R DataStructures

Data Types

Scalar types

numeric, character, logical, complex

Complex types

vectors,
matrices, arrays
data frames,
lists,
factors

Scalar Types

```
a = 10
b = "abc"
c = TRUE
d = 10 + i 20
```

Arithmetic Operators

| Operator | Description |
|-----------|-----------------------------|
| + | addition |
| - | subtraction |
| * | multiplication |
| 1 | division |
| ^ or ** | exponentiation |
| x %% y | modulus (x mod y) 5%%2 is 1 |
| x %/0/% y | integer division 5%/%2 is 2 |

Logical Operators

| Operator | Description |
|------------------------------|--------------------------|
| < | less than |
| <= | less than or equal to |
| > | greater than |
| >= | greater than or equal to |
| == | exactly equal to |
| != | not equal to |
| !x | Not x |
| $\mathbf{x} \mid \mathbf{y}$ | x OR y |
| x & y | x AND y |
| isTRUE(x) | test if x is TRUE |

Numeric Functions

| Function | Description |
|-----------------------------------|---|
| abs(x) | absolute value |
| $\mathbf{sqrt}(x)$ | square root |
| ceiling(x) | ceiling(3.475) is 4 |
| floor(x) | floor(3.475) is 3 |
| trunc(x) | trunc(5.99) is 5 |
| round(x, digits=n) | round(3.475, digits=2) is 3.48 |
| signif(x, digits=n) | signif(3.475, digits=2) is 3.5 |
| $\cos(x)$, $\sin(x)$, $\tan(x)$ | also $a\cos(x)$, $\cosh(x)$, $a\cosh(x)$, etc. |
| $\log(x)$ | natural logarithm |
| log10 (<i>x</i>) | common logarithm |
| exp(x) | e^ <i>x</i> |

Character Functions

Function

substr(x, start=n1, stop=n2)

Description

Extract or replace substrings in a character vector.

x <- "abcdef" substr(x, 2, 4) is "bcd" substr(x, 2, 4) <- "22222" is "a222ef"

grep(pattern, x ,
ignore.case=FALSE,
fixed=FALSE)

Search for *pattern* in *x*. If fixed =FALSE then *pattern* is a <u>regular expression</u>. If fixed=TRUE then *pattern* is a text string. Returns matching indices.

grep("A", c("b", "A", "c"), fixed=TRUE) returns

| <pre>sub(pattern, replacement, x, ignore.case =FALSE, fixed=FALSE)</pre> | Find <i>pattern</i> in <i>x</i> and replace with <i>replacement</i> text. If fixed=FALSE then <i>pattern</i> is a regular expression. If fixed = T then <i>pattern</i> is a text string. sub("\\s",".","Hello There") returns "Hello.There" |
|--|---|
| strsplit(x, split) | Split the elements of character vector <i>x</i> at <i>split</i> . strsplit("abc", "") returns 3 element vector "a","b","c" |
| paste(, sep=""") | Concatenate strings after using <i>sep</i> string to seperate them. paste("x",1:3,sep="") returns c("x1","x2" "x3") paste("x",1:3,sep="M") returns c("xM1","xM2" "xM3") paste("Today is", date()) |

toupper(x)

Uppercase

tolower(x)

Lowercase

Type Check & Conversion

Type Check

Use is. foo to test for data type foo.
 Returns TRUE or FALSE

• is.numeric(), is.character(), is.vector(), is.matrix(), is.data.frame()

Type Conversion

 Type conversions in R work as you would expect. For example, adding a character string to a numeric vector converts all the elements in the vector to character.

Use as. foo to explicitly convert it.

 as.numeric(), as.character(), as.vector(), as.matrix(), as.data.frame)

a. <u>Creating vectors</u>

- -concatenate function: c(), combines specified values in a vector
 - > v=c(1,2,3,20)
- -colon operator: :, generates an ordered sequence incremented by 1
 - > v=1:4
- -sequence function: seq(), generates an ordered sequence incremented by the specified value or a sequence of the specified length
 - > v=seq(from=5,to=6,by=0.1)
 - > v=seq(from=-10,to=-5,length.out=10)
- -repeat function: rep(), generates a vector of specified length containing the same value in each element of the vector
 - > v=rep(x=4,by=5)
- -numeric function: numeric(), generates a vector of specified length filled with 0's
 - > v=numeric(length=5)
- -vector function: vector(), generates a vector of specified length filled with FALSE's
 - > v=vector(length=5)

b. <u>Useful vector functions</u>

- -mathematical operators: +, -, *, /
- -logical operators: <, >, <=, >=, ==, !=
- -length(): returns the length of the vector
- -max(): returns the maximum value contained in the vector
- -min(): returns the minimum value contained in the vector
- -sum(): returns the sum of the values in the vector
- -cumsum(): returns the cumulative sum for each element of the vector
- *-mean()*: returns the mean of the vector
- -range(): returns the minimum and maximum values
- -var(): returns the variance of the vector
- -sd(): returns the standard deviation of the vector
- -sort(): returns a sorted version of the vector
- -order(): returns the numerical indices of vector elements in sorted order

c. <u>Vector indexing and subsetting</u>

Because vectors are an ordered list, a single element or subset of elements can be referred to using square brackets, [], and a numerical index.

```
> v=c(1,3,5,9,13)
> v[1]
[1] 1
> v[4]
[1] 9
> v[c(1,3,5)]
[1] 1 5 13
> v[-3]
[1] 1 3 9 13
```

An alternative means of indexing is a vector of logical values.

```
> v=c(1,3,5,9,13)
> v>3
[1] FALSE FALSE TRUE TRUE TRUE
>v[v>3]
[1] 5 9 13
```

The which() function creates numerical indices from a logical vector.

```
> v=c(1,3,5,9,13)
> v>3
[1] FALSE FALSE TRUE TRUE TRUE
> which(v>3)
[1] 3 4 5
```

Note that you cannot delete an element from a vector, but you can reassign a subset of a vector to the same variable.

```
> v=c(1,3,5,9,13)
> v
[1] 1 3 5 9 13
> v=v[v>3]
> v
[1] 5 9 13
```

All columns in a matrix must have the same type and the same length.

