R syntax in a nutshell - Part II

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- 1.7 Building simple databases: indexing/subsetting of vectors, arrays and data frames
- 1.8 Lists
- 1.9 Coercion of individual object types

1.6 Arrays, matrices, data frames

Several kinds of table-like objects exist in R. **Data frames** are data objects to be processed by statistics, with "observations" as columns (elements/oxides in geochemistry) and "cases" (samples) in rows. They can contain columns of any mode, even mixed modes; thus they are not meant for matrix operations.

On the other hand, all elements of a **matrix** can only be of a single mode (numeric, most commonly). **Arrays** are generalized matrices: they must have a single mode but can have any number of dimensions. Although superficially similar, these three types of objects must not be confused.

matrix(data = NA, nrow = 1, ncol = 1, byrow = FALSE)

This command defines a matrix of nrow rows and ncol columns, filled by the data (if data have several elements, they will be used down the columns, unless an extra parameter byrow=TRUE is provided). For instance:

In [2]: x <- matrix(1:12,3,4,byrow=TRUE)
print(x)</pre>

```
[1,1] [,2] [,3] [,4]
[1,] 1 2 3 4
[2,] 5 6 7 8
[3,] 9 10 11 12
```

As a default, filling a matrix with data — as well as matrix division by a vector — proceeds along columns, not rows!

array(data = NA, dim = length(data))

Defines a new data array and fills it with data. The argument dim is a vector of length one or more, giving maximal dimensions in each of the directions.

1.6.1 Matrix/data frame operations

Matrices can be subject to scalar operations using the common operators ($+-*/^{\circ}$). Similar to vectors, the shorter component is recycled as appropriate. Useful functions for matrix/data frame manipulations are summarized in the following Table:

Function	Meaning
nrow(x)	number of rows
ncol(x)	number of columns
rownames(x)	row names
colnames(x)	column names
<pre>rbind(x,y)</pre>	binds two objects (matrices or data frames) of the same ncol (or vectors of the same length) as rows
<pre>cbind(x,y)</pre>	binds two objects (matrices or data frames) of the same nrow (or vectors of the same length) as columns
t(x)	matrix transposition
apply(X,MARGIN,FUN)	applies function FUN (for vector manipulations) along the rows (MARGIN = 1) or columns (MARGIN = 2) of a data matrix X
x%*%y	matrix multiplication (does not work on a data frame!)
solve(A)	matrix inversion
dix(x)	diagonal elements of a matrix

It is worth noting that matrix multiplication is performed using the %*% operator.



The function `solve` can serve to solving a set of linear argebraic equations. Here is an example; we shal try to solve the following set of linear equations:

$$x + 2y - 0.7z = 21$$

$$3x + 0.2y - z = 24$$

$$0.9x + 7y - 2z = 27$$

Let's first specify the matrix of coefficients to the individual variables:

In [3]: A <- matrix(c(1,3,0.9,2,0.2,7,-0.7,-1,-2),3,3)
print(A)</pre>

[,1] [,2] [,3] [1,] 1.0 2.0 -0.7

[2,] 3.0 0.2 -1.0

[3,] 0.9 7.0 -2.0

And then the vector of right hand sides:

In [4]: y <- c(21,24,27)

 $21\cdot 24\cdot 27$

In [5]: x <- solve(A,y)</pre>

```
X
```

30 · 20 · 70

```
In [6]: # checking the result, back-calculating y
print(A%*%x)
```

```
[,1]
[1,] 21
[2,] 24
[3,] 27
```

Of the functions presented in the table, some explanation is required for apply:

apply(X, MARGIN, FUN,...)

If X is a matrix, it is split into vectors along rows (if MARGIN = 1) or columns (if MARGIN = 2). To each of these vectors is applied the function FUN with optional parameters ... passed to it. For instance, we can calculate row sums of a matrix:

1.7 Building simple databases: indexing/subsetting of vectors, arrays and data frames

In real life, one often needs to select some elements of a vector or a matrix, fulfilling certain criteria. This can be achieved using logical conditions or logical variables placed in **square brackets** after the object name. Subsets can be also used on the left hand side of the assignments when replacement of selected elements by certain values is desired.

1.7.1 Vectors

Subsets of a vector may be selected by appending to the name of the vector an index vector in square brackets. For example, we can first create a named vector:

```
In [8]: x <- c(1,12,15,NA,16,13,0,NA,NA)
    names(x) <- c("Pl","Bt","Mu","Q","Kfs","Ky","Ol","Px","C")
    print(x)

Pl Bt Mu Q Kfs Ky Ol Px C
    1 12 15 NA 16 13 0 NA NA</pre>
```

Index vectors can be of several types: logical, numeric (with positive or negative values), and character:

1. Logical vector

```
In [9]: \#x[x>10] # all elements of x higher than 10 (or NA)
x[!is.na(x)] # all elements of x that are available
```

Pl: 1 **Bt:** 12 **Mu:** 15 **Kfs:** 16 **Ky:** 13 **Ol:** 0

2. Numeric vector with positive values

```
In [10]: x[1:5]  # the first five elements
    x[c(1,5,7)] # 1st, 5th and 7th elements
```

