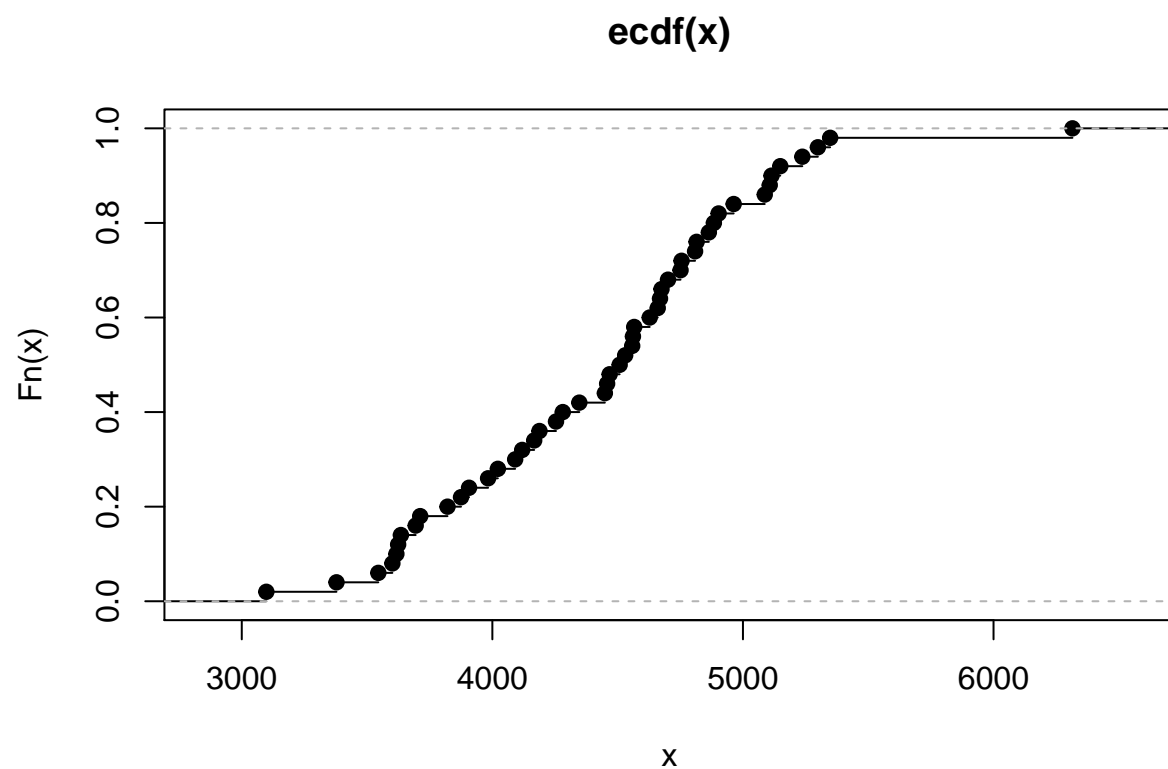
```
y <- quakes$depth                                     # 1000 earthquake depths
hist(y, seq(0, 1000, by = 70), freq = FALSE, col = "pink",
     xlab="Earthquake Depth", main="Histogram of Earthquake Depths")
hist(y+ 200, seq(0, 1000, by = 70), freq = FALSE, col = rgb(0,0.5, 0.5, 0.5), add = TRUE)
```



### Univariate data: Empirical CDF

Function `plot.ecdf()` provides data for empirical cdf

```
plot.ecdf(x)
```

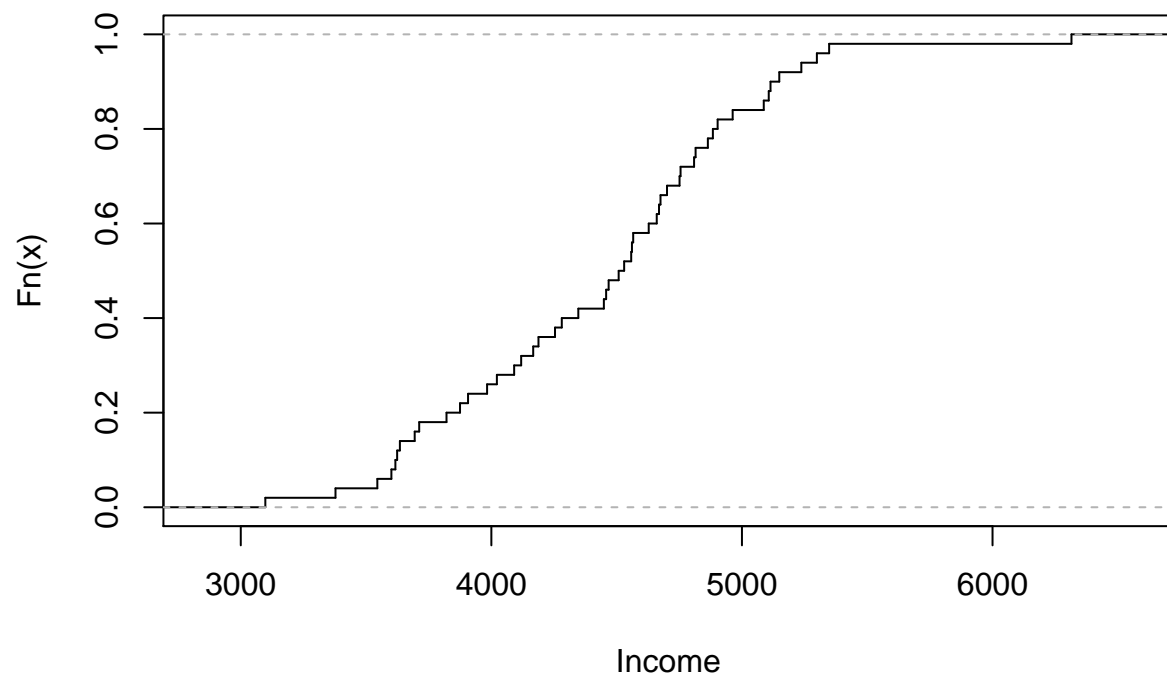


## Univariate data: Empirical CDF

Can add vertical lines and remove dots

```
plot.ecdf(x, verticals = T, pch = "", xlab = "Income",  
          main = "ECDF of State Income in 1977")
```

## ECDF of State Income in 1977

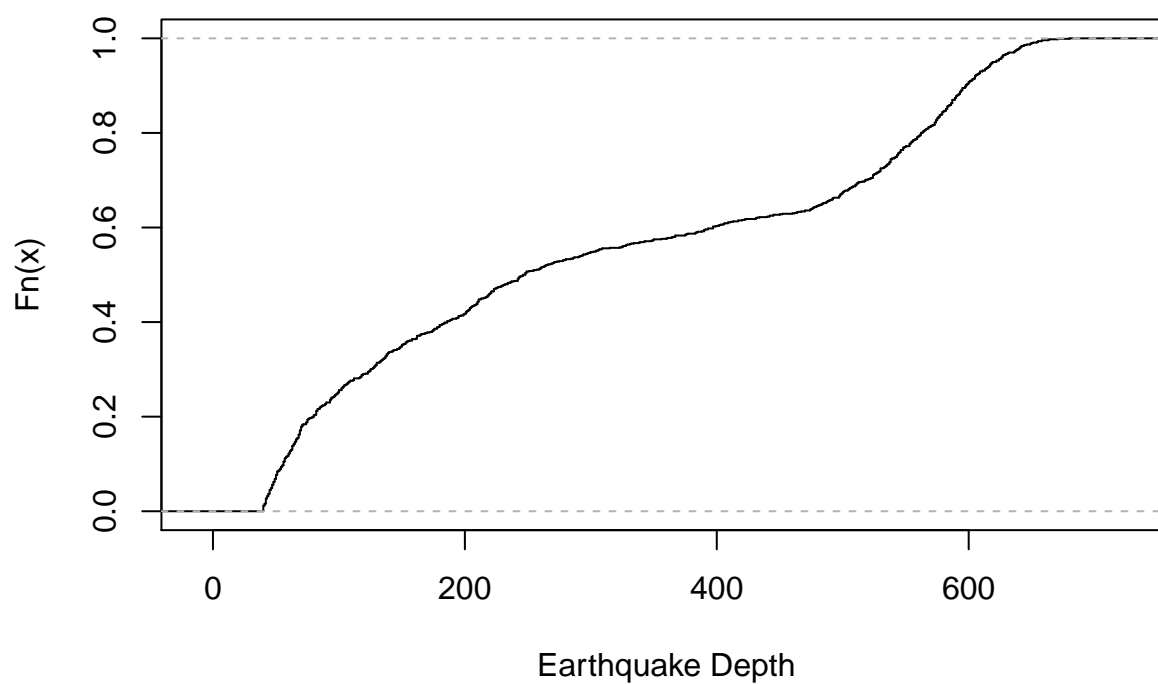


## Univariate data: Empirical CDF

```
plot.ecdf(y, verticals = T, pch = "", xlab = "Earthquake Depth",  
          main = "ECDF of Earthquake Depths")
```

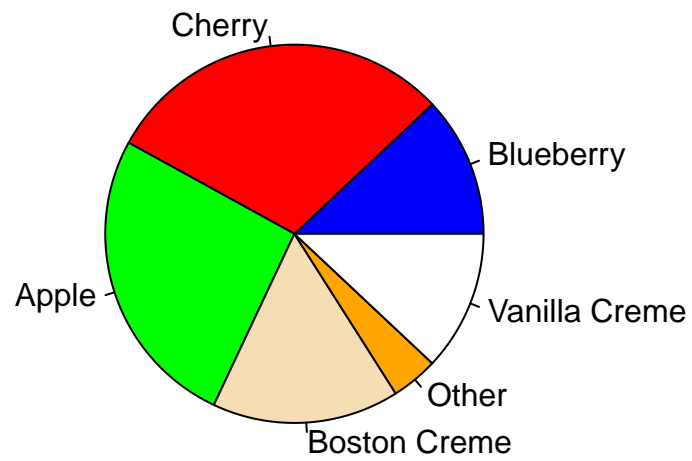


## ECDF of Earthquake Depths



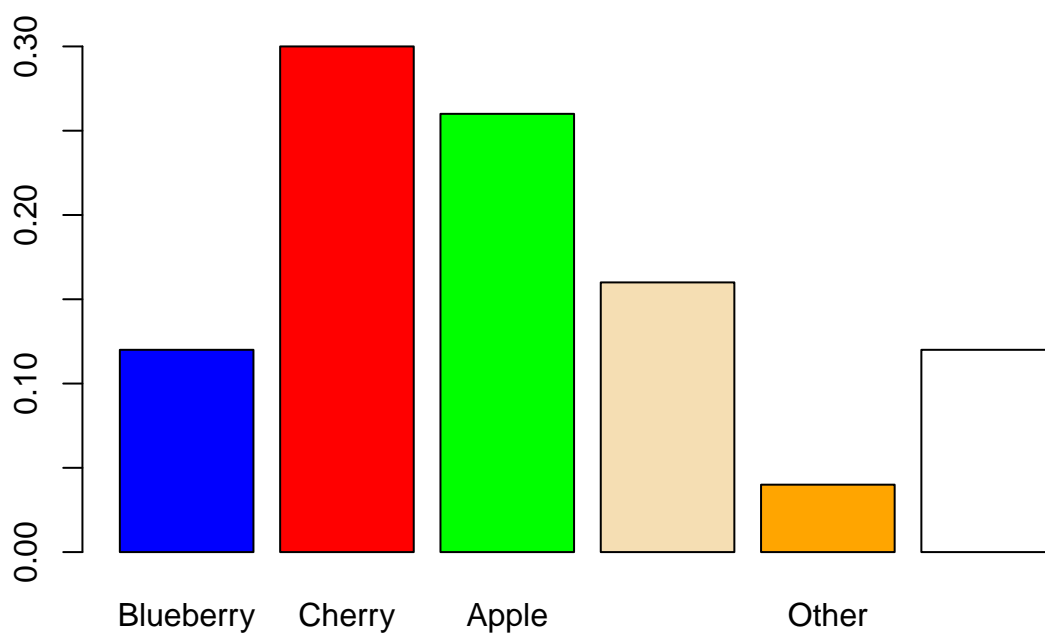
## Univariate data: Pie charts

```
pie.sales = c(0.12, 0.30, 0.26, 0.16, 0.04, 0.12)
names(pie.sales) = c("Blueberry", "Cherry", "Apple", "Boston Creme",
                    "Other", "Vanilla Creme")
pie(pie.sales, col = c("blue", "red", "green", "wheat", "orange", "white"))
```

