## Plotting Tools

Statistical Computing, 36-350

Thursday October 31, 2024

#### Last week: Text manipulation

- Strings are, simply put, sequences of characters bound together
- Text data occurs frequently "in the wild", so you should learn how to deal with it!
- nchar(), substr(): functions for substring extractions and replacements
- strsplit(), paste(): functions for splitting and combining strings
- Reconstitution: take lines of text, combine into one long string, then split to get the words
- table(): function to get word counts, useful way of summarizing text data
- Zipf's law: word frequency tends to be inversely proportional to (a power of) rank

#### Part I

Plot basics

### Plotting in R

Base R has a set of powerful plotting tools. An overview:

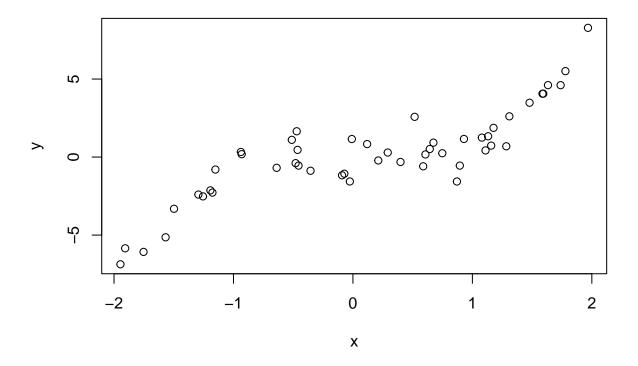
- plot(): generic plotting function
- points(): add points to an existing plot
- lines(), abline(): add lines to an existing plot
- text(), legend(): add text to an existing plot
- rect(), polygon(): add shapes to an existing plot
- hist(), image(): histogram and heatmap
- heat.colors(), topo.colors(), etc: create a color vector
- density(): estimate density, which can be plotted
- contour(): draw contours, or add to existing plot
- curve(): draw a curve, or add to existing plot

The ggplot2 package also provides very nice (and very different) plotting tools; we won't cover it in this course (it tends to be the focus in Statistical Graphics, 36-315)

## Scatter plot

To make a scatter plot of one variable versus another, use plot()

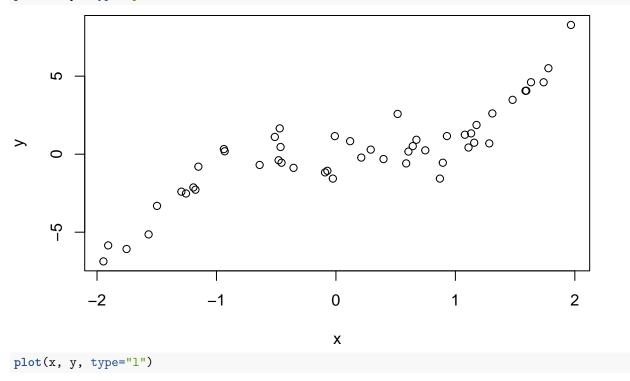
```
n = 50
set.seed(0)
x = sort(runif(n, min=-2, max=2))
y = x^3 + rnorm(n)
plot(x, y)
```

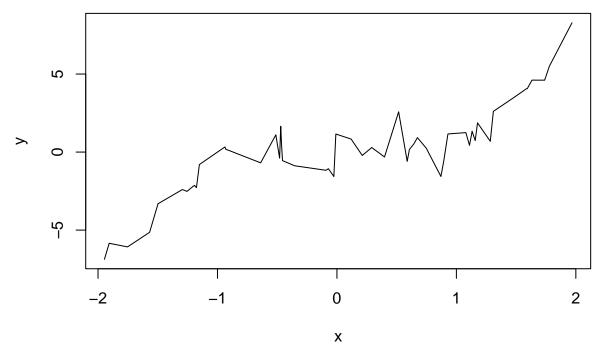


## Plot type

The type argument controls the plot type. Default is p for points; set it to 1 for lines

plot(x, y, type="p")





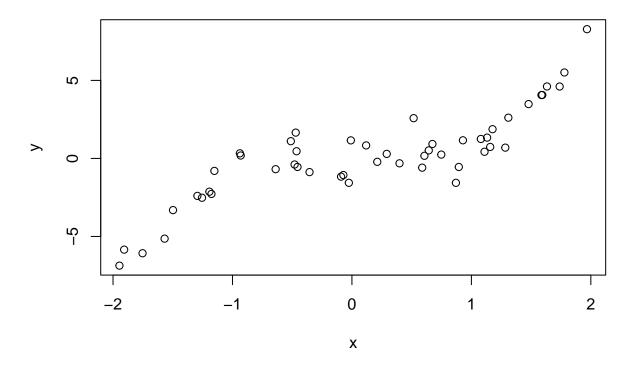
Try also  ${\tt b}$  or  ${\tt o},$  for both points and lines