PI: 1 **Bt:** 12 **Mu:** 15 **Q:** <NA> **Kfs:** 16

PI: 1 **Kfs:** 16 **OI:** 0

3. Numeric vector with negative values (specifies elements to be excluded)

```
In [11]: x x[-(1:5)] # all elements except for the first five

Pl: 1 Bt: 12 Mu: 15 Q: <NA> Kfs: 16 Ky: 13 Ol: 0 Px: <NA> C: <NA>
```

4. Character vector (referring to the element names)

```
In [12]: x[c("Q","Bt","Mu")]
```

Q: <NA> **Bt:** 12 **Mu:** 15

Ky: 13 **OI:** 0 **Px:** <NA> **C:** <NA>

1.7.2 Matrices/data frames

Elements of a matrix are presented in the order [row,column]. If nothing is given for a row or column, it means no restriction. For instance:

```
x[1,] # (all columns) of the first row

x[,c(1,3)] # (all rows) of the first and third columns

x[1:3,-2] # all columns (apart from the 2nd) of rows 1-3
```

If the result is a single row or column, it is automatically converted to a vector. To prevent such a behaviour, one can supply an optional parameter drop=FALSE, e.g.:

```
x[1,,drop=FALSE] # (all columns) of the 1st row, keep as matrix

Moreover, matrices can be manipulated using index arrays. This concept is best explained on an example. Let's assume a matrix defined as:
```

```
x \leftarrow matrix(1:20,4,5)
```

If the elements [1,3], [2,2] and [3,1] in x are to be replaced by zeroes, we can create an index array i containing the element coordinates:

```
1 5 9
               13
[1,]
                   17
[2,]
     2 6 10
               14
                   18
        7 11
[3,]
     3
               15
                   19
       8 12
               16
                   20
[4,]
```

```
In [14]: print(x[1,])
         print(x[1,,drop=FALSE])
          [1] 1 5 9 13 17
               [,1] [,2] [,3] [,4] [,5]
                    5 9
                               13
          [1,]
In [15]: print(x[,c(1,3)])
               [,1] [,2]
         [1,]
                  1
          [2,]
                  2
                      10
                  3
         [3,]
                      11
         [4,]
                      12
In [16]: print(x[1:3,-2])
              [,1] [,2] [,3] [,4]
          [1,]
                 1
                      9
                           13
                                17
                  2
                      10
                           14
                                18
          [2,]
         [3,]
                  3
                      11
                           15
                                19
In [17]: x <- matrix(1:20,4,5)
         print(x)
         i <- matrix(c(1,2,3,3,2,1),3,2)</pre>
         print(i)
               [,1] [,2] [,3] [,4] [,5]
         [1,]
                 1
                            9
                                13
                                     17
                       5
          [2,]
                  2
                       6
                           10
                                14
                                     18
                      7
                  3
                           11
                                15
                                     19
         [3,]
                 4
                      8
                           12
                                16
                                     20
         [4,]
               [,1] [,2]
          [1,]
                  1
                       3
                       2
         [2,]
                  2
         [3,]
                  3
                       1
In [18]: x[i]
         9 \cdot 6 \cdot 3
In [19]: x[i] <- 0
         print(x)
               [,1] [,2] [,3] [,4] [,5]
          [1,]
                1
                    5
                          0
                                13
                                     17
          [2,]
                  2
                       0
                                     18
                           10
                                14
                      7
         [3,]
                  0
                           11
                                15
                                     19
          [4,]
                  4
                       8
                           12
                                16
                                      20
         The situation for multidimensional arrays is analogous — just the relevant number of
         dimensions is higher than two.
```

R involves numerous built in datasets that can be used to demonstrate its capabilities.

The object `islands` is a vector with areas of landmasses (i.e. islands and continents) exceeding 10 000 sq. miles. Before we can start working, we need to attach the data object using the command:

```
In [20]: data(islands)
```

To-do list:

- Display the whole vector. What is the area of Luzon?
- What is the average value of the whole vector?
- Which continent is the largest and what is its area?
- Which continents/islands have an area larger than 5000 sq. miles?
- Display the names of 15 smallest and largest continents/islands.
- Assuming that these are British miles, recalculate the data to km2 (1 sq mi = 2.59 km2)

```
In [21]: head(islands) # print just a few observations
  islands["Luzon"]
```

Africa: 11506 Antarctica: 5500 Asia: 16988 Australia: 2968 Axel Heiberg: 16 Baffin: 184