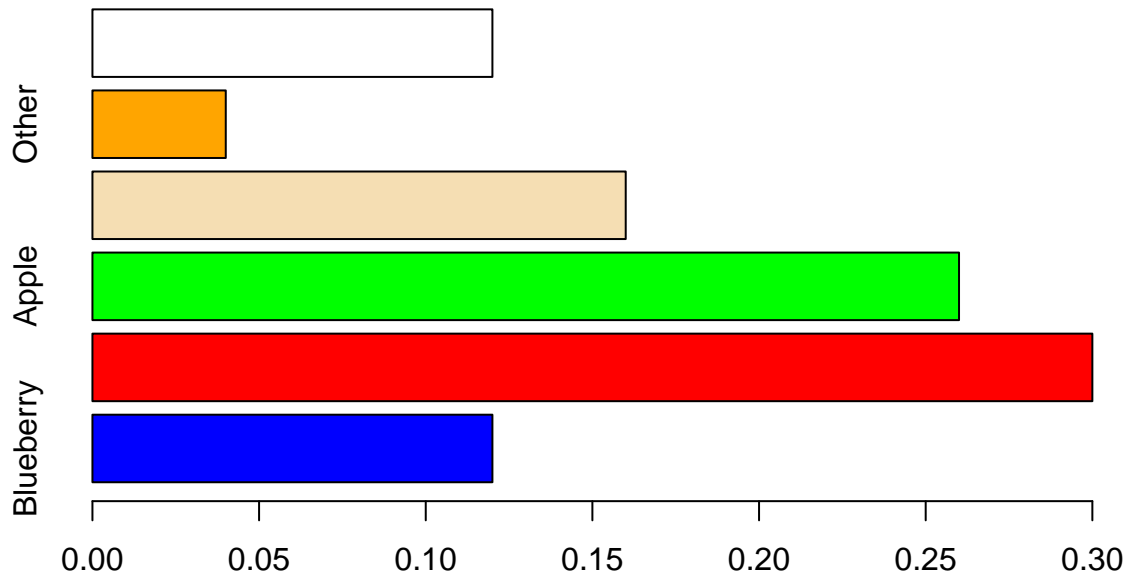


Univariate data: bar charts

```
barplot(pie.sales, col = c("blue", "red", "green", "wheat", "orange", "white"))
```



```
barplot(pie.sales, col = c("blue", "red", "green", "wheat", "orange", "white"), horiz = TRUE)
```

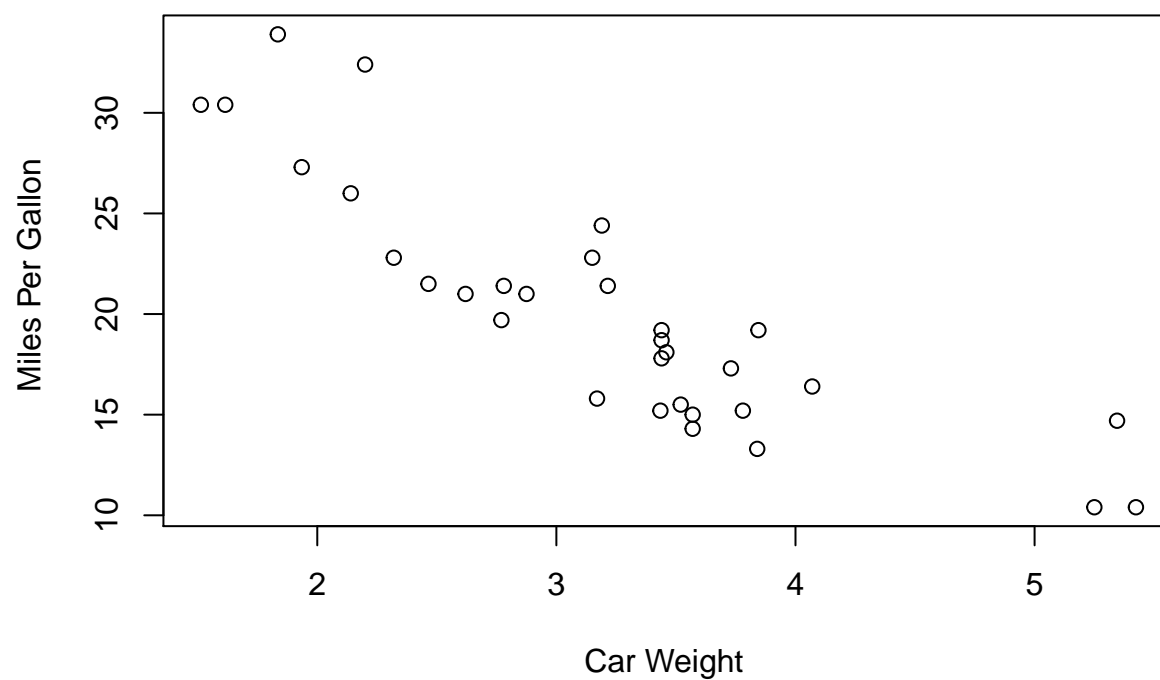


## Bivariate data

Scatterplots: `plot(x, y)`

```
# Simple Scatterplot  
attach(mtcars)  
plot(wt, mpg, main="Scatterplot Example",  
      xlab="Car Weight ", ylab="Miles Per Gallon ")
```

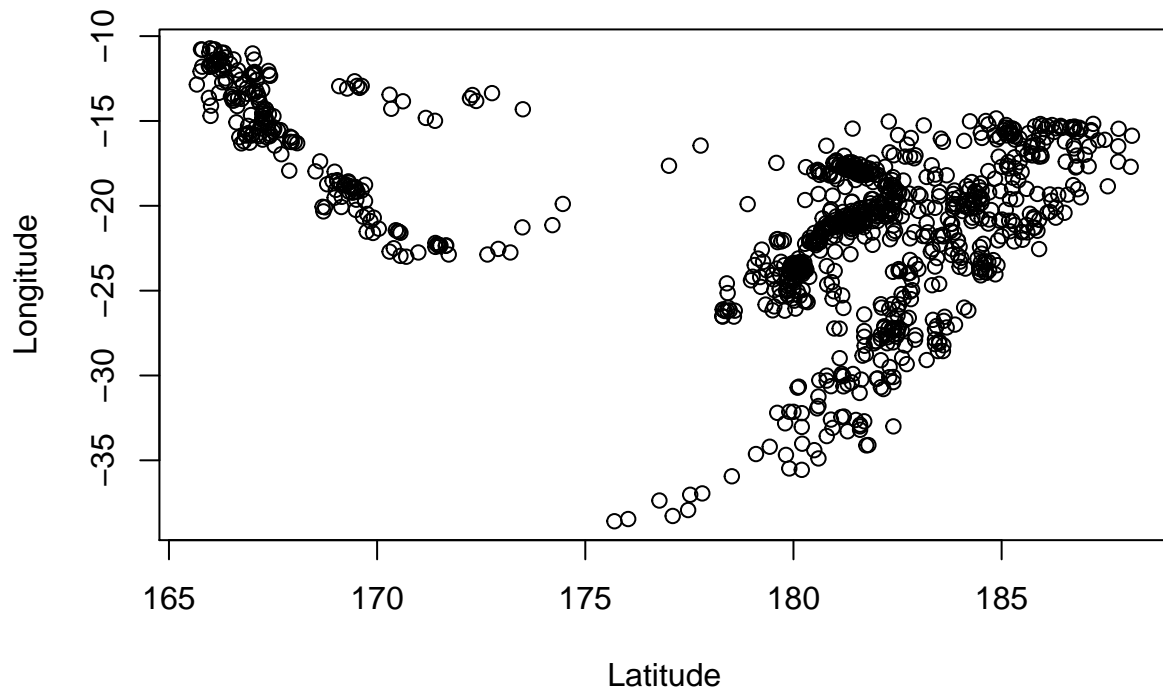
## Scatterplot Example



```
detach(mtcars)
```

```
plot(quakes$long, quakes$lat, xlab="Latitude", ylab="Longitude",  
     main="Location of Earthquake Epicenters")
```

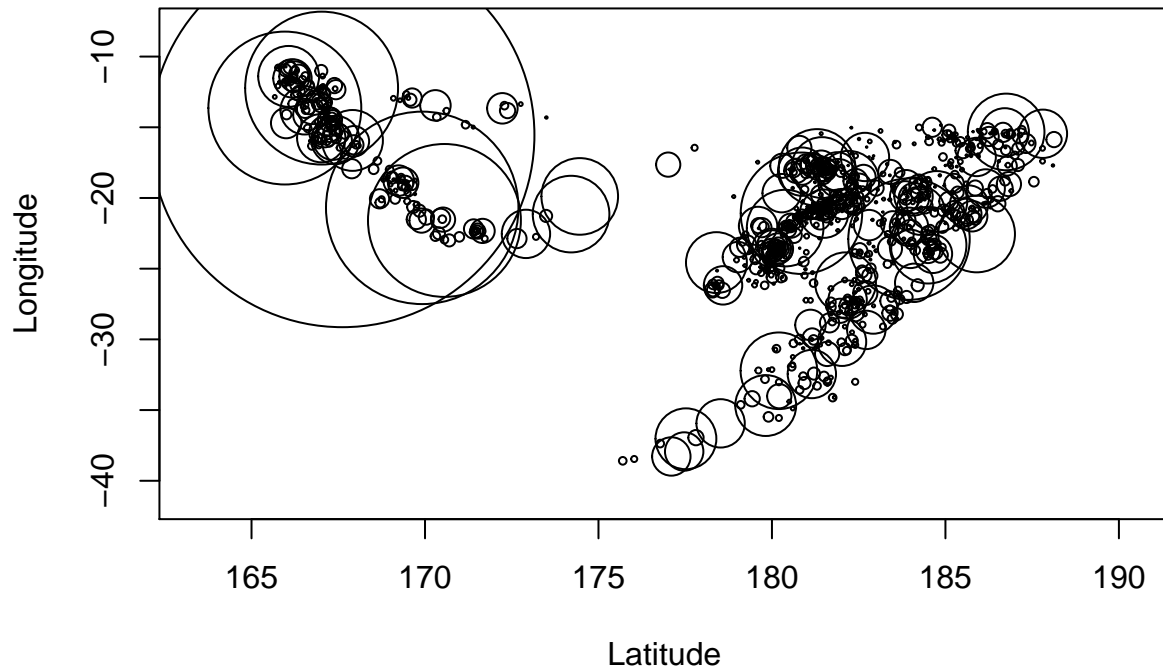
## Location of Earthquake Epicenters



Scatterplots: `plot(x, y)`

```
symbols(quakes$long, quakes$lat, circles = 10 ^ quakes$mag,  
        xlab="Latitude", ylab="Longitude",  
        main="Location of Earthquake Epicenters")
```