## Labels

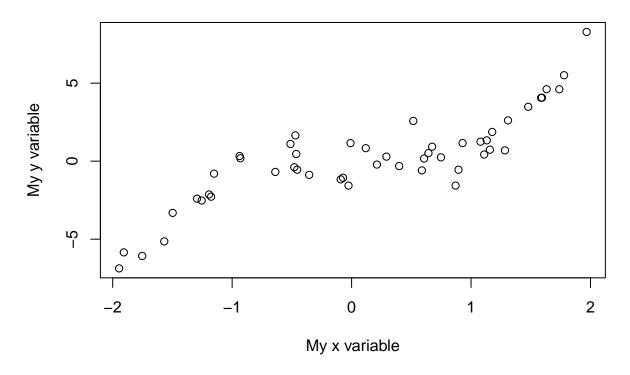
The  $\mathtt{main}$  argument controls the title;  $\mathtt{xlab}$  and  $\mathtt{ylab}$  are the  $\mathtt{x}$  and  $\mathtt{y}$  labels

plot(x, y, main="A noisy cubic") # Note the default x and y labels

# A noisy cubic



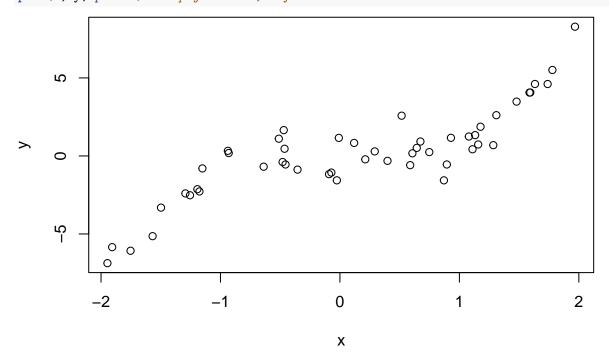
# A noisy cubic



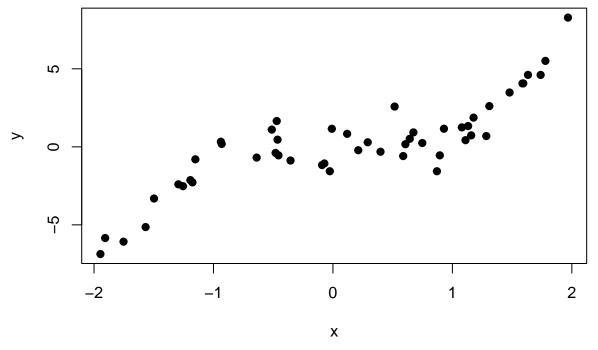
## Point type

Use the pch argument to control point type

plot(x, y, pch=21) # Empty circles, default





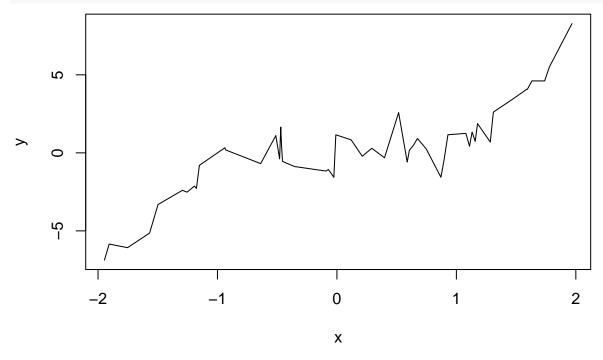


Try also 20 for small filled circles, or  $"\,.\,"$  for single pixels

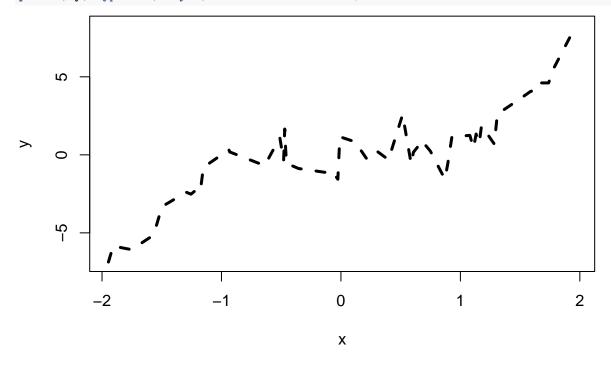
# Line type

Use the lty argument to control the line type, and lwd to control the line width

plot(x, y, type="l", lty=1, lwd=1) # Solid line, default width



plot(x, y, type="1", lty=2, lwd=3) # Dashed line, 3 times as thick



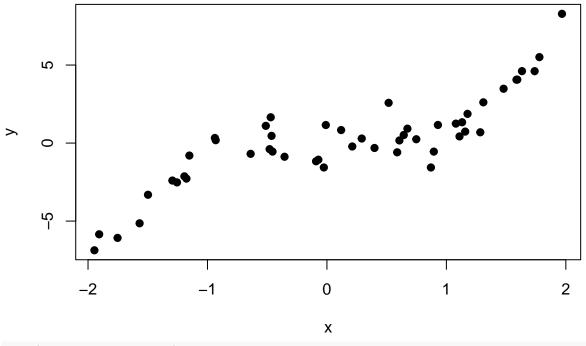
## Color

Use the col argument to control the color. Can be:

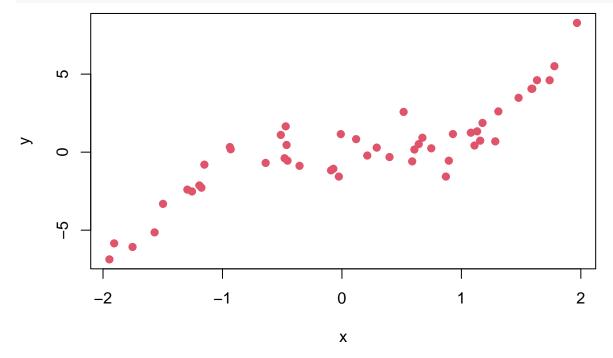
- An integer between 1 and 8 for basic colors
- A string for any of the 657 available named colors

The function colors() returns a string vector of the available colors

plot(x, y, pch=19, col=1) # Black, default



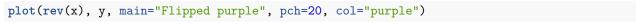


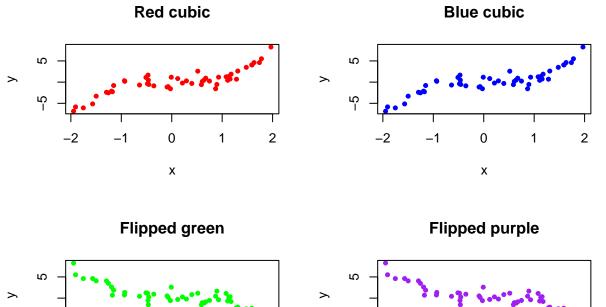


## Multiple plots

To set up a plotting grid of arbitrary dimension, use the par() function, with the argument mfrow. Note: in general this will affect all following plots! (Except in separate R Markdown code chunks ...)

```
par(mfrow=c(2,2)) # Grid elements are filled by row
plot(x, y, main="Red cubic", pch=20, col="red")
plot(x, y, main="Blue cubic", pch=20, col="blue")
plot(rev(x), y, main="Flipped green", pch=20, col="green")
```





## Margin

-5

-2

-1

0

rev(x)

Default margins in R are large (and ugly); to change them, use the par() function, with the argument mar. Note: in general this will affect all following plots! (Except in separate R Markdown code chunks ...)

2

1

-5

-2

-1

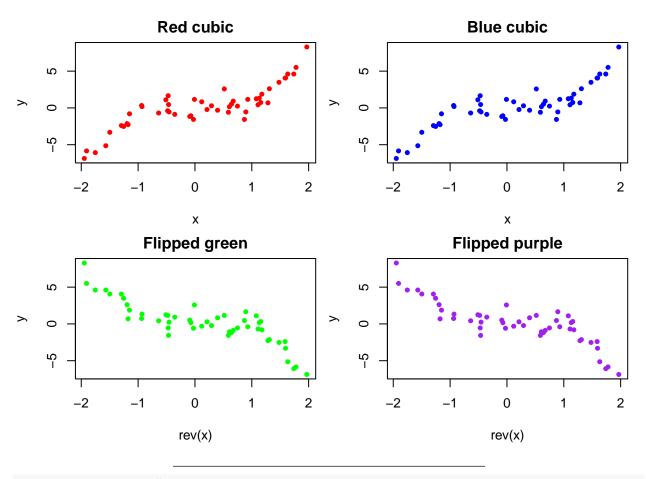
0

rev(x)

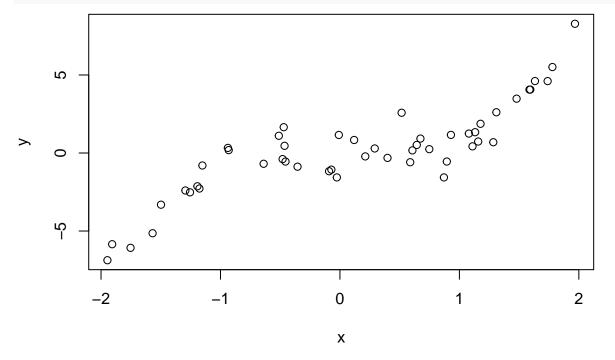
1

2

```
par(mfrow=c(2,2), mar=c(4,4,2,0.5))
plot(x, y, main="Red cubic", pch=20, col="red")
plot(x, y, main="Blue cubic", pch=20, col="blue")
plot(rev(x), y, main="Flipped green", pch=20, col="green")
plot(rev(x), y, main="Flipped purple", pch=20, col="purple")
```



