# The data.frame class

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### Learning Objectives

- understand the concept of a data.frame
- use sequences
- know how to access any element of a data.frame

#### What are data frames?

Data frames are the de facto data structure for most tabular data, and what we use for statistics and plotting.

A data frame is a collection of vectors of identical lengths. Each vector represents a column, and each vector can be of a different data type (e.g., characters, integers, factors). The str() function is useful to inspect the data types of the columns.

A data frame can be created by hand, but most commonly they are generated by the functions read.csv() or read.table(); in other words, when importing spreadsheets from your hard drive (or the web).

By default, when building or importing a data frame, the columns that contain characters (i.e., text) are coerced (=converted) into the factor data type. Depending on what you want to do with the data, you may want to keep these columns as character. To do so, read.csv() and read.table() have an argument called stringsAsFactors which can be set to FALSE:

```
some_data <- read.csv("data/some_file.csv", stringsAsFactors=FALSE)</pre>
```

You can also create a data frame manually with the function data.frame(). This function can also take the argument stringsAsFactors. Compare the output of these examples, and compare the difference between when the data are being read as character, and when they are being read as factor.

```
#> 'data.frame': 4 obs. of 3 variables:
#> $ animal: chr "dog" "cat" "sea cucumber" "sea urchin"
#> $ feel : chr "furry" "furry" "squishy" "spiny"
#> $ weight: num 45 8 1.1 0.8
```

#### Challenge

1. There are a few mistakes in this hand crafted data.frame, can you spot and fix them? Don't hesitate to experiment!

- 2. Can you predict the class for each of the columns in the following example? Check your guesses using str(country\_climate):
  - Are they what you expected? Why? Why not?
  - What would have been different if we had added stringsAsFactors = FALSE to this call?
  - What would you need to change to ensure that each column had the accurate data type?

r ## Answers ## \* missing quotations around the first names of the authors ## \* the year column is missing one value, 1859 (the year of publication of ## the origin of species)

r ## Answers ## \* 'country', 'climate', 'temperature', and 'northern\_hemisphere' are ## factors; 'has\_kangaroo' is numeric. ## \* using 'stringsAsFactors=FALSE' would have made them character instead of ## factors ## \* removing the quotes in temperature, northern\_hemisphere, and replacing 1 ## by TRUE in the 'has\_kangaroo' column would probably what was originally ## intended.

The automatic conversion of data type is sometimes a blessing, sometimes an annoyance. Be aware that it exists, learn the rules, and double check that data you import in R are of the correct type within your data frame. If not, use it to your advantage to detect mistakes that might have been introduced during data entry (a letter in a column that should only contain numbers for instance.).

## Inspecting data.frame Objects

We already saw how the functions head() and str() can be useful to check the content and the structure of a data.frame. Here is a non-exhaustive list of functions to get a sense of the content/structure of the data.

- Size:
  - dim() returns a vector with the number of rows in the first element, and the number of columns as the second element (the dim ensions of the object)
  - nrow() returns the number of rows
  - ncol() returns the number of columns

- Content:
  - head() shows the first 6 rowstail() shows the last 6 rows
- Names:
  - names() returns the column names (synonym of colnames() for data.frame objects)
  - rownames() returns the row names
- Summary:
  - str() structure of the object and information about the class, length and content of each column
  - summary() summary statistics for each column

Note: most of these functions are "generic", they can be used on other types of objects besides data.frame.

## Indexing, Sequences, and Subsetting

If we want to extract one or several values from a vector, we must provide one or several indices in square brackets. For instance:

```
animals <- c("mouse", "rat", "dog", "cat")
animals[2]

#> [1] "rat"
animals[c(3, 2)]

#> [1] "dog" "rat"
animals[2:4]

#> [1] "rat" "dog" "cat"

more_animals <- animals[c(1:3, 2:4)]
more_animals
#> [1] "mouse" "rat" "dog" "rat" "dog" "cat"
```

R indexes start at 1. Programming languages like Fortran, MATLAB, and R start counting at 1, because that's what human beings typically do. Languages in the C family (including C++, Java, Perl, and Python) count from 0 because that's simpler for computers to do.