## Location of Earthquake Epicenters



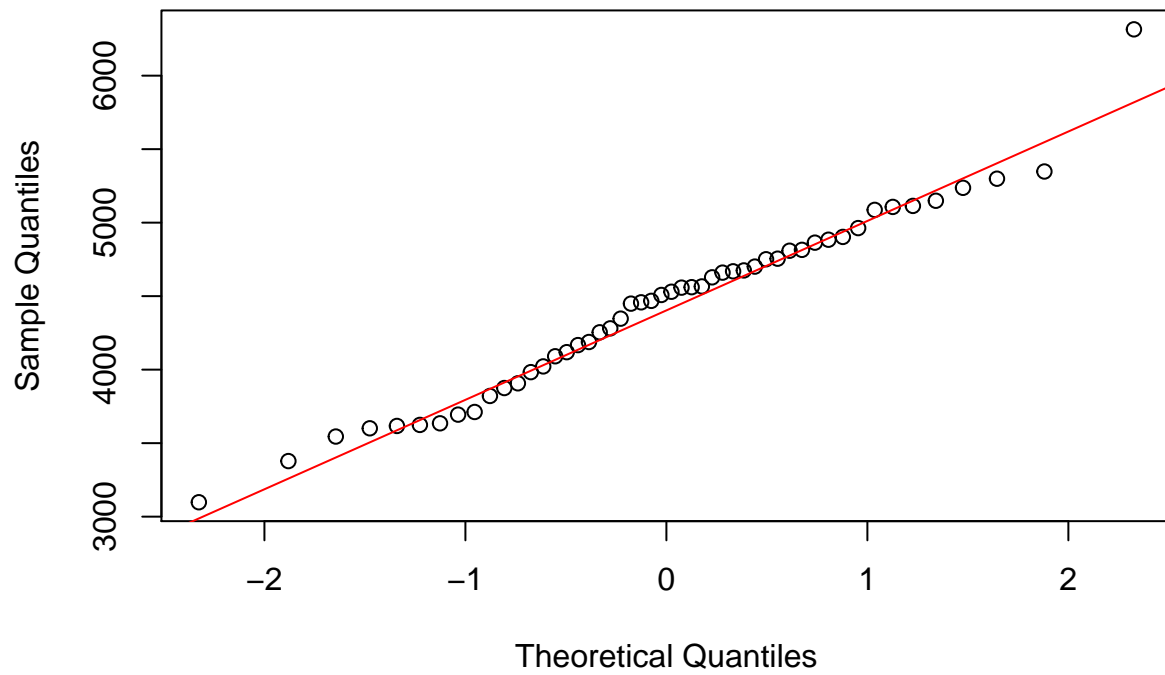
### qqnorm() and qqplot()

- qqnorm() plots the quantiles of a data set against the quantiles of a Normal distribution
- qqplot() plots the quantiles of a first data set against the quantiles of a second data set

### qqnorm() and qqplot()

```
qqnorm(x)                # qq plot for the earthquake depths
qqline(x, col = "red")    # red reference line
```

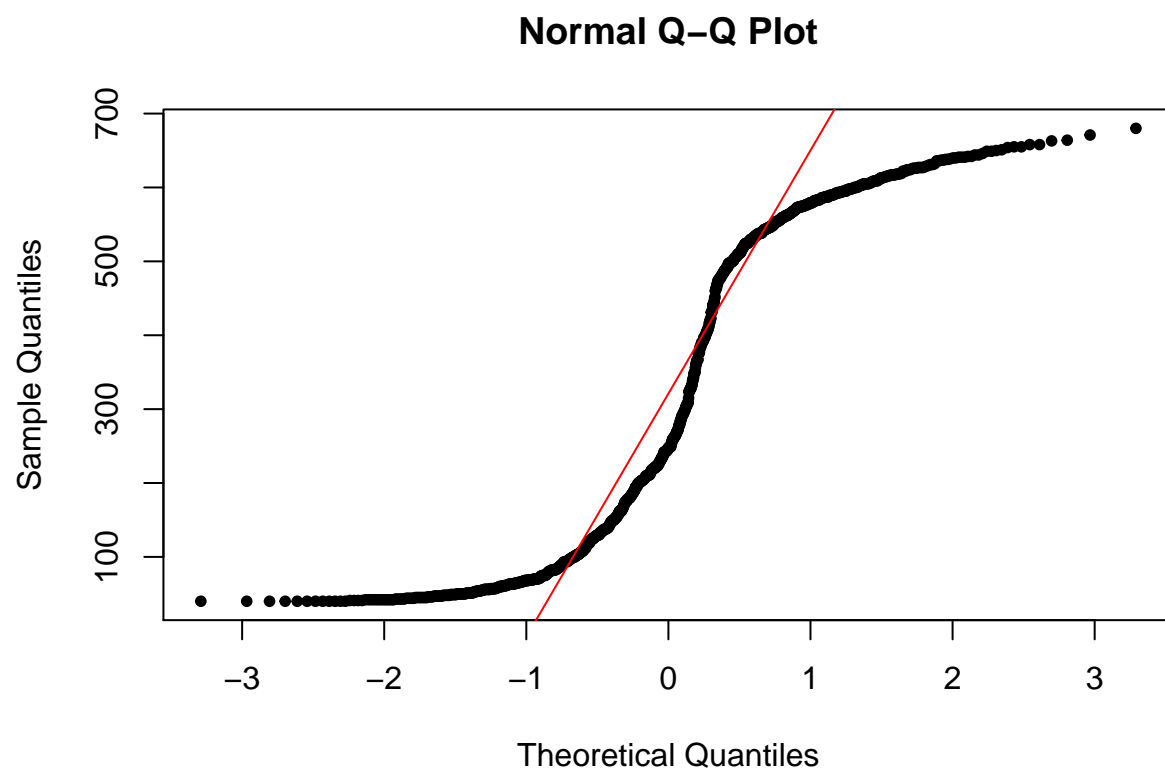
## Normal Q-Q Plot



`qqnorm()` and `qqplot()`

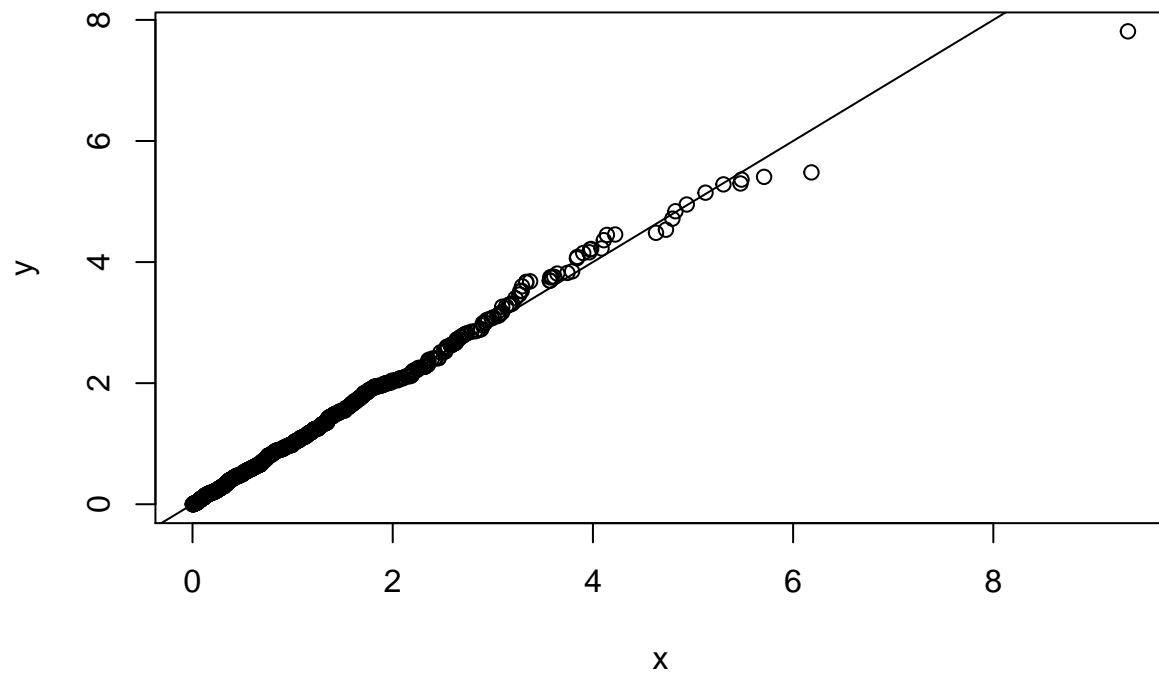
```
qqnorm(y, pch=20)           # qq plot for the earthquake depths
qqline(y, col = "red")      # red reference line
```





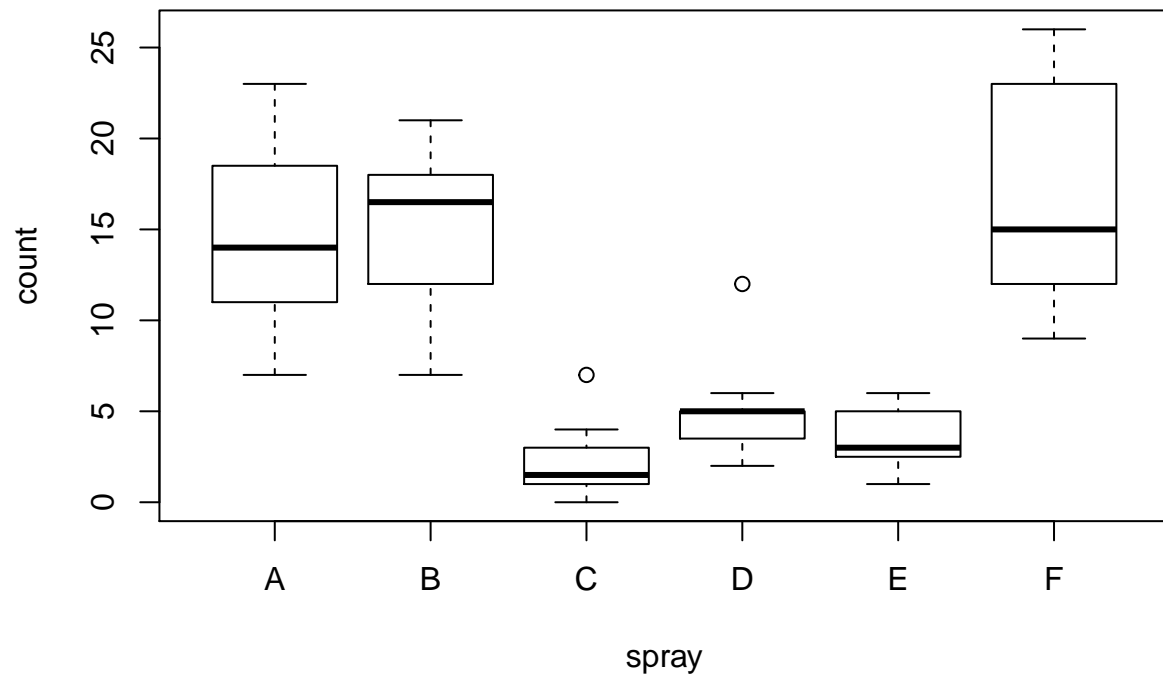
`qqnorm()` and `qqplot()`

```
x <- rexp(1000)
y <- rexp(1000)
qqplot(x,y)
abline(a=0,b=1)
```