Luzon: 42

```
In [22]: mean(islands)
```

1252.72916666667

```
In [23]: max(islands)
which(islands==max(islands))
```

16988

Asia: 3

```
In [24]: names(islands)[islands>5000]
```

'Africa' · 'Antarctica' · 'Asia' · 'North America' · 'South America'

```
In [25]: z <- sort(islands)
    print(z[1:15])</pre>
```

```
Hainan Prince of Wales
   Vancouver
                                                            Timor
          12
                           13
                                                               13
                                    New Britain
      Kyushu
                       Taiwan
                                                      Spitsbergen
                           14
                                             15
          14
                                                               15
Axel Heiberg
                     Melville
                                    Southampton Tierra del Fuego
                                                               19
          16
                            16
                                             16
                                          Celon
       Devon
                        Banks
          21
                                              25
                            23
```

```
In [26]: out <- z[(length(z)-14):length(z)]
    print(out)</pre>
```

Britain	Honshu	Sumatra	Baffin	Madagascar
84	89	183	184	227
Borneo	New Guinea	Greenland	Australia	Europe
280	306	840	2968	3745
Antarctica	South America	North America	Africa	Asia
5500	6795	9390	11506	16988

```
In [27]: # Or, alternatively:
   out <- rev(sort(islands))[1:15]
   print(out)</pre>
```

Antarctica	South America	North America	Africa	Asia
5500	6795	9390	11506	16988
Borneo	New Guinea	Greenland	Australia	Europe
280	306	840	2968	3745
Britain	Honshu	Sumatra	Baffin	Madagascar
84	89	183	184	227

In [28]:	print(islands*2.59)	#	recalculate	the	data 1	to km2	(1 sa	mi =	= 2.59 km2)

Africa	Antarctica	Asia	Australia
29800.54	14245.00	43998.92	7687.12
Axel Heiberg	Baffin	Banks	Borneo
41.44	476.56	59.57	725.20
Britain	Celebes	Celon	Cuba
217.56	189.07	64.75	111.37
Devon	Ellesmere	Europe	Greenland
54.39	212.38	9699.55	2175.60
Hainan	Hispaniola	Hokkaido	Honshu
33.67	77.70	77.70	230.51
Iceland	Ireland	Java	Kyushu
103.60	85.47	126.91	36.26
Luzon	Madagascar	Melville	Mindanao
108.78	587.93	41.44	93.24
Moluccas	New Britain	New Guinea	New Zealand (N)
75.11	38.85	792.54	113.96
New Zealand (S)	Newfoundland	North America	Novaya Zemlya
150.22	111.37	24320.10	82.88
Prince of Wales	Sakhalin	South America	Southampton
33.67	75.11	17599.05	41.44
Spitsbergen	Sumatra	Taiwan	Tasmania
38.85	473.97	36.26	67.34
Tierra del Fuego	Timor	Vancouver	Victoria
49.21	33.67	31.08	212.38

1.8 Lists

Lists are ordered collections of other objects, known as components, which do not have to be of the same mode or type. Thus lists can be viewed as very loose groupings of R objects, involving various types of vectors, data frames, arrays, functions and even other lists.

Components are numbered and hence can be addressed using their sequence number given in double square brackets, x[[3]]. Moreover, components may be named and referenced using an expression of the form $list_name$component_name$. Subsetting is similar to that of other objects, described above.

list.name <- list(component_name_1=, component_name_2=...)</pre>

Builds a list with the given components.

Here is a simple real-life example of a list definition:

```
In [29]: x1 <- c("Luckovice","9 km E of Blatna","disused quarry")
    x2 <- "melamonzonite"
    x3 <- c(47.31,1.05,14.94,7.01,8.46,10.33)
    names(x3) <- c("SiO2","TiO2","Al2O3","FeO","MgO","CaO")
    luc <- list(ID="Gbl-4",Locality=x1,Rock=x2,major=x3)
    print(luc)</pre>
```

```
$ID
[1] "Gbl-4"

$Locality
[1] "Luckovice" "9 km E of Blatna" "disused quarry"

$Rock
[1] "melamonzonite"

$major
    Si02    Ti02    Al203    Fe0    Mg0    Ca0
47.31    1.05    14.94    7.01    8.46    10.33
```

As well as some examples of subsetting:

```
In [30]: luc[[1]]

'Gbl-4'

In [31]: luc$major # or Luc[[3]]

SiO2: 47.31 TiO2: 1.05 Al2O3: 14.94 FeO: 7.01 MgO: 8.46 CaO: 10.33

In [32]: luc[[2]][3]

'disused quarry'

In [33]: luc$major[c("SiO2","Al2O3")]
```

SiO2: 47.31 Al2O3: 14.94

1.9 Coercion of individual object types

R is generally reasonably good at dealing seamlessly with data types, converting them on the fly when needed. When necessary, there are a series of functions for testing the mode (or type) of an object: is.numeric(x), is.character(x), is.logical(x), is.matrix(x), is.data.frame(x)

At times there is a need to explicitly convert between data types/modes using functions such as: as.numeric(x), as.character(x), as.expression(x).

Less straightforward are: as.matrix(x), as.data.frame(x) which attempt to convert an object x to a matrix or data frame, respectively.

A more user-friendly way of converting data frames to matrices is provided by the function data.matrix that converts all the variables in a data frame x to numeric mode and then binds them together as the columns of a matrix.

```
In [34]: as.numeric("1.5444")*2

3.0888
```