Object types

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Prof. Dr. Claudius Gräbner-Radkowitsch Europa-University Flensburg, Department of Pluralist Economics

www.claudius-graebner.com | @ClaudiusGraebner | claudius@claudius-graebner.com





Prologue:



PrologueFeedback and exercises

- XX of you filled out the feedback survey. Main take-aways:
 - TBA
- What were the main problems with the exercises?

Goals for today

- I. Understand the main object types in R and their practical relevance
- II. Learn how to transform object types into each other
- III. Hear about some useful helper functions and the concept of vectorisation

Basic object types in R



Object types in R

To understand computations in R, two slogans are helpful:

Everything that exists is an **object**.

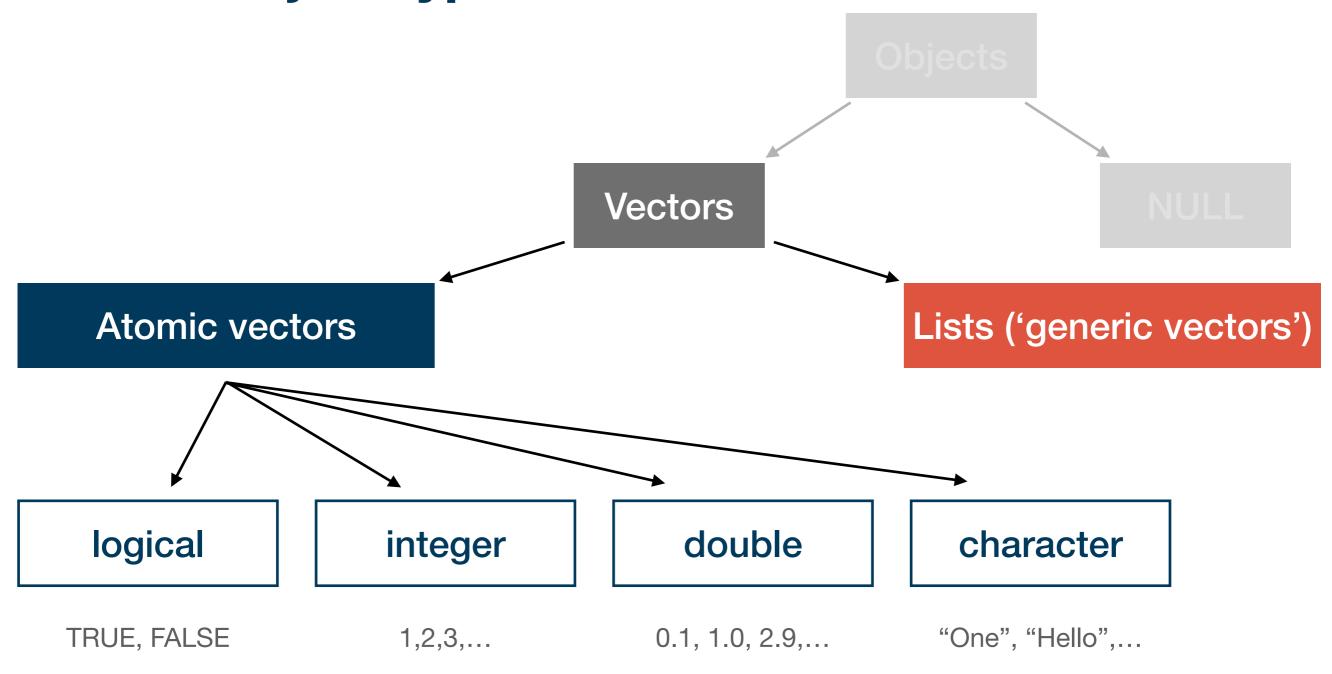
Everything that happens is a function call.