## Box plots

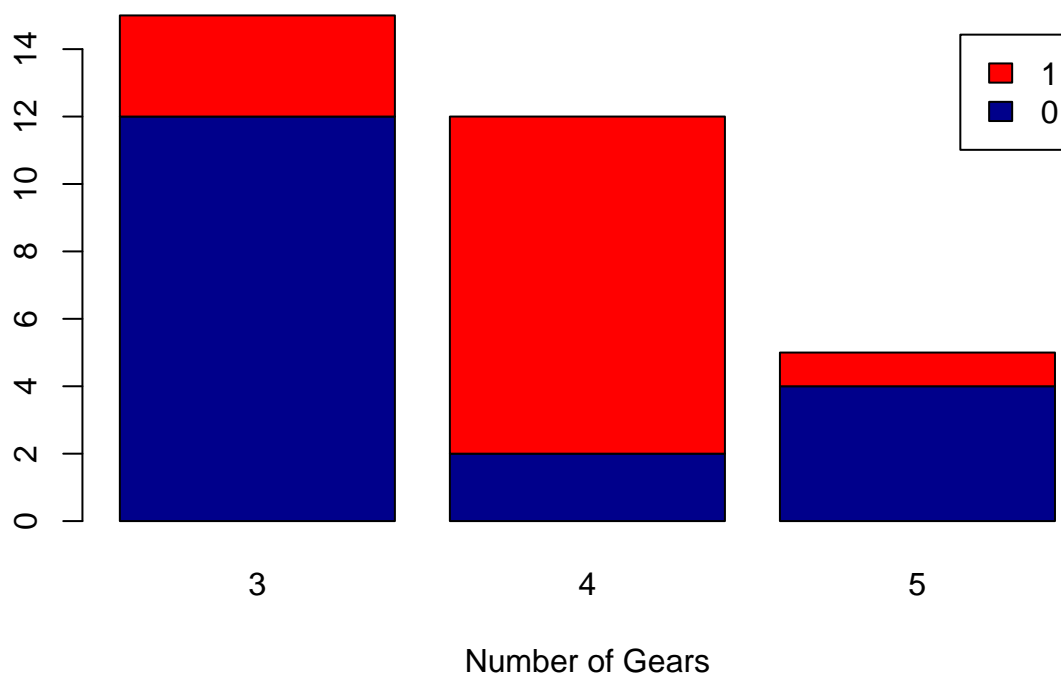
```
boxplot(count ~ spray, data = InsectSprays)
```



### Stacked Bar plot

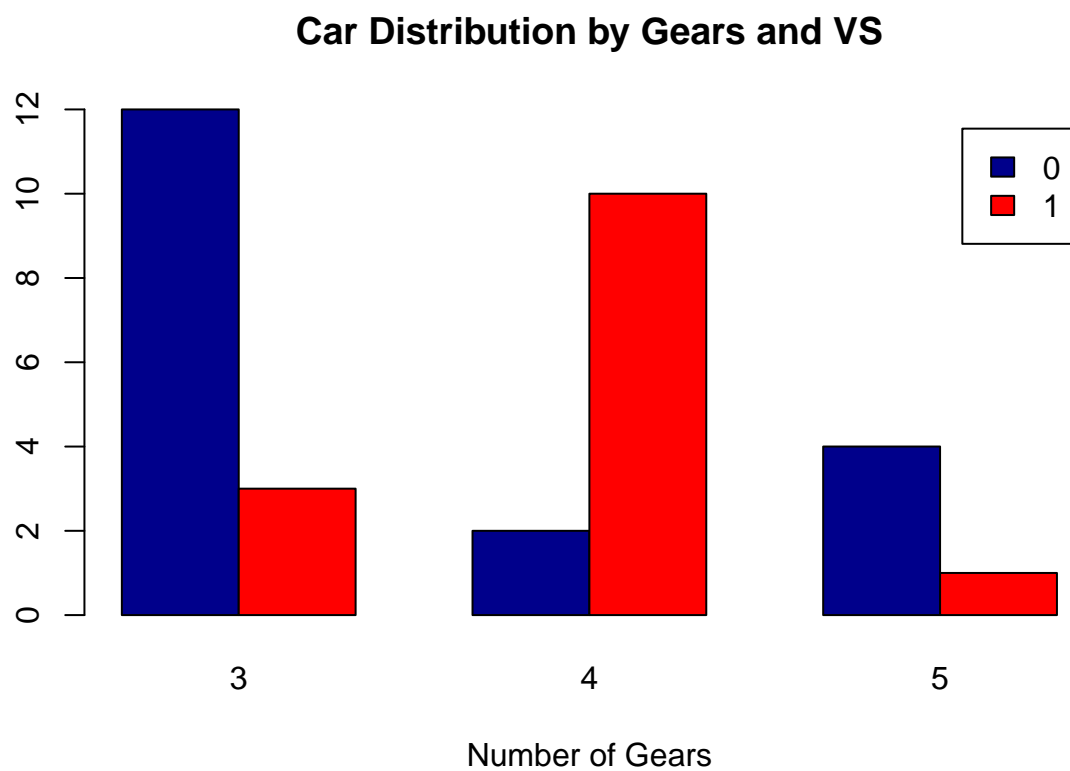
```
counts <- table(mtcars$vs, mtcars$gear)
barplot(counts, main="Car Distribution by Gears and VS",
        xlab="Number of Gears", col=c("darkblue","red"),
        legend = rownames(counts))
```

## Car Distribution by Gears and VS



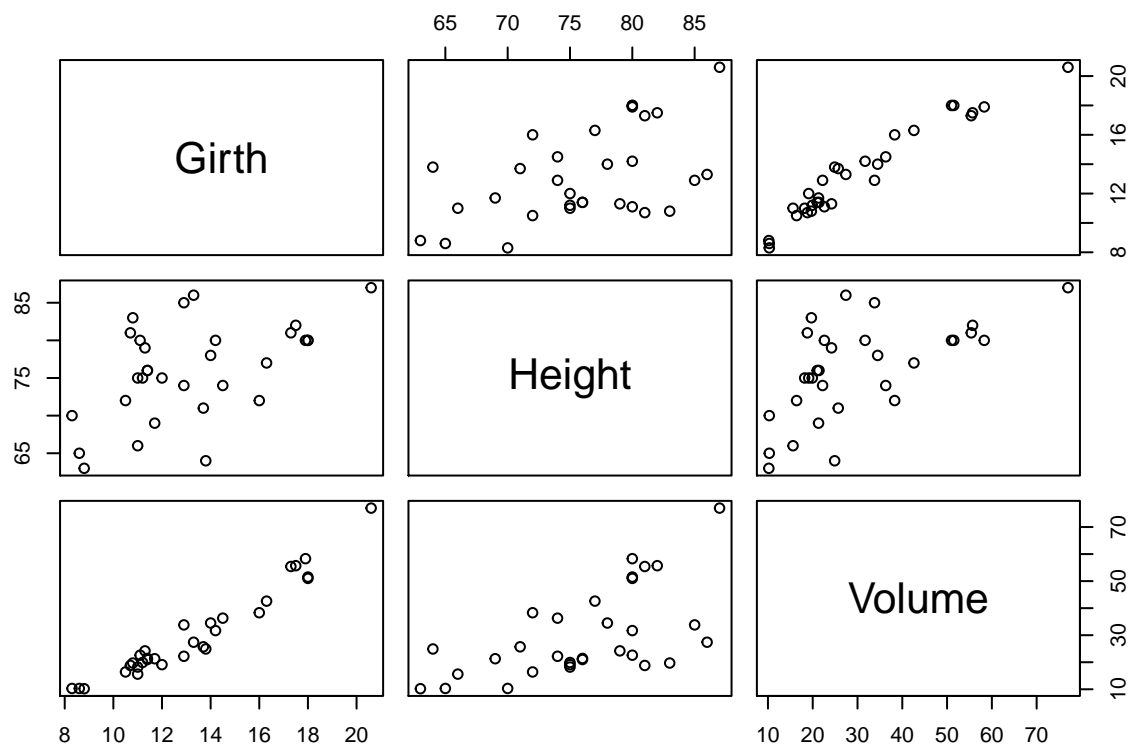
### Grouped Bar Plot

```
counts <- table(mtcars$vs, mtcars$gear)
barplot(counts, main="Car Distribution by Gears and VS",
        xlab="Number of Gears", col=c("darkblue","red"),
        legend = rownames(counts), beside=TRUE)
```



Three-dimensional data: `pairs(x)`

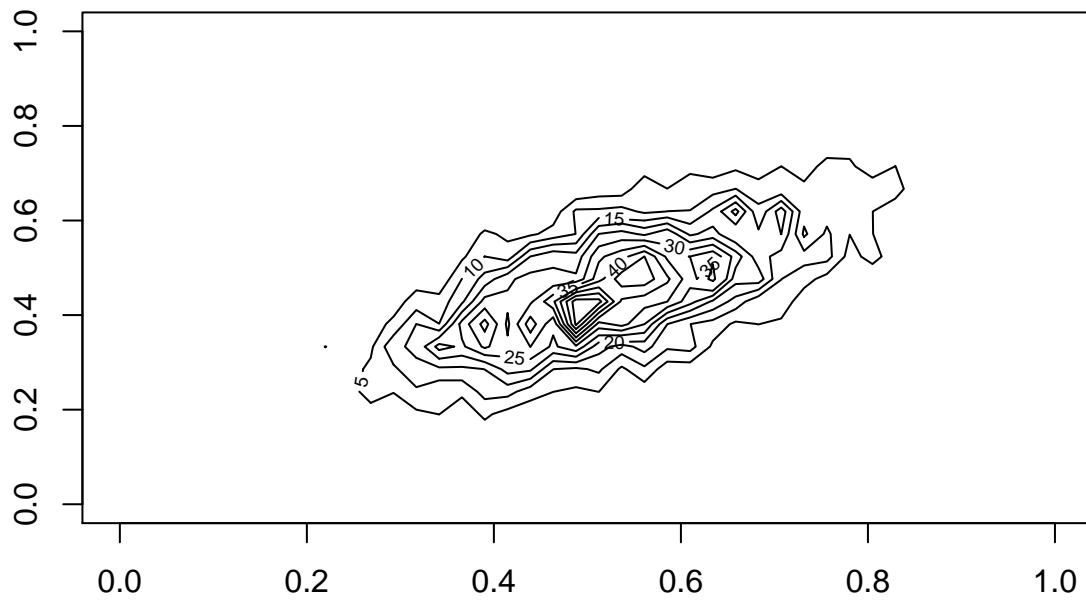
```
pairs(trees)
```



Three dimensional plots: `contour()`

```
contour(crimtab, main="Contour Plot of Criminal Data")
```