# Evidence that par() does not carry over to separate R Markdown code chunks plot(x, y)



#### Saving plots

Use the pdf() function to save a pdf file of your plot, in your R working directory. Use getwd() to get the working directory, and setwd() to set it

```
getwd() # This is where the pdf will be saved
```

## [1] "/Users/townesf/cmu-gdrive/36-350/lectures/09\_plotting"

```
pdf(file="noisy_cubics.pdf", height=7, width=7) # Height, width are in inches
par(mfrow=c(2,2), mar=c(4,4,2,0.5))
plot(x, y, main="Red cubic", pch=20, col="red")
plot(x, y, main="Blue cubic", pch=20, col="blue")
plot(rev(x), y, main="Flipped green", pch=20, col="green")
plot(rev(x), y, main="Flipped purple", pch=20, col="purple")
graphics.off()
```

Also, use the jpg() and png() functions to save jpg and png files

#### Adding to plots

The main tools for this are:

- points(): add points to an existing plot
- lines(), abline(): add lines to an existing plot
- text(), legend(): add text to an existing plot
- rect(), polygon(): add shapes to an existing plot

You'll get practice with this on lab. Pay attention to layers—they work just like they would if you were painting a picture by hand

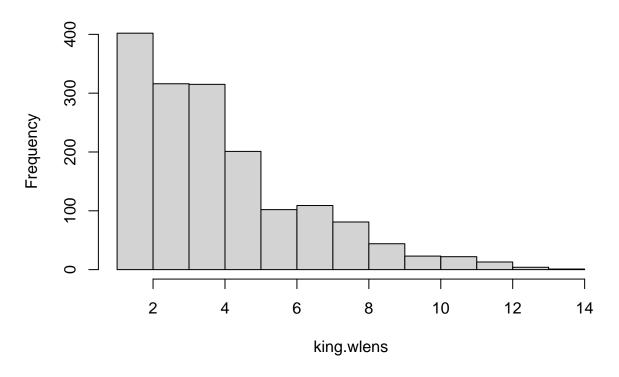
#### Part II

Histograms and heatmaps

## Plotting a histogram

To plot a histogram of a numeric vector, use hist()

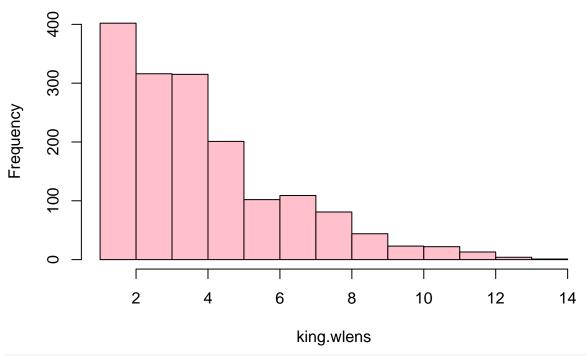
# Histogram of king.wlens



## Histogram options

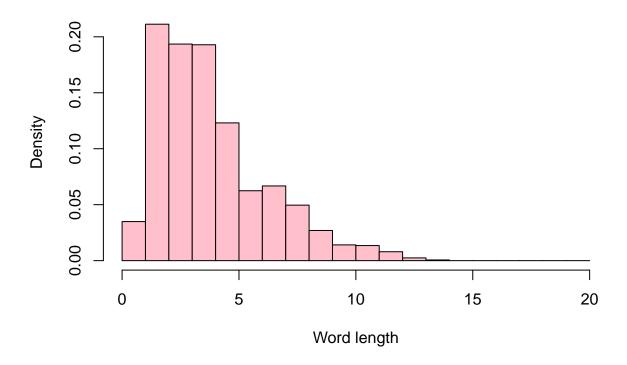
Several options are available as arguments to hist(), such as col, freq, breaks, xlab, ylab, main hist(king.wlens, col="pink", freq=TRUE) # Frequency scale, default

# Histogram of king.wlens



hist(king.wlens, col="pink", freq=FALSE, # Probability scale, and more options breaks=0:20, xlab="Word length", main="King word lengths")

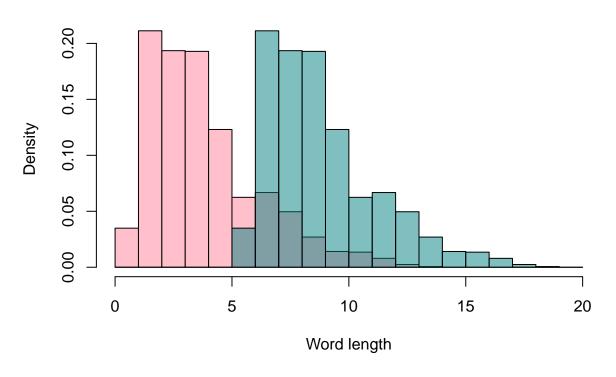
## King word lengths



#### Adding a histogram to an existing plot

To add a histogram to an existing plot (say, another histogram), use hist() with add=TRUE

