Professor Bear - R Data structures

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1.1: Data structures in R

Data Structures are the programmatic way of storing data so that data can be used efficiently. One can think of a data structure as a container for data.

1.1.1: Vectors

Usually when you create variables in R, you create **vectors**. A vector is simply a set of elements *of the same class* (e.g. character, numeric, integer, or logical -as in True/False). It is the basic data structure in R. Most commonly, you will use the `c()` function (c stands for concatenate) to create vectors. :

```
v1 <- c(1,2,3,4,5) #creates a numeric vector
v1

## [1] 1 2 3 4 5

v2 <- c(1L, 2L) #creates an integer vector
v2

## [1] 1 2

v3 <- c(TRUE, FALSE) #creates a logical vector
v3

## [1]  TRUE FALSE

v4 <- c("a", "b", "c") #creates a character vector
v4

## [1] "a" "b" "c"

v5 <- c(1+0i, 3+5i) #creates a complex vector
v5

## [1] 1+0i 3+5i
```

Operations on vectors

Once you have a vector (or a list of numbers) in memory most basic operations are available. This makes R very powerful.

```
v1
```

```
## [1] 1 2 3 4 5
```

```
v1 + 5 # add 5 to each of the numbers
```

```
## [1]  6  7  8  9 10
```

```
v1*3 # Multiply each of the numbers by 3
```

```
## [1]  3  6  9 12 15
```

```
v1/3 # Divide each of the numbers by 3
```

```
## [1] 0.3333333 0.6666667 1.0000000 1.3333333 1.6666667
```

```
log(v1) # Take the log of each of the numbers
```

```
## [1] 0.0000000 0.6931472 1.0986123 1.3862944 1.6094379
```

We can even add vectors to vectors.

```
a <- c(1,2,3)
a
```

```
## [1] 1 2 3
```

```
b <-  c(1,2,3,4,5)
b
```

```
## [1] 1 2 3 4 5
```

```
a+b
```

```
## Warning in a + b: longer object length is not a multiple of shorter object
## length
```

```
## [1] 2 4 6 5 7
```

If the lengths of the vectors differ then you may get an error message, a warning message and unpredictable results. It is best if they are the same length.

```
a <- c(1,2,3)
a
```

```
## [1] 1 2 3
```

```
length(a)
```

```
## [1] 3
```

```
b <-  c(3,4,5)
b
```

```
## [1] 3 4 5
```

```
length(b)
```

```
## [1] 3
```

```
a+b
```

```
## [1] 4 6 8
```

If you mix in a vector elements that are of a different class (for example numerical and logical), R will **coerce** to the minimum common denominator, so that every element in the vector is of the same class.

```
b <- c(33, "bear")
class(b)
```

```
## [1] "character"
```

Note what happens when a vector is placed in another vector.

```
x <- c(2, 3, 5)
x
```

```
## [1] 2 3 5
```

```
class(x)
```

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

```
y <- c(6, x, 2)
y
```

```
## [1] 6 2 3 5 2
```

```
class(y)
```

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

1.1.2: Lists

A list is a generic data structure containing other objects. Unlike a vector it allows us to create a mixed data sequence.

```
w <-  c(1, 2, 3)
w
```

```
## [1] 1 2 3
```

```
class(w)
```

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

```
x <- c("a", "b", "c", "d", "e")
x
```

```
## [1] "a" "b" "c" "d" "e"
```

```
class(x)
```

```
## [1] "character"
```

```
y <- c(TRUE, FALSE, TRUE, FALSE)
y
```

```
## [1]  TRUE FALSE  TRUE FALSE
```

```
z <- list(w, y, x, 33, "bear", FALSE) # create a mixed data type list
z
```

```
## [[1]]
## [1] 1 2 3
##
## [[2]]
## [1]  TRUE FALSE  TRUE FALSE
##
## [[3]]
## [1] "a" "b" "c" "d" "e"
##
## [[4]]
## [1] 33
##
## [[5]]
## [1] "bear"
##
## [[6]]
## [1] FALSE
```

```
class(z)
```

```
## [1] "list"
```