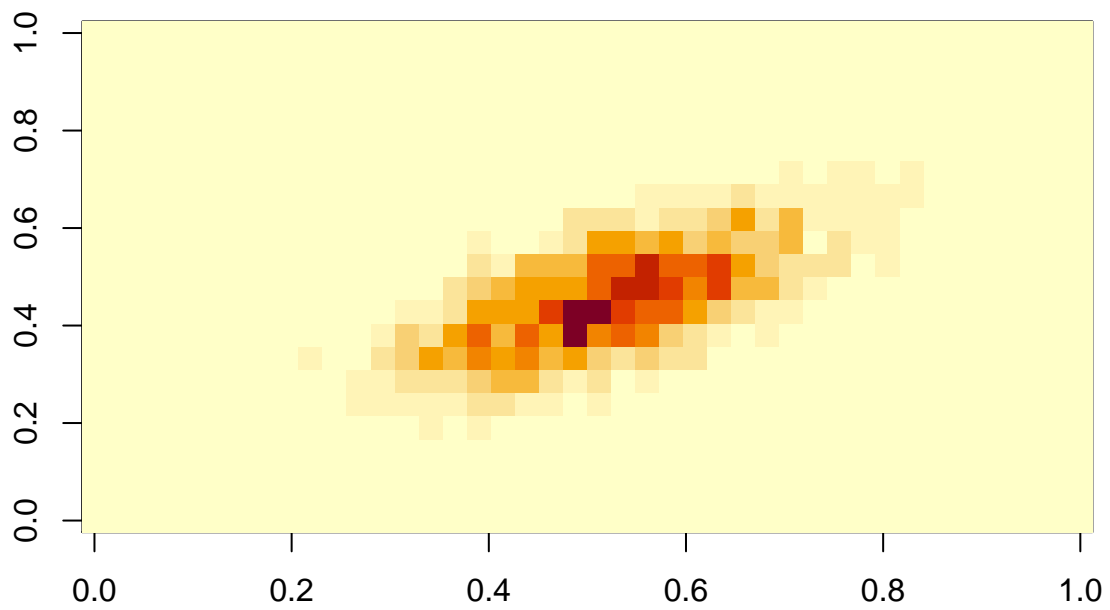
## Contour Plot of Criminal Data



Three dimensional plots: `image()`

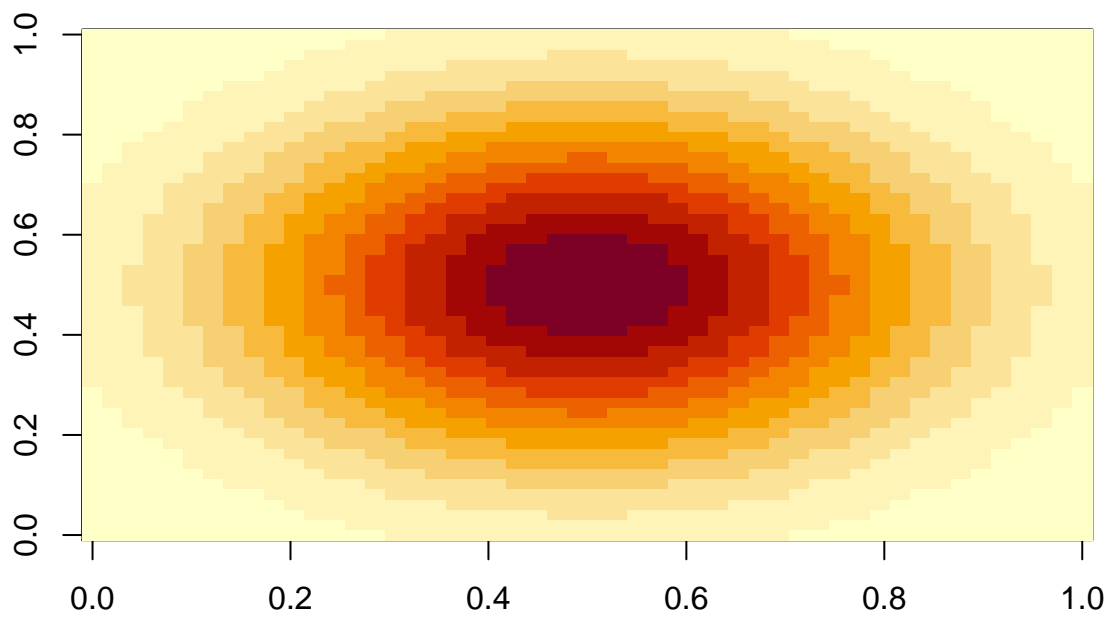
```
image(crimtab, main="Image Plot of Criminal Data")
```

## Image Plot of Criminal Data

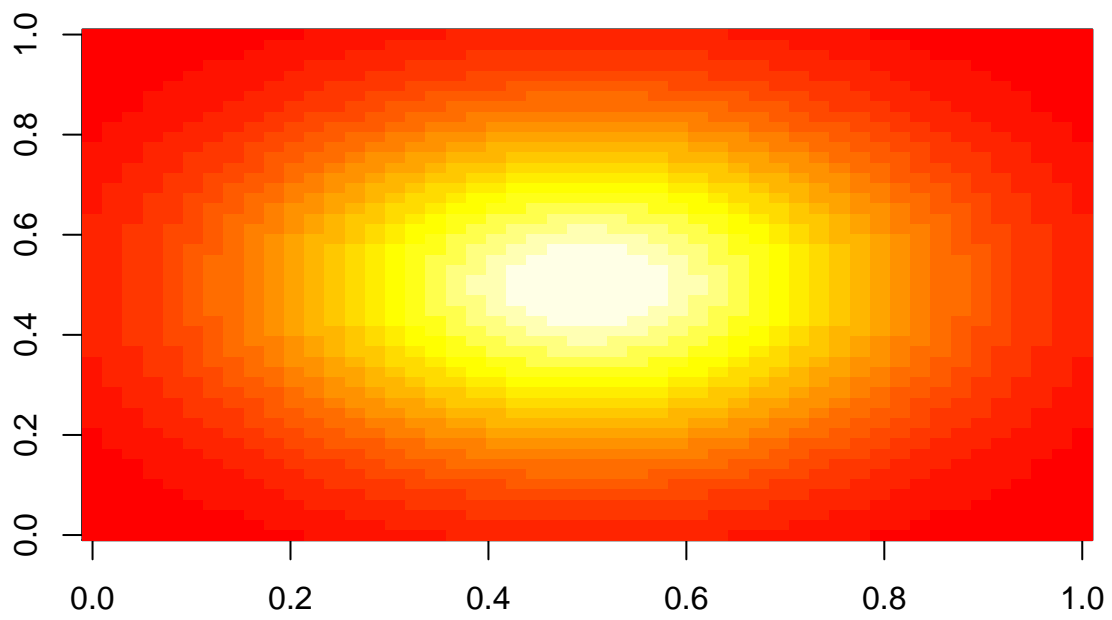


```
phi <- dnorm(seq(-2,2,length=50))  
normal.mat <- phi %o% phi  
image(normal.mat)
```

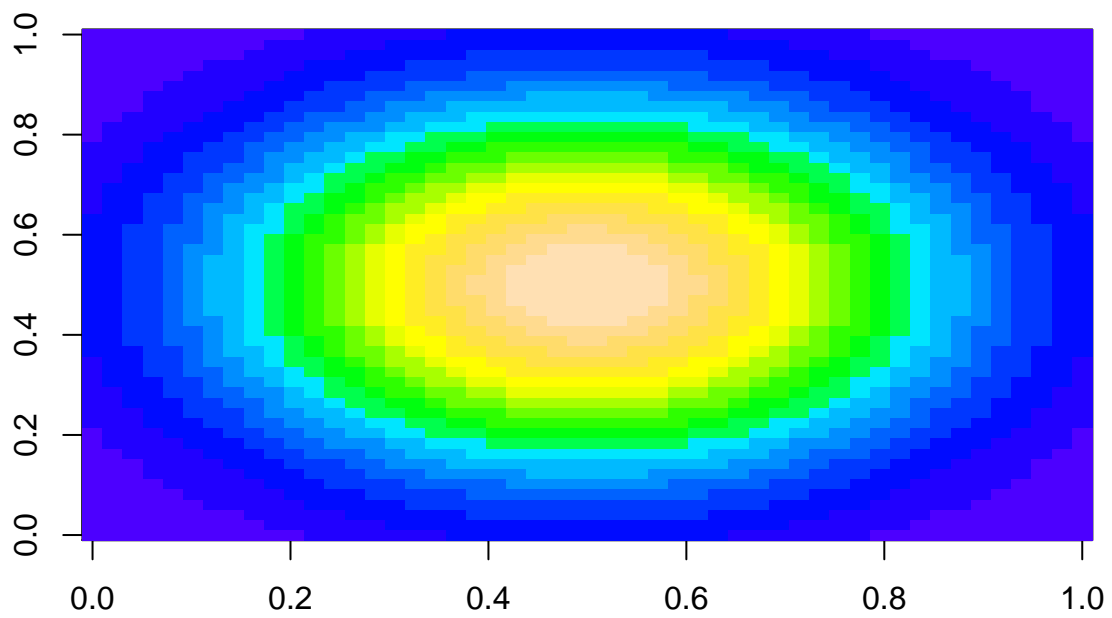




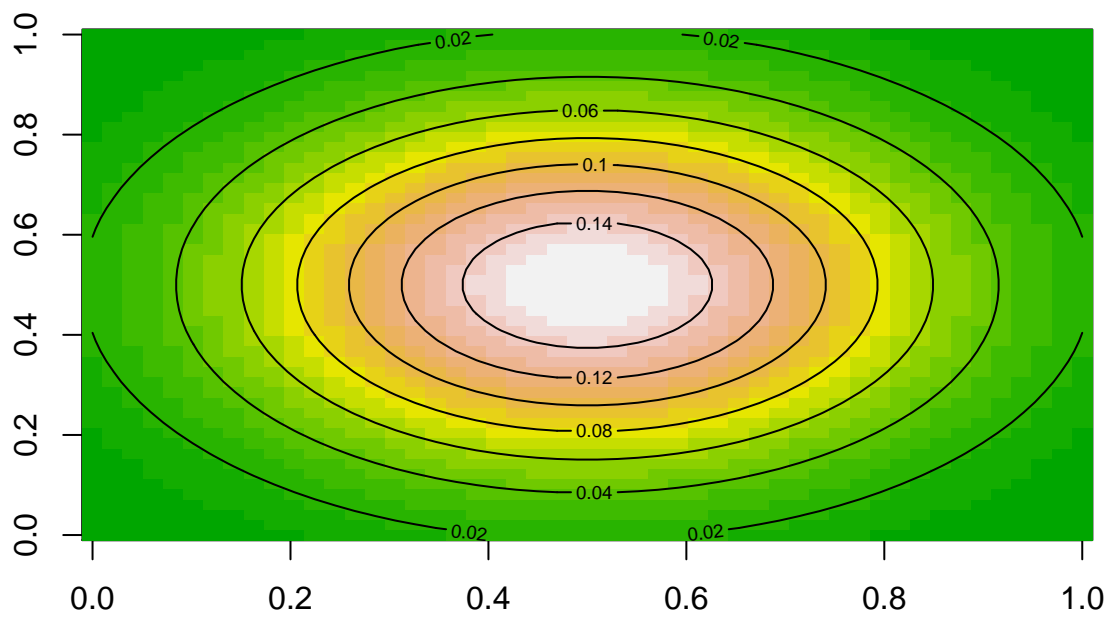
```
image(normal.mat, col=heat.colors(20))
```



```
image(normal.mat, col=topo.colors(20))
```



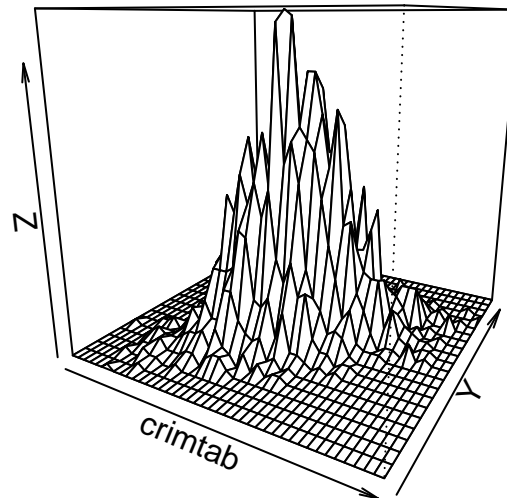
```
image(normal.mat, col=terrain.colors(20))  
contour(normal.mat, add = TRUE)
```



### Three dimensional plots

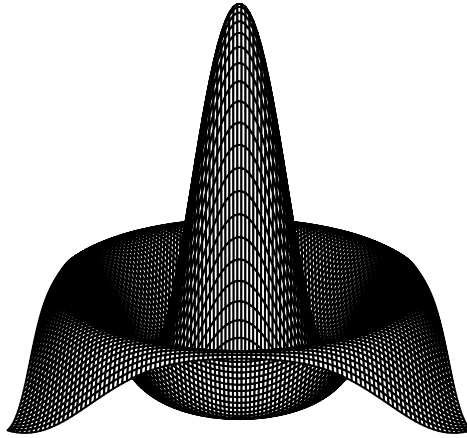
```
persp(crimtab, theta=30, main="Perspective Plot of Criminal Data")
```