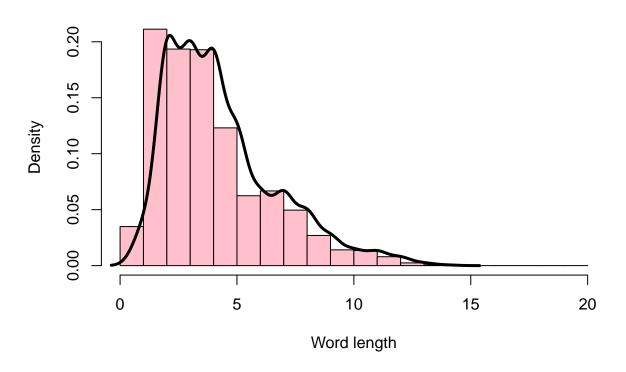
### King word lengths



### Adding a density curve to a histogram

To estimate a density from a numeric vector, use density(). This returns a list; it has components x and y, so we can actually call lines() directly on the returned object

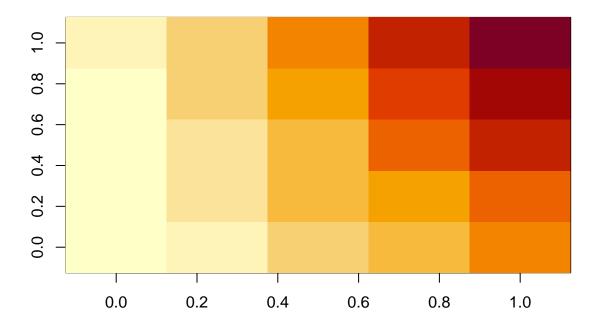
## King word lengths



## Plotting a heatmap

To plot a heatmap of a numeric matrix, use image()

```
(mat = 1:5 %o% 6:10) # %o% gives for outer product
##
        [,1] [,2] [,3] [,4] [,5]
## [1,]
                 7
                            9
           6
                      8
                                10
##
   [2,]
          12
                14
                     16
                           18
                                20
   [3,]
          18
                21
                           27
                                30
##
                     24
##
  [4,]
          24
                28
                     32
                          36
                                40
## [5,]
          30
                35
                     40
                                50
image(mat) # Red means low, white means high
```



### Orientation of image()

The orientation of image() is to plot the heatmap according to the following order, in terms of the matrix elements:

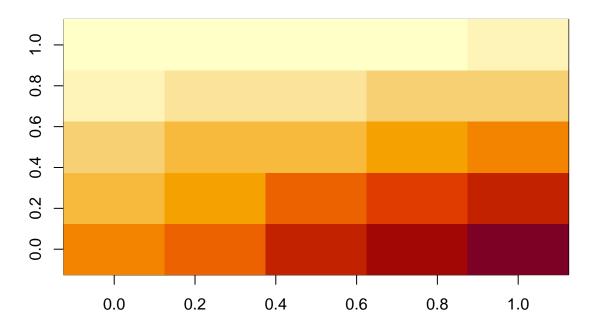
$$\begin{array}{cccc} (1, ncol) & (2, ncol) & \dots & (nrow, ncol) \\ \vdots & & & \\ (1, 2) & (2, 2) & \dots & (nrow, 2) \\ (1, 1) & (2, 1) & \dots & (nrow, 1) \end{array}$$

This is a 90 degrees counterclockwise rotation of the "usual" printed order for a matrix:

$$(1,1)$$
  $(1,2)$  ...  $(1, ncol)$   $(2,1)$   $(2,2)$  ...  $(2, ncol)$   $\vdots$   $(nrow,1)$   $(nrow,2)$  ...  $(nrow,ncol)$ 

Therefore, if you want the displayed heatmap to follow the usual order, you must rotate the matrix 90 degrees clockwise before passing it in to image(). (Equivalently: reverse the row order, then take the transpose.) Convenient way of doing so:

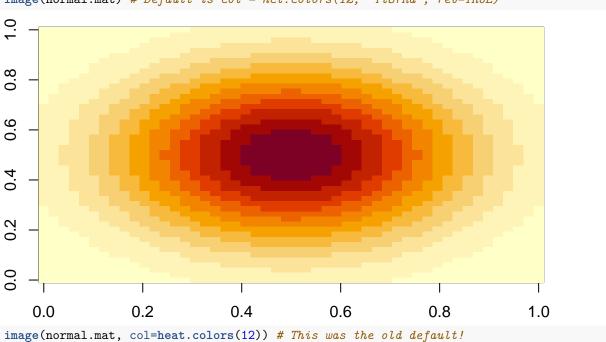
```
clockwise90 = function(a) { t(a[nrow(a):1,]) } # Handy rotate function
image(clockwise90(mat))
```