List Slicing

We use the single square bracket "[]" operator to access elements at the first level of a list. The index 1 is the first element. We use the doubkle square bracket "[]" operator to access elements at the second level, etc.

```
z
```

```
## [[1]]
## [1] 1 2 3
##
## [[2]]
```

```
## [1]  TRUE FALSE  TRUE FALSE
##
## [[3]]
## [1] "a" "b" "c" "d" "e"
##
## [[4]]
## [1] 33
##
## [[5]]
## [1] "bear"
##
## [[6]]
## [1] FALSE
```

```
z[1] # First element
```

```
## [[1]]
## [1] 1 2 3
```

```
z[[2]] # cond element
```

```
## [1]  TRUE FALSE  TRUE FALSE
```

```
z[[2]][1] # First element of the second element
```

```
## [1] TRUE
```

1.1.3: Arrays

Vectors are one-dimensional arrays in R and matrices are two-dimensional arrays in R. We can create n-dimensional arrays as a set of stacked matrices of identical dimensions. This will be discuss in the matrices section below. The term arrays is discussed here becuase most programming languages use the term array. The reason R uses the term vector, is the basic operations can be applied to vectors whereas most programming languages use functions and loops to apply operations to arrays.

1.1.4: Matrices

Matrices are n-dimensional vectors (usually two-dimensional). We build matrices from vectors (one-dimensional arrays). You can create a matrix in two ways. By using the command *matrix*.

```
x<-matrix(c(1,2,3,4,5,6), nrow = 2, ncol = 3)
x
```

```
##      [,1] [,2] [,3]
## [1,]    1    3    5
## [2,]    2    4    6
```

```
class(x)
```

```
## [1] "matrix"
```

Or using *cbind()* or *rbind()* to add columns or rows to a vectors.

```
x <- c(1, 2, 3)         # Creates a vector `x' of 3 values.
x

## [1] 1 2 3

class(x)

## [1] "numeric"

y <- c(55, 33, 11)         # Creates another vector `y' of 3 values.
y

## [1] 55 33 11

class(y)

## [1] "numeric"

a<-rbind(x, y) # Creates a 2 x 3 matrix.  Note that the rows are appended
a

##   [,1] [,2] [,3]
## x    1    2    3
## y   55   33   11

class(a)

## [1] "matrix"

b<-cbind(x, y)   # Creates a 3 x 2 matrix.  Note that the columns are
appended
b

##      x  y
## [1,] 1 55
## [2,] 2 33
## [3,] 3 11

class(b)

## [1] "matrix"
```

Matrix element access

As with vectors, square brackets extract specific values from a matrix but more values are used within the brackets seperated by commas for each diminasion.

```
b
```

```
##      x  y
## [1,] 1 55
## [2,] 2 33
## [3,] 3 11
```

```
b[,2] # Extracts the second column
```

```
## [1] 55 33 11
```

```
b[1,]  # Extracts the first row
```

```
##  x  y
##  1 55
```

```
b[1,2] # Extracts element in the first row and the second column
```

```
##  y
## 55
```

```
b[1,2]<=55 # Recplaces the element in the first row and the second column
with 55
```

```
##    y
## TRUE
```

```
b
```

```
##      x  y
## [1,] 1 55
## [2,] 2 33
## [3,] 3 11
```

```
b[1,] <- c(3,3) # Replaces the first row
b
```

```
##      x  y
## [1,] 3  3
## [2,] 2 33
## [3,] 3 11
```

Matrix operations and functions

R supports a variety of matrix functions, including: det(), which returns the matrix's determinant; t(), which transposes the matrix; solve(), which inverts the the matrix; dim() command returns the dimensions of your matrix.

```
b
```

```
##      x  y
## [1,] 3  3
## [2,] 2 33
## [3,] 3 11
```

```
dim(b)

## [1] 3 2
```

1.1.5: Data frames

A data frame is used for storing data tables of mixed data type. Typically, data from an excel sheet will be imported in to R as a data frame. We will use the built-in data set "InsectSprays" to discuss data frames.

```
data(InsectSprays)
names(InsectSprays)

## [1] "count" "spray"

head(InsectSprays,3)

##   count spray
## 1    10     A
## 2     7     A
## 3    20     A

dim(InsectSprays)

## [1] 72  2

nrow(InsectSprays)

## [1] 72

ncol(InsectSprays)

## [1] 2

levels(InsectSprays$spray)

## [1] "A" "B" "C" "D" "E" "F"

summary(InsectSprays)

##      count       spray
##  Min.   : 0.00   A:12
##  1st Qu.: 3.00   B:12
##  Median : 7.00   C:12
##  Mean   : 9.50   D:12
##  3rd Qu.:14.25   E:12
##  Max.   :26.00   F:12
```

1.17: Further resources

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