## Perspective Plot of Criminal Data



Plot using `persp()` for wire mesh

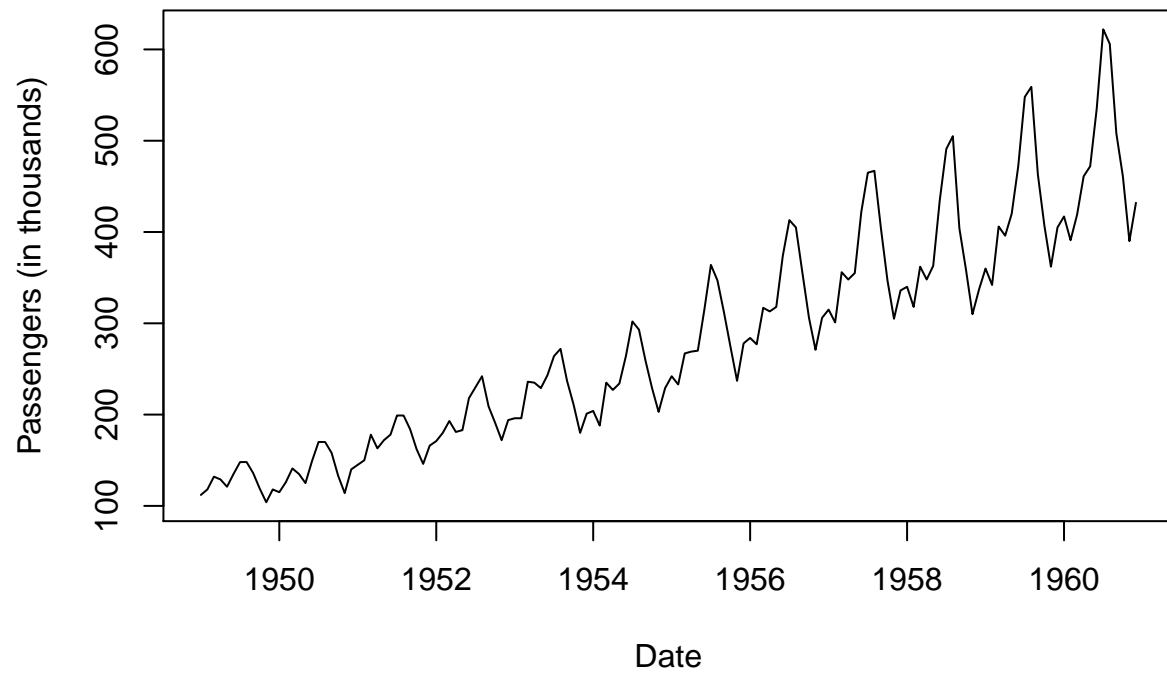
```
x <- seq(-8, 8, length = 100)
y <- x
f <- function(x, y) sin(sqrt(x ^ 2 + y ^ 2)) / (sqrt(x ^ 2 + y ^ 2))
z <- outer(x, y, f)
persp(x, y, z, xlab = "", ylab = "", zlab = "", axes = F, box = F)
```



## Time series plots

```
ts.plot(AirPassengers, xlab="Date", ylab="Passengers (in thousands)",  
        main="International Airline Passengers")
```

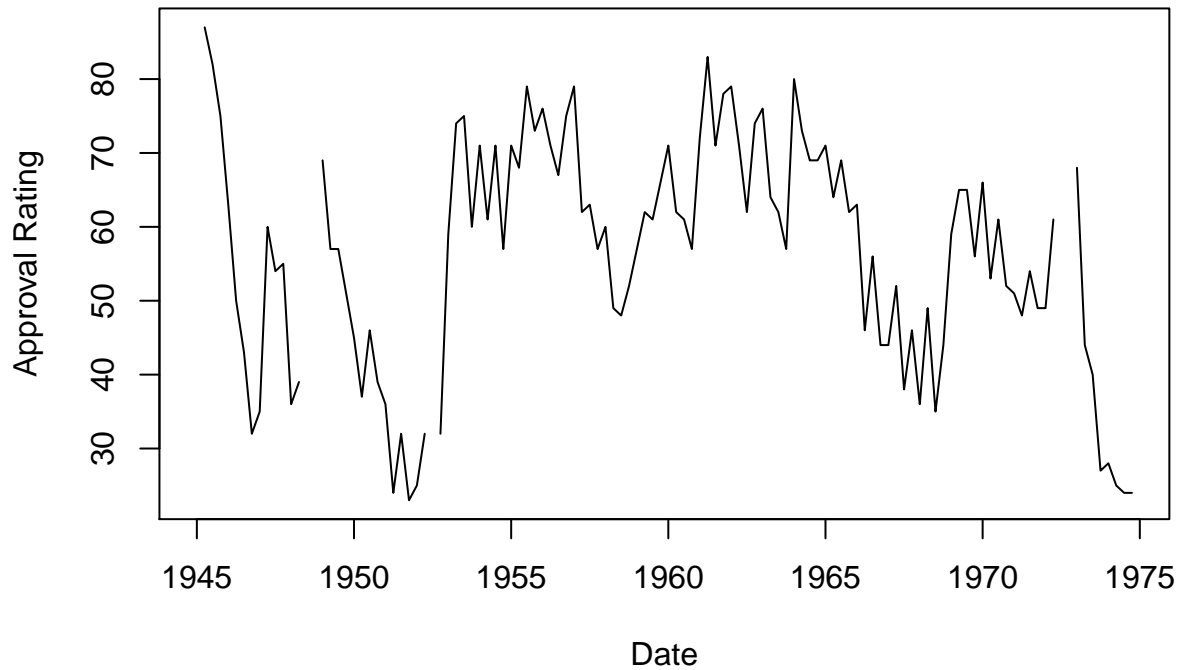
## International Airline Passengers



### Time series plots

```
ts.plot(presidents, xlab="Date", ylab="Approval Rating",  
        main="Presidential Approval Ratings")
```

## Presidential Approval Ratings

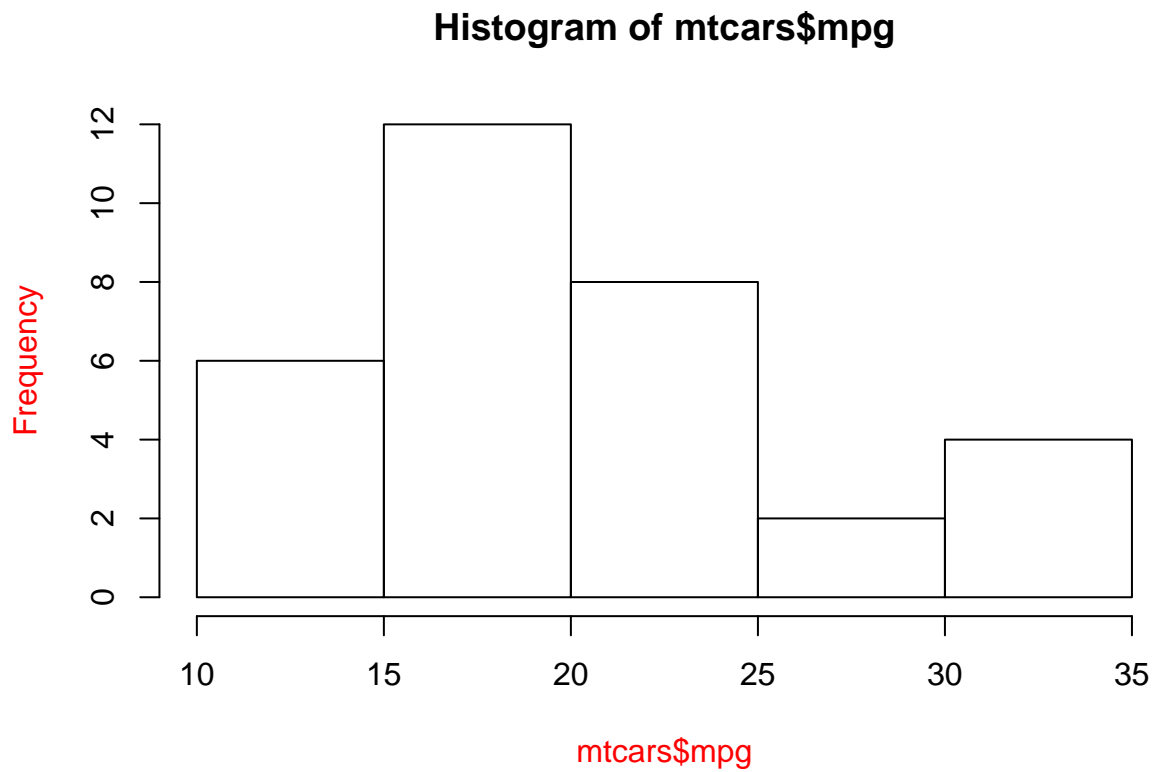


### Custom graphics

- `par()` can be used to set or query graphical parameters (fonts, colors, axes, titles)
- We've already used some of these
- `col`, `col.axis`, `col.lab`, ...: color specification
- `lty`: line type, e.g. dashed, dotted, solid (default), longdash, ...
- `lwd`: line width (helpful to increase for presentation plots)
- `pch`: point types
- ...

```
opar <- par()
par(col.lab="red")
hist(mtcars$mpg)
```





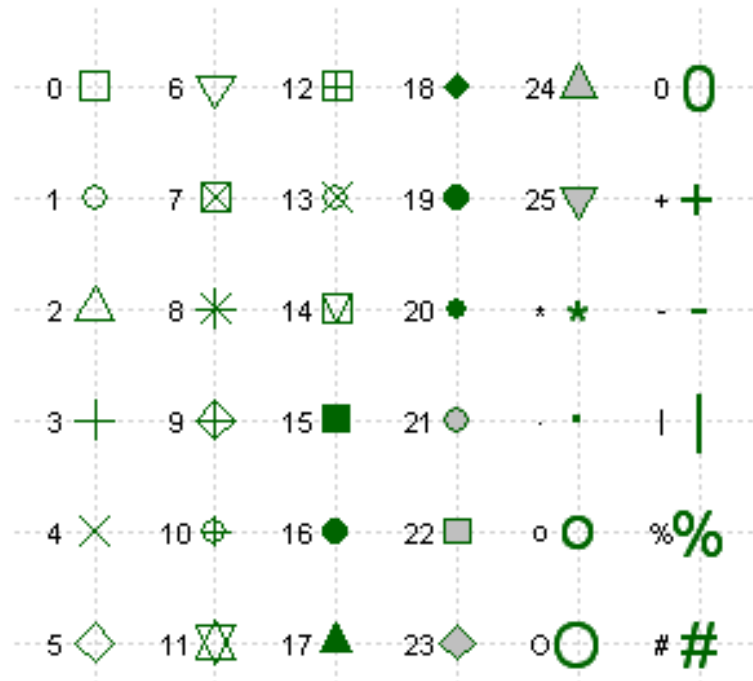
### Text and symbol size

- `cex`: number indicating the amount by which plotting text and symbols should be scaled relative to the default.
- 1=default,
- 1.5 is 50% larger,
- 0.5 is 60% smaller,
- ...
- `cex.axis`
- `cex.lab`
- `cex.main`