John Chambers

- We have learned quite a bit about functions, now we turn to objects
- We must distinguish different object types because functions operate differently depending on the type of the object we are processing
 - E.g.: 'adding up' numbers is different than 'adding up' words
- Fortunately, there are only a few basic types you must know about
 - More complex types are natural modifications of these basic types
- The most general type of object in R is a vector



Basic object types in R



 Among the more specific vector types, we will learn about factors and data frames later



Atomic vectors

- Atomic vectors are composed only of objects of the same type
 - We say that an atomic vector is of the same type as are its elements
 - We can test for this type using the function typeof()
- There are four main types of atomic vector that are most important:

Logical values: logical

- Only two* options: TRUE
 or FALSE
- Often the result of logical operations (e.g. 4>2)

Whole numbers: integer

- A whole number, followed by L:
- 1L, 2L, 100L, etc.
- Often the result of counting

Decimal numbers: double

- A number with the decimal sign.
- 2.0, 0.8, -7.5, etc.
- The 'standard' number you will use

Letters and words: character

- Might contain all kinds of tokens and start and end with "
 - "2", "Hello!", "vec 1", etc.

^{*:} We will see later that missing values are also considered logical in some instances, but this is basically irrelevant now.



Creating atomic vectors

• The easiest way to create atomic vectors is the function c() ('concatenate')

```
t_{vec} < c(1, 2, 3)
```

- The number of elements that are part of a vector are its length:
 - You can test for the length of a vector using length():

```
length(t_vec)
```

• c() can also be used to merge atomic vectors or arbitrary length:

```
t_vec_2 <- c(4, 5, 6)

t_vec_full <- c(t_vec, t_vec_2)</pre>
```



Coercion

- Sometimes we might want to change the type of an atomic vector
- In this context, the functions as.*() and is.*() are useful
 - Substitute the * for the type of vector, and you can test and transform them:

```
is.double(xx)

yy <- as.double(xx)

is.double(yy)</pre>
```

- But beware of some counter-intuitive transformation behaviour:
 - as.integer(22.9)
 - as.logical(99)

Intermediate exercises

Do the following tasks with you neighbour(s) and discuss open questions!

- 1. Create a vector containing the numbers 2, 5, 2.4 and 11.
- 2. What is the type of this vector?
- 3. Transform this vector into the type integer. What happens?
- 4. Do you think you can create a vector containing the following elements: "2", "Hallo", 4.0, and TRUE? Why? Why not?

Some useful helper functions

- There are some types of atomic vectors that you create frequently
 - Sequences of numbers, concatenated words, or repetitions
- For case 1 you may use the function seq() with the following arguments:
 - from, to: starting and end values of the sequence
 - by: increment steps of the sequences (must be numeric)
 - length.out: desired length of final sequence
 - along.with: creates sequence of same length as object
- Only one of the arguments (ii), (iii), and (iv) can be used, e.g.:
 - seq(-5, 5, by=2.5); seq(1, 4, length.out=10)



Some useful helper functions

- There are some types of atomic vectors that you create frequently
 - Sequences of numbers, concatenated words, or repetitions
- For case 2 you may use the function paste() with the argument sep:
 - sep: How should the input vectors be separated?
- This is useful, for instance, if you want to create file names:

```
paste("file_", seq(1,4), ".pdf", sep = "")
```

• Finally, if you want to repeat something, use rep():

```
rep("Cool!", 5)
```

Indexing

- Indexing means referencing a particular position of a vector
 - You do this by adding the position in square brackets to the end of the vector
 - v_c[3], for instance, returns the third element of the vector v_c
 - You can also use this logic to replace these elements:

```
v_c <- c("First", "Second", "Second", "Fourth")
v c[3] <- "Third!"</pre>
```

But you cannot use this to add new elements to a vector:

Add a fifth element to the vector v_c!

Vectorisation

- One reason why atomic vectors are so popular is that they allow for very fast computations
 - For the computer it is much easier to work with sets of objects that all behave the same
- Vectorisation means that an operation is applied to each element of a vector:

- "To vectorise" a task means to write it in a way that operations are applied to atomic vectors → in R, you should do that whenever possible
 - A slower alternative are loops, which we learn about later and which are unavoidable in certain situations

Intermediate practice

Do the following tasks with you neighbour(s) and discuss open questions!