: is a special function that creates numeric vectors of integers in increasing or decreasing order, test 1:10 and 10:1 for instance. The function seq() (for \_\_\_seq\_\_\_uence) can be used to create more complex patterns:

```
seq(1, 10, by=2)
```

```
#> [1] 1 3 5 7 9
```

```
seq(5, 10, length.out=3)

#> [1] 5.0 7.5 10.0

seq(50, by=5, length.out=10)

#> [1] 50 55 60 65 70 75 80 85 90 95

seq(1, 8, by=3) # sequence stops to stay below upper limit
```

#> [1] 1 4 7

Our temperature data frame has rows and columns (it has 2 dimensions), if we want to extract some specific data from it, we need to specify the "coordinates" we want from it. Row numbers come first, followed by column numbers.

```
temperature[1]  # first column in the data frame
temperature[1, 1]  # first element in the first column of the data frame
temperature[1, 6]  # first element in the 6th column
temperature[1:3, 7]  # first three elements in the 7th column
temperature[3, ]  # the 3rd element for all columns
temperature[, 8]  # the entire 8th column
head_temperature <- temperature[1:6, ] # temperature[1:6, ] is equivalent to head(temperature)</pre>
```

As well as using numeric values to subset a data.frame (or matrix), columns can be called by name, using one the three following notations:

```
temperature[, "City"]
temperature[["City"]]
temperature$City
```

For our purposes, these three notations are equivalent. However, the last one with the \$ does partial matching on the name. So you could also select the column "month" by typing temperature\$m. It's a shortcut, as with all shortcuts, they can have dangerous consequences, and are best avoided. Besides, with auto-completion in RStudio, you rarely have to type more than a few characters to get the full and correct column name.

### Challenge

1. The function nrow() on a data.frame returns the number of rows. Use it, in conjuction with seq() to create a new data.frame called surveys\_by\_10 that includes every 10th row of the survey data frame starting at row 10 (10, 20, 30, ...)

```
### The function `nrow()` on a `data.frame` returns the number of
### rows. Use it, in conjuction with `seq()` to create a new
### `data.frame` called `temperature_by_10` that includes every 10th row
### of the temperature data frame starting at row 10 (10, 20, 30, ...)
```

```
## Answer
temperature_by_10 <- temperature[seq(10, nrow(temperature), by=10), ]</pre>
```

#### Conditional subsetting

Besides using the index position of an element in a vector to extract its value as we saw earlier, we can also use logical vectors:

```
animals <- c("mouse", "rat", "dog", "cat")
animals[c(TRUE, FALSE, TRUE, TRUE)]</pre>
```

But typically, those logical vectors are not typed by hand but the result of a logical test:

```
animals != "rat"
animals[animals != "rat"]
animals[animals == "cat"]
```

If you can combine multiple tests using & (both conditions are true, AND) or | (at least one of the conditions if true, OR):

```
animals[animals == "cat" & animals == "rat"] # returns nothing
animals[animals == "cat" | animals == "rat"] # returns both rat and cat
```

If you are trying to combine many conditions, it can become tedious to type. The function %in% allows you to test if a value if found in a vector:

```
animals %in% c("rat", "cat", "dog", "duck")
animals[animals %in% c("rat", "cat", "dog", "duck")]
```

In addition to testing equalities, you can also test whether the elements of your vector are less than or greater than a given value:

```
dates <- c(1960, 1963, 1974, 2015, 2016)
dates >= 1974
dates[dates >= 1974]
dates[dates > 1970 & dates <= 2015]
dates[dates < 1975 | dates > 2016]
```

### Challenge

• Can you figure out why "four" > "five" returns TRUE?

```
# * Can you figure out why `"four" > "five"` returns `TRUE`?

## Answers
## * When using ">" or "<" on strings, R compares their alphabetical order. Here
## "four" comes after "five", and therefore is "greater than" it.</pre>
```