### Plotting symbols

- `pch`

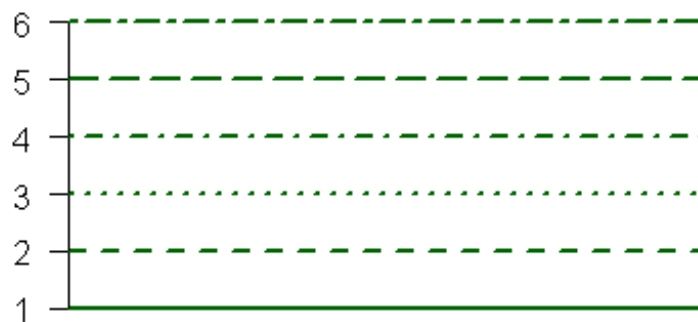
### plot symbols : pch =



### Line type

- lty

### Line Types: lty=

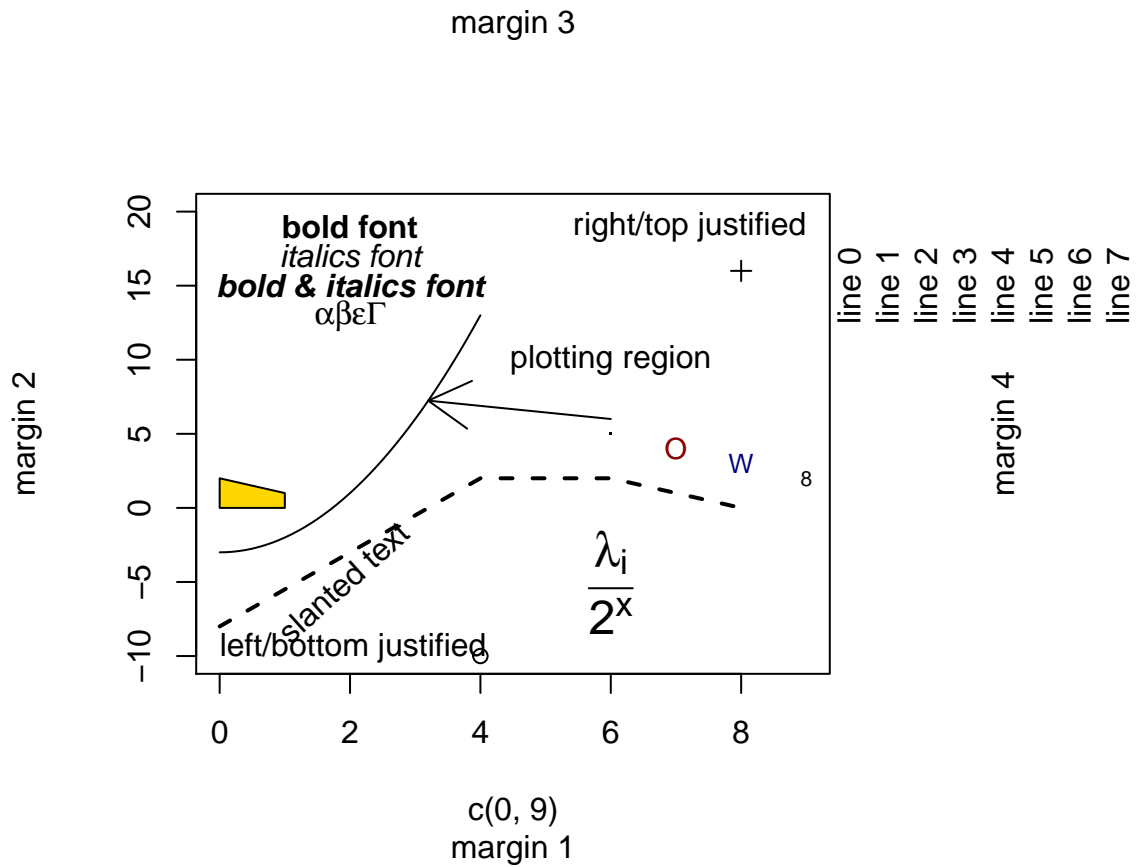


### Fonts

- font, font.axis, font.lab, font.main
- 1 = plain
- 2 = bold
- 3 = italic
- 4 = bold italic
- 5 = symbol
- ps: font point size

- family: font family such as “serif”, “sans”.

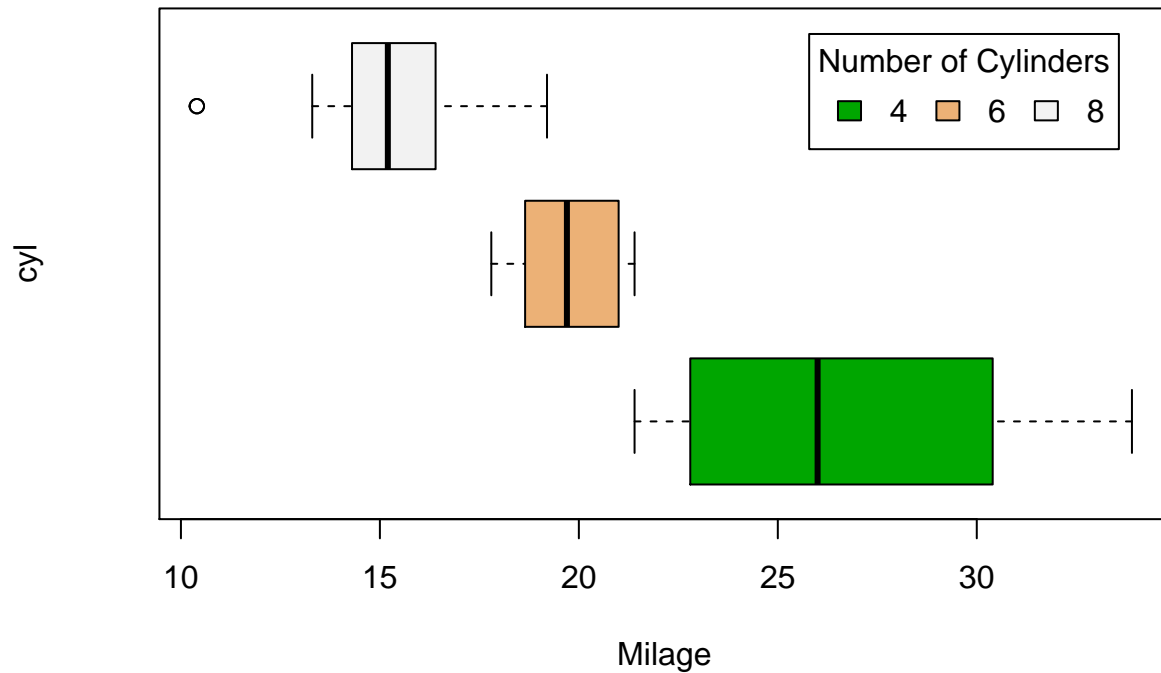
Custom plot:



Add legend

```
boxplot(mpg~cyl, data = mtcars, main="Milage by Car Weight",
  yaxt="n", xlab="Milage", horizontal=TRUE,
  col=terrain.colors(3))
legend("topright", inset=.05, title="Number of Cylinders",
  c("4","6","8"), fill=terrain.colors(3), horiz=TRUE)
```

## Milage by Car Weight



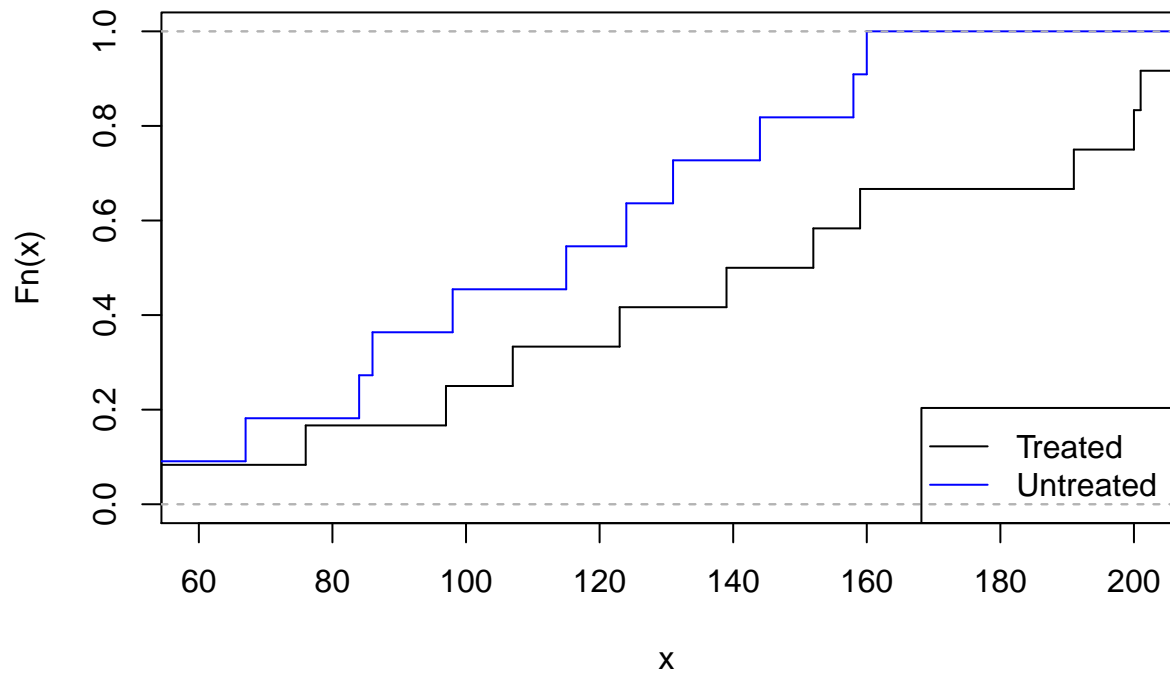
## Multiple plots: Puromycin dataset

```
x <- Puromycin$rate[Puromycin$state == "treated"]
y <- Puromycin$rate[Puromycin$state == "untreated"]
```

## Multiple plots: Puromycin dataset

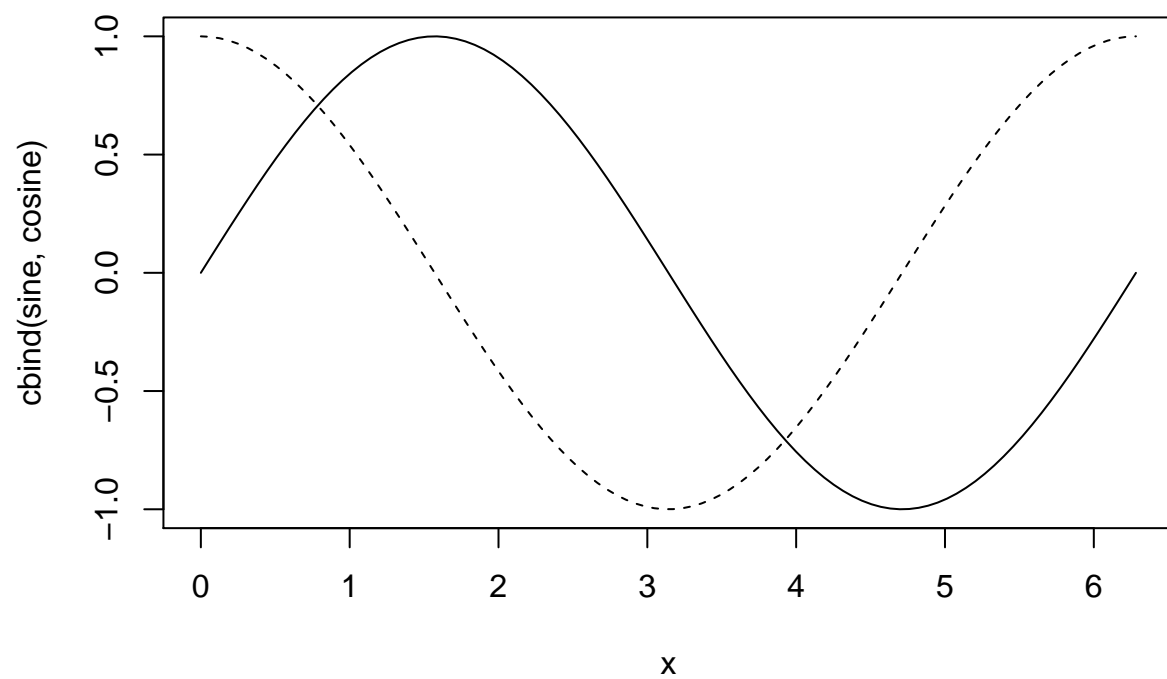
```
plot.ecdf(x, verticals = TRUE, pch = "", xlim = c(60, 200), main="Treated versus Untreated")
lines(ecdf(y), verticals = TRUE, pch = "", xlim = c(60, 200), col="blue")
legend("bottomright", c("Treated", "Untreated"), pch = "", col=c("black", "blue"), lwd = 1)
```