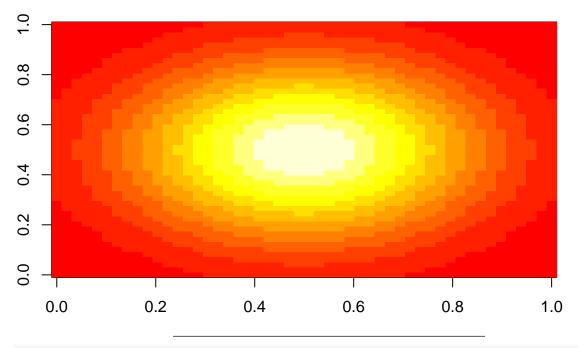


#### Color scale

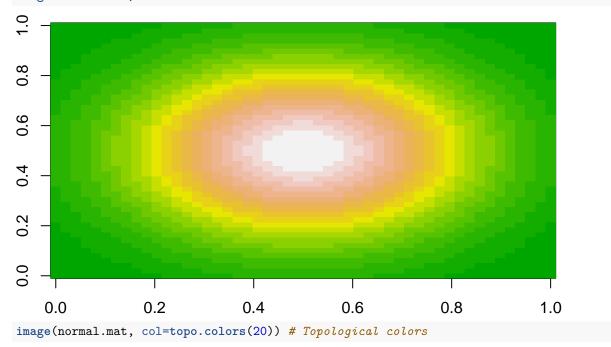
The default is to use a red-to-white color scale in image(). But the col argument can take any vector of colors. Built-in functions gray.colors(), rainbow(), heat.colors(), topo.colors(), terrain.colors(), cm.colors() all return continguous color vectors of given length

```
phi = dnorm(seq(-2,2,length=50))
normal.mat = phi %o% phi
image(normal.mat) # Default is col = hcl.colors(12, "YlOrRd", rev=TRUE)
```

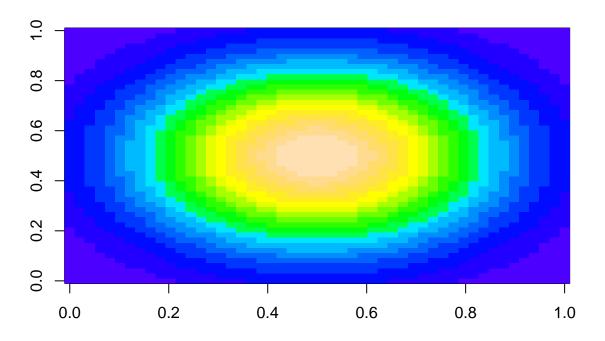




image(normal.mat, col=terrain.colors(20)) # Terrain colors

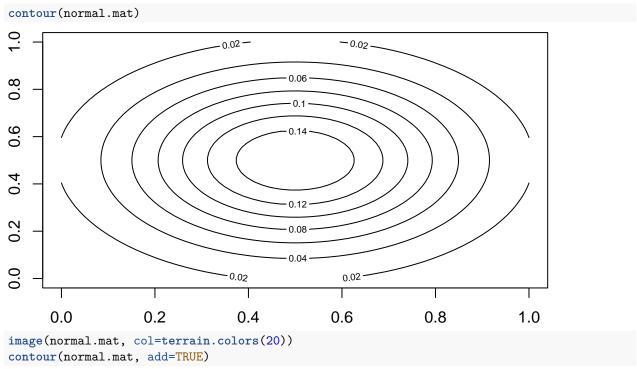


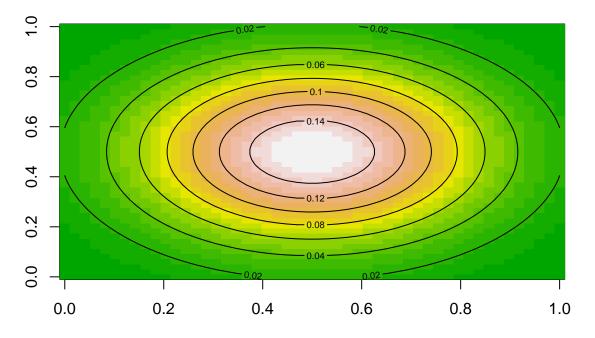
17



## Drawing contour lines

To draw contour lines from a numeric matrix, use contour(); to add contours to an existing plot (like, a heatmap), use contour() with add=TRUE