- I. Create a vector with the numbers from -2 to 19 (step size: 0.75)
- II. Create an index vector for this first vector (note: an index vector is a vector with all possible indices of the original vector)
- III. Compute the log of each element of the first vector using vectorisation. Anything that draws your attention?
- IV. What happens if you concatenate vectors of different types using c()? Can you derive a systematization?
 - Remember that you can check for the type of an atomic vector using typeof()



Lists

- The second major type of vectors → sometimes called generic vectors
- Difference to atomic vectors: lists may contain objects of different types
 - Thus, the type of a list is always...

```
l_1 <- list(c(1,2), c("a", "b"), c(TRUE, FALSE, FALSE)); typeof(l_1)</pre>
```

Lists can be complex → get an overview using str():

```
Types of the elements

> str(l_1)
List of 3

$: num [1:2] 1 2
$: chr [1:2] "a" "b"
$: logi [1:3] TRUE FALSE FALSE

Preview of the elements
```

Naming and indexing of lists

The different elements of lists can be named:

```
l_2 <- list("numbers"=c(1,2), "letters"=c("a", "b"),
"logics"=c(TRUE, FALSE, FALSE))</pre>
```

• You can retrieve the names using names():

```
names(1_2)
```

You can subset the list using the names:

```
1_2["letters"]
```

And access the elements of the sublists with [[:

```
1_2[["letters"]]
```

• Alternatively use the shortcut \$: 1 2\$letters



Practical differences to atomic vectors

- There are two very important differences to atomic vectors:
 - Vectorisation does not work for lists
 - Indexing works differently for lists
- To illustrate the first issue compare:

To illustrate the latter:

```
typeof(l_[1])
typeof(l_[[1]])
```

• Lists are fundamental to more complex data structures we will encounter later

Final remarks on basic object types

- There are two "strange" data types: NA and NULL
- NA is used to represent absent elements of vectors
 - Happens frequently when vectors contain observations
 - Many functions behave differently when NAs are present (remember the na.rm argument!):

```
mean(c(1,2,NA)); mean(c(1,2,NA), na.rm = TRUE)
```

You test for NA using is.na():

```
is.na(c(1, 2, NA))
```

To check whether a vector contains missing values, use anyNA():

```
anyNA(c(1,2,NA))
```