## Treated versus Untreated



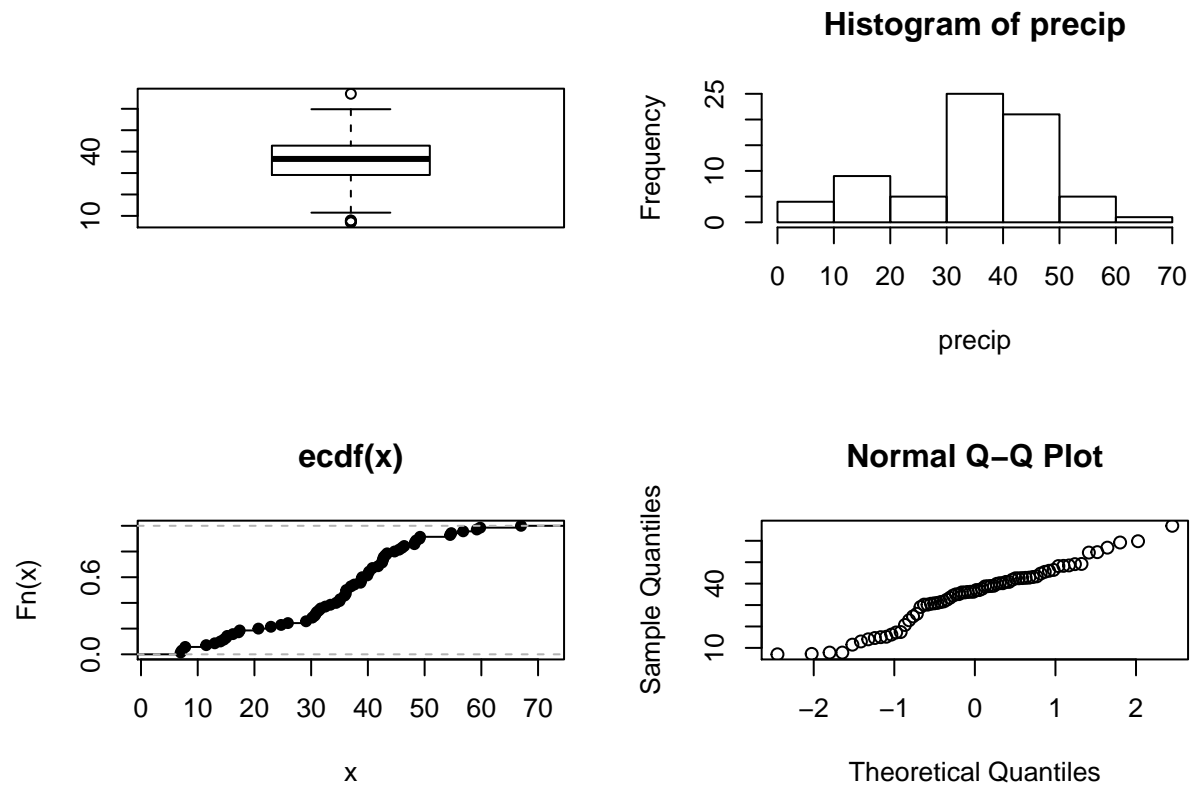
## Multiple plots on one set of axes

```
x <- seq(0, 2 * pi, length = 100)
sine <- sin(x)
cosine <- cos(x)
matplot(x, cbind(sine, cosine), col = c(1, 1), type = "l")
```



### Multiple frame plots: mfrow

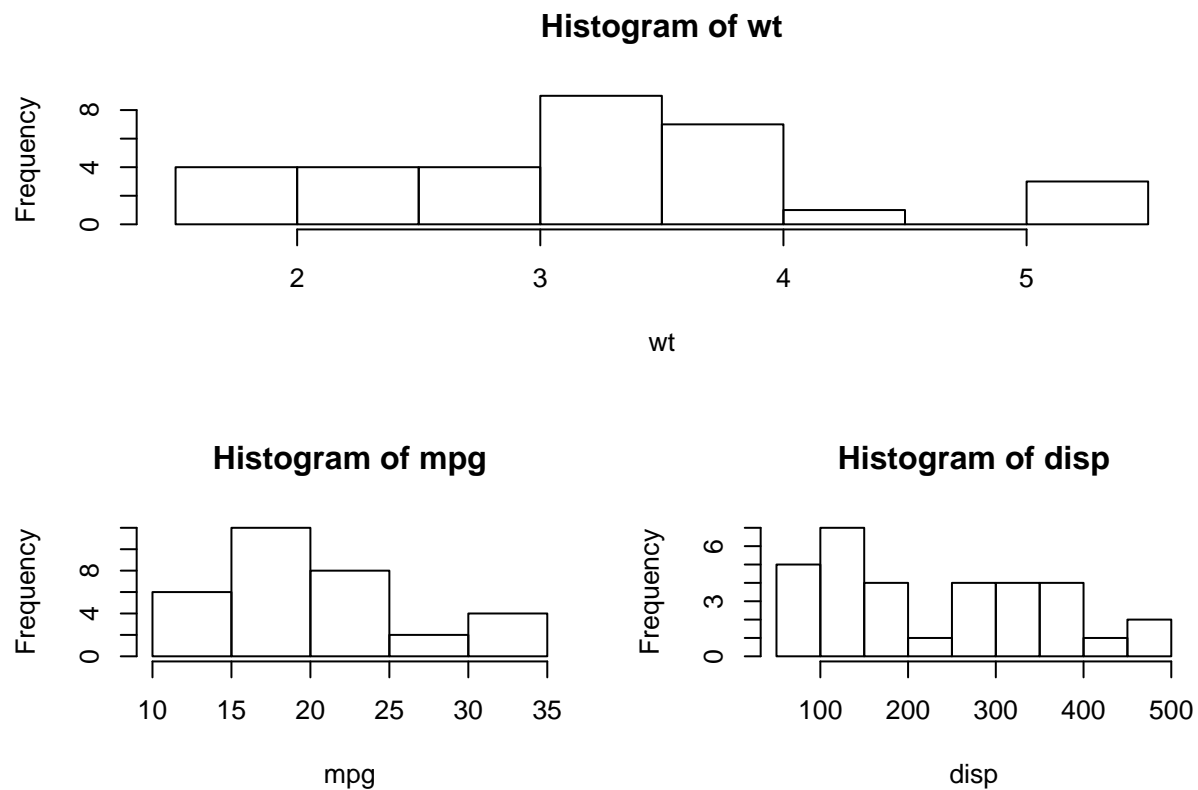
```
par(mfrow = c(2, 2))  
boxplot(precip)  
hist(precip)  
plot.ecdf(precip)  
qqnorm(precip)
```



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

Multiple frame plots: `layout()`

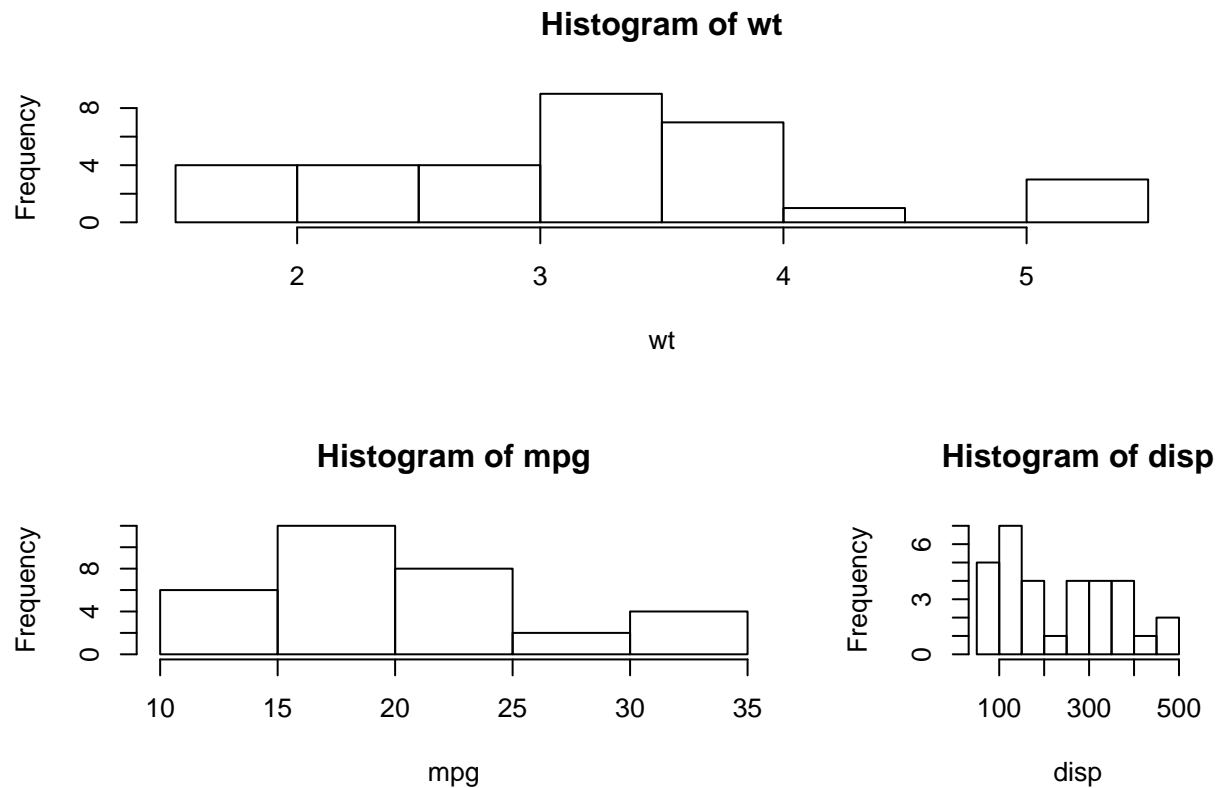
```
attach(mtcars)
layout(matrix(c(1,1,2,3), 2,2, byrow = TRUE))
hist(wt)
hist(mpg)
hist(displ)
```



```
detach(mtcars)
```

```
attach(mtcars)
layout(matrix(c(1,1,2,3), 2,2, byrow = TRUE), widths = c(2,1), heights = c(1,1))
hist(wt)
hist(mpg)
hist(dis)
```





```
detach(mtcars)
```

## Saving a plot to a file

- Begin with functions `postscript()`, `pdf()`, `tiff()`, `jpeg()`, ...
- ... put all your plotting commands here ...
- Finish with `dev.off()`

```
pdf("2cdfs.pdf", width=6, height=4)
plot.ecdf(x, verticals = TRUE, pch = "", xlim = c(60, 200), main="Treated versus Untreated")
lines(ecdf(y), verticals = TRUE, pch = "", xlim = c(60, 200), col="blue")
legend("bottomright", c("Treated", "Untreated"), pch = "", col=c("black", "blue"), lwd = 1)
dev.off()
```

```
## pdf
## 2
```