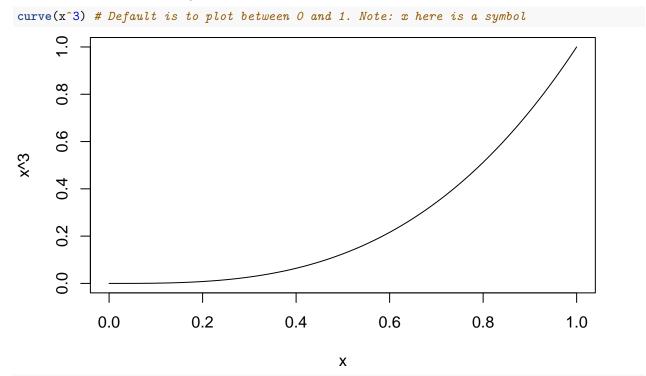


Part III

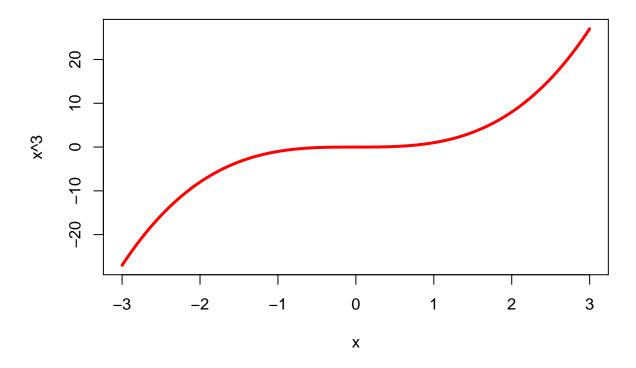
Curves, surfaces, and colors

## Drawing a curve

To draw a curve of a function, use curve()



curve(x^3, from=-3, to=3, lwd=3, col="red") # More plotting options

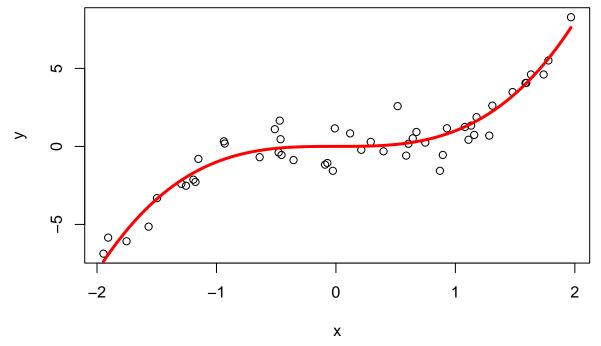


## Adding a curve to an existing plot

To add a curve to an existing plot, use curve() with add=TRUE

```
n = 50
set.seed(0)
x = sort(runif(n, min=-2, max=2))
y = x^3 + rnorm(n)

plot(x, y)
curve(x^3, lwd=3, col="red", add=TRUE)
```

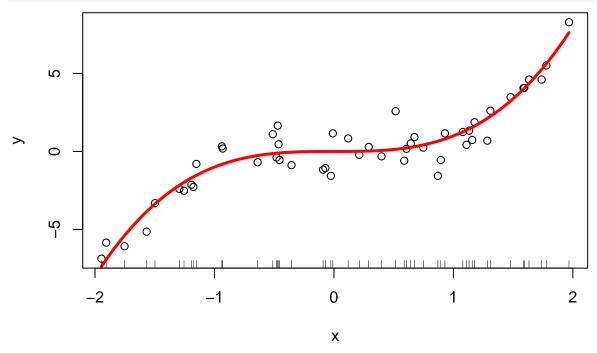


```
\# Note: the x argument here and the x vector we defined above are different! \# Reminder: x here is a symbol
```

## Adding a rug to an existing plot

To add a rug to an existing plot (just tick marks, for where the x points occur), use rug()

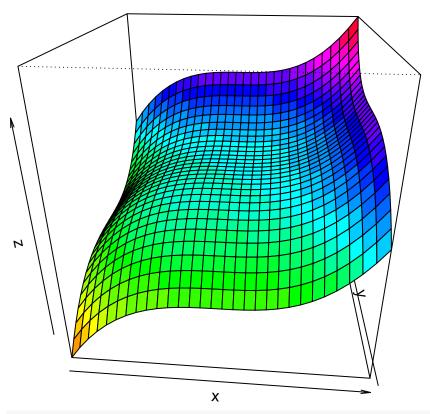
```
plot(x, y)
curve(x^3, lwd=3, col="red", add=TRUE)
rug(x)
```

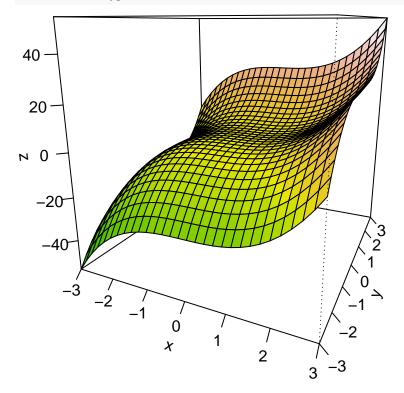


## Drawing a surface

To draw a surface, use surface(), available on canvas. This is a function written by Professor Ryan Tibshirani, relying on the built-in persp() function.