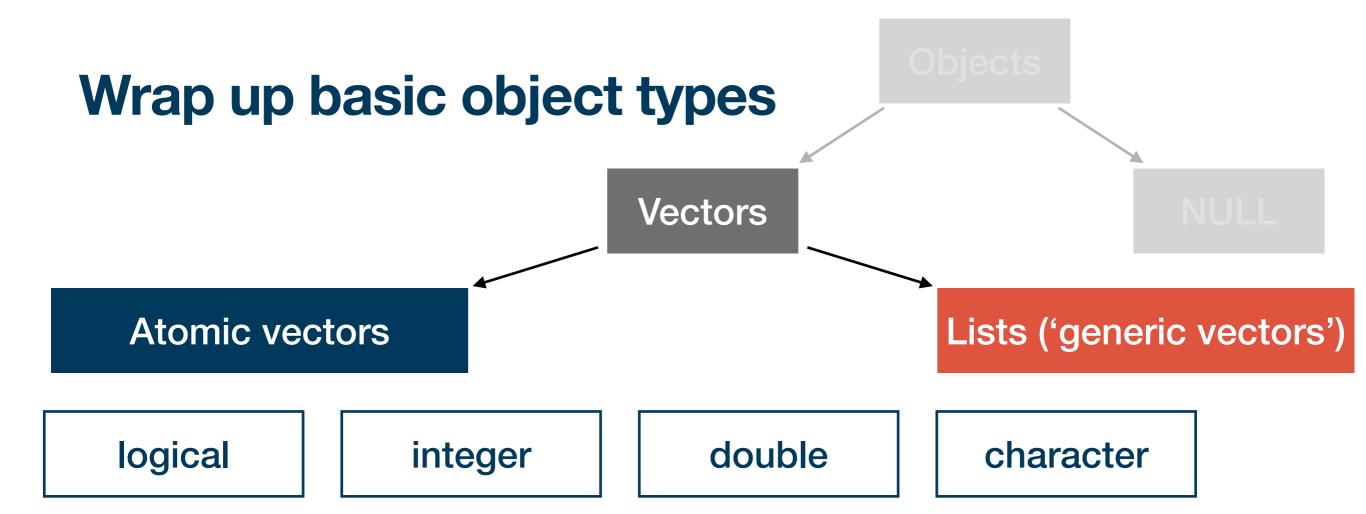


Final remarks on basic object types

- There are two "strange" data types: NA and NULL
- NULL is in fact a data type in itself, but in practice its best thought of as a vector of length zero:

```
c()
typeof(NULL)
length(NULL)
is.null(NULL)
```

- You might use NULL mainly in two instances:
 - Represent an empty vector of arbitrary type
 - Represent and absent vector (≠ NA, which represents absent elements of vectors)



- The central take-aways concern:
 - How to test for and transform these types: typeof(), is.*(), as.*()
 - How to index them: [, [[, \$
 - How to create typical instances: rep(), paste(), seq()
- We learned about vectorisation and its attractiveness in R
- We also encountered "strange" types such as NA, NULL and NaN



Break

Get together in groups of 2 to 5 people and talk about what were your main take aways, and resolve open questions!

Advanced object types in R



On more advanced object types

- While there are many object types in R, understanding the basics is key
 - These are by far the most common ones
 - All other object types are somehow 'built upon' the basic types by adding attributes
- Among the special types, two stand out in their prevalence:

Categorical data: factor

- Can also take a pre-specified number of values: levels
- Classical example: Male, Female,
 Diverse
- Created using the function factor()

Data frames: data.frame & tibble A kind of 'table' in which different variables are stored as vectors A table-like form of gender height Tibbles as a new va male 189 that "do less and co2 male 175 Created using data 3 male 180 tibble::tibble(166 5 female 150

Others that we will not cover here are, e.g., matrices, durations, or dates



Digression: some remarks on attributes

- To turn our basic object types into something more fancy we can give them attributes, one of which is called class
 - This changes their behaviour when functions are applied to them
 - Technically, adding a class attribute changes the class but not the type:

```
ff <- factor(c("F", "M", "M"), levels = c("F", "M", "D"))
typeof(ff)
class(ff)</pre>
```

The class factor is an integer with two attributes:

```
attributes(ff)
```

Not too important for us right now, but good to keep in mind!

Factors

- Factors are used to represent ordinal or categorial data
 - Elements of factors can take one out of several pre-specified values: levels
 - Factors are integers with the attributes levels and class
- We create factors using the function factor(), which takes a vector and an optional argument levels:

- Your turn:
 - What happens if we do not specify levels explicitly?
 - What happens if the vector contains elements not pre-specified as levels?

Factors

 Usually levels are not ordered, but for ordinal data you can use the argument ordered:

- There are some useful factor-specific functions such as table().
 - What does it do? Try it on f_1 and f_2!
- In general, its usually better to store categorial data as character, and only transform them to factors if necessary

Data frames

- Data frames are special lists of vectors where the length of each vector is equal!
 - → Most list operations also work for data.frames
- We usually represent data frames as tables:

```
gender height Names of the vectors

male 189
male 175 vector 1 &
male 180 vector 2
female 166
female 150
```

```
To create a data frame from scratch use
data.frame():

df_1 <- data.frame(
   "gender" = c(rep("male", 3), rep("female", 2)),
   "height" = c(189, 175, 180, 166, 150)
)</pre>
```

- To create a data frame from a list use as.data.frame()
- If you read in data into R, it almost always starts off as a data.frame
- How to transform them is the main subject of the sessions on data wrangling



Data frames and tibbles

- A modern version of the data.frame is the tibble (from the package tibble)
 - We will mostly use tibbles in this course, but make sure you familiarise yourself with the differences to the data.frame, which continues to be widespread (see the tutorial reading)
- To transform a data.frame (or a list) into a tibble, use tibble::as_tibble():
 tb_1 <- tibble::as_tibble(df_1)
- To extract single columns use the [or [[operators
 - What's the difference between the two?
 - How do you think you cam test for the type of a column vector?