The general format is:

```
mymatrix <- matrix(vector, nrow=r, ncol=c,
    byrow=FALSE,dimnames=list(char_vector_rownames,
    char_vector_colnames))</pre>
```

byrow=TRUE indicates that the matrix should be filled by rows.

byrow=FALSE indicates that the matrix should be filled by columns (the default).

dimnames provides optional labels for the columns and rows.

```
# generates 5 x 4 numeric matrix
  y<-matrix(1:20, nrow=5,ncol=4)
# another example
  cells < -c(1,26,24,68)
  rnames <- c("R1", "R2")
  cnames <- c("C1", "C2")
  mymatrix <- matrix(cells, nrow=2, ncol=2,
  byrow=TRUE, dimnames=list(rnames, cnames))
#Identify rows, columns or elements using subscripts.
  x[,4] # 4th column of matrix
  x[3,] # 3rd row of matrix
  x[2:4,1:3] # rows 2,3,4 of columns 1,2,3
```

Useful matrix functions

- ***many of these will work on higher dimensional arrays too
- -dim(): returns the dimensions (number of rows and columns) of the matrix
- -nrow(): returns the number of rows in the matrix
- -ncol(): returns the number of columns in the matrix
- -rownames(): returns the row names of the matrix; can also be used for assignment
- -colnames(): returns the column names of the matrix; can also be used for assignment
- -rbind(): add a vector to a specified matrix as a new row at the bottom of the matrix
- -cbind(): add a vector to a specified matrix as a new column at the furthest right
- -%*%: matrix multiplication
- -t(): transpose the matrix
- -colMeans(): calculate the mean of each column of the matrix
- -colSums(): calculate the sum of each column of the matrix
- -apply(): applies a function that works on a vector to each row or column of a matrix

dimnames()

Element Access in Matrices

```
> M=matrix(1:4,nrow=2,ncol=2)
> M[2,1]
[1] 2
> M[2,2]
[1] 4
> M[4]
[1] 4
> M[,2]
[1] 3 4
```

Matrix operations

```
# matrix addition
matC = matA + matB
# matrix subtraction
matC = matA - matB
# scalar multiplication
matA = matrix(c(3,-1,0,5),2,2,byrow=TRUE)
matC = 2*matA
# matrix multiplication
matA = matrix(1:4,2,2,byrow=TRUE)
matB = matrix(c(1,2,1,3,4,2),2,3,byrow=TRUE)
matC = matA%*%matB
matC
```

Matrix operations

```
# create identity matrix
matI = diag(2)
matl
matA = matrix(c(1,2,3,4), 2, 2, byrow=TRUE)
matl%*%matA
matA%*%matI
# matrix inversion
matA
matA.inv = solve(matA)
matA.inv
matA%*%matA.inv
matA.inv%*%matA
```

Vector to matrix

- names(xvec) = c("x1", "x2", "x3")
- dim(xvec)

coerce vector to class matrix: note #column vector is created

- xvec = as.matrix(xvec)
- xvec
- class(xvec)

Data frames

Data frames

A data frame is more general than a matrix, in that different columns can have different modes (numeric, character, factor, etc.).

```
d <- c(1,2,3,4)
e <- c("red", "white", "red", NA)
f <- c(TRUE,TRUE,TRUE,FALSE)
mydata <- data.frame(d,e,f)
names(mydata) <- c("ID","Color","Passed")
#variable names</pre>
```

Element access in Data frames

There are a variety of ways to identify the elements of a dataframe.

myframe[3:5] # columns 3,4,5 of dataframe myframe[c("ID","Age")] # columns ID and Age from dataframe myframe\$X1 # variable x1 in the dataframe

Operations on Dataframes

b. <u>Useful data frame functions</u>

- -rbind(): add a vector to a specified matrix as a new row at the bottom of the matrix
- -cbind(): add a vector to a specified matrix as a new column at the furthest right
- -colMeans(): calculate the mean of each column of the matrix
- -colSums(): calculate the sum of each column of the matrix
- -apply(): applies a function that works on a vector to each row or column of a matrix
- -merge(): joins two data frames together using a shared column as an index
- -lapply(): analogous to apply(), but operates on lists and returns a list
- -sapply(): the same functionality as lapply(), but returns a matrix or vector

Factors

Factors

The factor stores the nominal values as a vector of integers in the range [1... k] (where k is the number of unique values in the nominal variable), and an internal vector of character strings (the original values) mapped to these integers.

Creating factors

stores gender as 20 1s and 30 2s and associates # 1=female, 2=male internally (alphabetically)

An ordered collection of objects (components). A list allows you to gather a variety of (possibly unrelated) objects under one name.

example of a list with 4 components - a string, a numeric vector, a matrix, and a scalar

w = list(name="Fred", mynumbers=a, mymatrix=y, age=5.3)

example of a list containing two lists
v = c(list1,list2)

Identify elements of a list using the [[]] convention.

mylist[[2]] # 2nd component of the list

Arrays

Arrays

Arrays are similar to matrices but can have more than two dimensions. See **help(array)** for details.