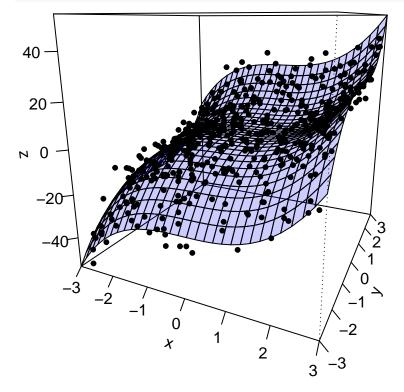
```
source("surface.R")
surface(x^3 + y^3, from.x=-3, to.x=3, from.y=-3, to.y=3)
```





### Adding points to a surface

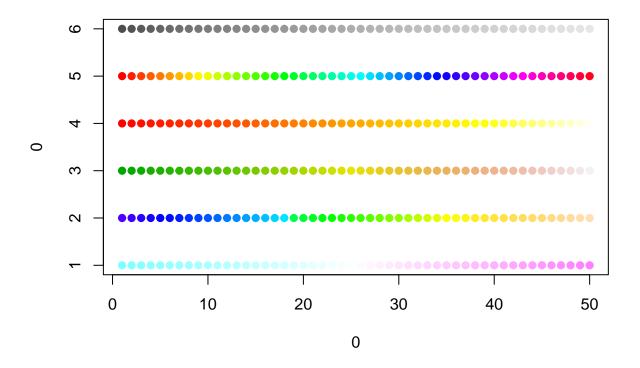
To add points to a surface, save the output of surface(). Then use trans3d(), to transform(x,y,z) coordinates to (x,y) coordinates that you can pass to points()



### Color palettes

Color palettes are functions for creating vectors of contiguous colors, just like gray.colors(), rainbow(), heat.colors(), terrain.colors(), topo.colors(), cm.colors(). Given a number n, each of these functions just returns a vector of colors (names, stored as strings) of length n

```
n = 50
plot(0, 0, type="n", xlim=c(1,n), ylim=c(1,6))
points(1:n, rep(6,n), col=gray.colors(n), pch=19)
points(1:n, rep(5,n), col=rainbow(n), pch=19)
points(1:n, rep(4,n), col=heat.colors(n), pch=19)
points(1:n, rep(3,n), col=terrain.colors(n), pch=19)
points(1:n, rep(2,n), col=topo.colors(n), pch=19)
points(1:n, rep(1,n), col=cm.colors(n), pch=19)
```

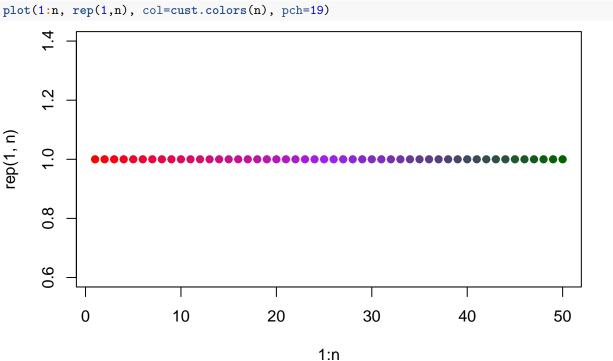


## Creating a custom color palette

To create a custom palette, that interpolates between a set of base colors, colorRampPalette()

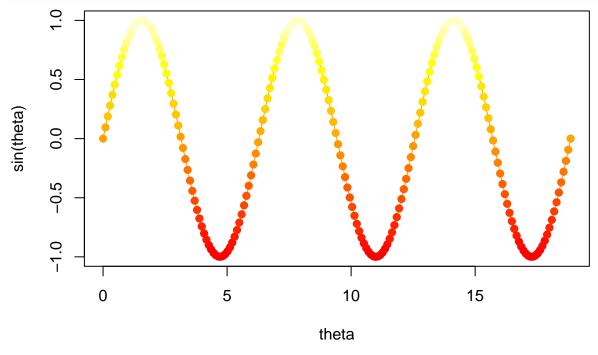
```
cust.colors = colorRampPalette(c("red","purple","darkgreen"))
class(cust.colors)
```

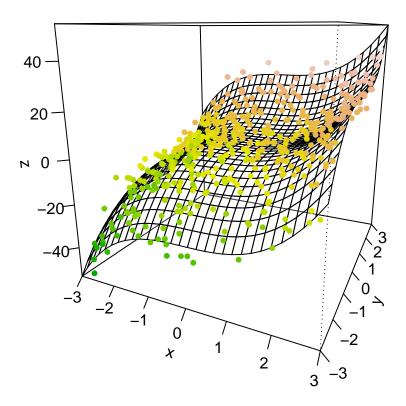
## [1] "function"



### Coloring points by value

Coloring points according to the value of some variable can just be done with a bit of indexing, and the tools you already know about colors





## Summary

- plot(): generic plotting function
- points(): add points to an existing plot
- lines(), abline(): add lines to an existing plot
- text(), legend(): add text to an existing plot
- rect(), polygon(): add shapes to an existing plot
- hist(), image(): histogram and heatmap
- heat.colors(), topo.colors(), etc: create a color vector
- density(): estimate density, which can be plotted
- contour(): draw contours, or add to existing plot
- curve(): draw a curve, or add to existing plot