Data frames and tibbles

- To get a quick overview about the content, use dplyr::glimpse() or head()
- A complete overview can be obtained via View()
- Data frames are among the most widely used data types
 - There different approaches of how to handle and transform them, each associated with an R dialect
 - We mainly rely on the tidyverse dialect, which is the easiest to learn and comprehend → built upon tibbles
 - Alternatives are the base (classical) and data.table (fastest) dialect, which mainly use data.frames and data.tables
- This is useful to keep in mind when searching help in the internet

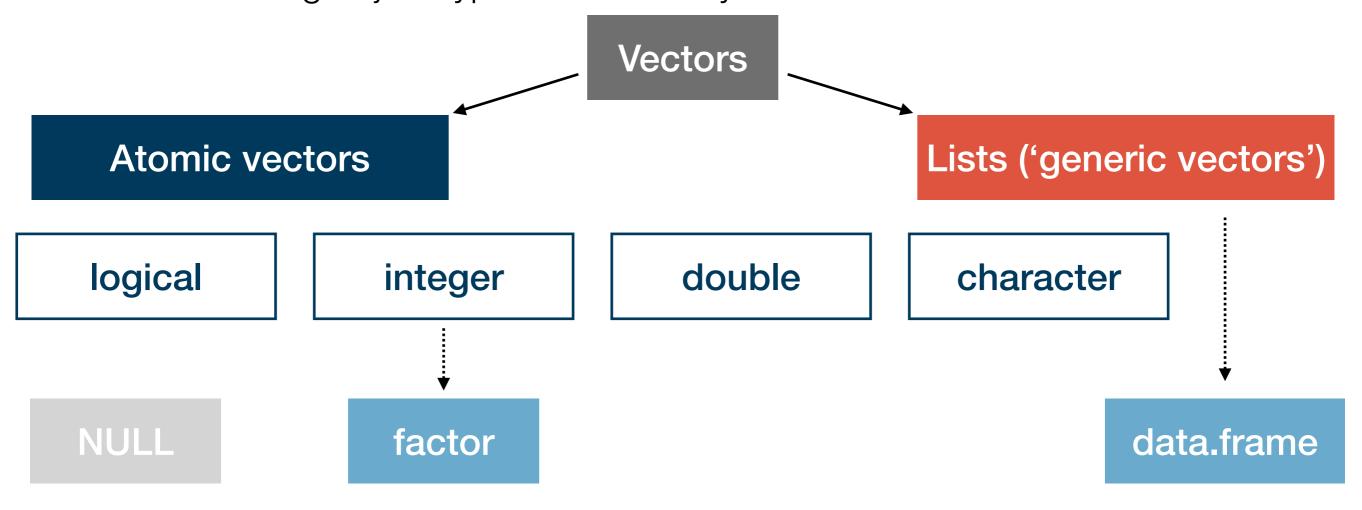


Quick exercises

- Create a factor with the levels "still", "medium" and "sparkling", and arbitrary instances of the three levels
- Transform this factor into a character vector
- Get the relative frequencies for "medium" of this factor
- Create a data frame with two columns, one called "nb" containing the numbers 1 to 5 as double, the other called "char" containing the numbers 6 to 10 as character
- Transform this data.frame into a tibble!
- Extract the second column of this tibble such that you have a vector

Summary and outlook

- This was the last session on the fundamentals of R
- We learned about the most important object types in R
- Functions do different things when applied to different objects → understanding object types is absolutely fundamental



Summary and outlook

- Next week we will learn how to visualise data that is stored in data frames
- This will be the first big intro session into data science fundamentals, succeeded by sessions on data wrangling and project management

Tasks until next week:

- 1. Fill in the quick feedback survey on Moodle
- 2. Read the **tutorials** posted on the course page
- Do the exercises provided on the course page and discuss problems and difficulties via the Moodle forum

