

How to load and save data in various formats

In this subchapter, we will present how to load tabular data from various sources. We will discuss all kinds of formats, since we understand that people who use R have various needs. For typical cases, you will only need to read the first three sections that explain how to load data from packages, text files and Excel files.

Generally speaking, data sets can have various structures. They can be expressed as graphs, trees, they may describe photos or multimedia files. All of these structures can be loaded into R and analysed. Most of the time, however, the data sets that we work with have a tabular structure with columns and rows clearly marked. We will put focus on this sort of structure in the following subchapter.

Loading data from text files

Loading data from packages is the easiest case. You only need to turn the given package on using `library()`, and then use `data()` to load a given file. A single package can store multiple datasets, which is why packages are often used as storages of useful data.

One of such packages is the `PogromcyDanych` package (Polish for Data Busters). It contains various datasets which we use in the present book, such as `cats_birds`. Once we install the `PogromcyDanych` package, we only need two lines of code to use this dataset:

```
library("PogromcyDanych")
cats_birds <- read.csv("/home/krz/cats_birds.csv", sep=";", dec=",")
```

A new symbol for the dataset should appear in the `Environment` of RStudio. We can see the first few rows using the `head()` function.

```
head(cats_birds)
```

##	species	weight	length	velocity	habitat	lifespan	team
## 1	Tiger	300	2.5	60	Asia	25	Cat
## 2	Lion	200	2.0	80	Africa	29	Cat
## 3	Jaguar	100	1.7	90	America	15	Cat
## 4	Puma	80	1.7	70	America	13	Cat
## 5	Panthera	70	1.4	85	Asia	21	Cat
## 6	Cheetah	60	1.4	115	Africa	12	Cat

The `cats_birds` dataset contains 7 columns that describe various parameters of sample animals. To show the list of all datasets in a given package, we can also use the `data()` function:

```
data(package="PogromcyDanych")
```

Loading data from text files

Text files are a very popular format to store data. Their contents can be displayed in any text file, or in RStudio. They usually have the `.txt`, `.csv` (comma separated values) or `.tsv` (tab separated values) extension. In such files, subsequent column values are separated with some sort of separators, such as tabs, commas, semicolons, or other characters.

We will now analyse an example data set available on-line at http://biecek.pl/R/dane/cats_birds.csv. The file contents is presented below:

```
species;weight;length;velocity;habitat;lifespan;team
Tiger;300;2,5;60;Asia;25;Cat
Lion;200;2,0;80;Africa;29;Cat
Jaguar;100;1,7;90;America;15;Cat
Puma;80;1,7;70;America;13;Cat
```

```

Panthera;70;1,4;85;Asia;21;Cat
Cheetah;60;1,4;115;Africa;12;Cat
Snow leopard;50;1,3;65;Asia;18;Cat
Swift;0,05;0,2;170;Eurasia;20;Bird
Ostrich;150;2,5;70;Africa;45;Bird
Golden eagle;5;0,9;160;North;20;Bird
Peregrine falcon;0,7;0,5;110;North;15;Bird
Gerfalcon;2;0,7;100;North;20;Bird
Albatross;4;0,8;120;South;50;Bird

```

We can see that the file contains a table of values separated with semicolons. The first row is the header with column (variable) names:

```
species;weight;lenght;velocity;habitat;lifespan;team
```

Further rows contain the values of these variables. A description of the logical structure of this file is presented in Figure @ref(fig:wy216).

species	weight	lenght	velocity	habitat	lifespan	team
Tiger	300	2,5	60	Asia	25	Cat
Lion	200	2	80	Africa	29	Cat
Jaguar	100	1,7	90	America	15	Cat
Puma	80	1,7	70	America	13	Cat
Panthera	70	1,4	85	Asia	21	Cat
Cheetah	60	1,4	115	Africa	12	Cat
Snow	50	1,3	65	Asia	18	Cat
Swift	0,05	0,2	170	Euroasia	20	Bird
Ostrich	150	2,5	70	Africa	45	Bird
Golden eagle	5	0,9	160	North	20	Bird
Peregrine falcon	0,7	0,5	110	North	15	Bird
Gerfalcon	2	0,7	100	North	20	Bird
Albatross	4	0,8	120	South	50	Bird

Figure 1: The logical structure of the ‘cats-birds.csv’ file.

To load data in this format into R, we can use the `read.table()` function. It has many parameters which are described in detail later. In the example below, the table of values that we loaded is assigned to the `cats_birds` symbol using the `<-` symbol.

Variable name	Assignment operator <- or =	Function that loads data						
cats_birds	<-	read.table("cats_birds.csv", header=TRUE, sep=";")						
		species	weight	lenght	velocity	habitat	lifespan	team
		Tiger	300	2,5	60	Asia	25	Cat
		Lion	200	2	80	Africa	29	Cat
		Jaguar	100	1,7	90	America	15	Cat
		Puma	80	1,7	70	America	13	Cat
		Panthera	70	1,4	85	Asia	21	Cat
		Cheetah	60	1,4	115	Africa	12	Cat
		Snow	50	1,3	65	Asia	18	Cat
		Swift	0,05	0,2	170	Euroasia	20	Bird
		Ostrich	150	2,5	70	Africa	45	Bird
		Golden eagle	5	0,9	160	North	20	Bird
		Peregrine falcon	0,7	0,5	110	North	15	Bird
		Gerfalcon	2	0,7	100	North	20	Bird
		Albatross	4	0,8	120	South	50	Bird

Figure 2: Logical structure of reading data from a file.

```
cats_birds <- read.table(file="http://biecek.pl/R/cats_birds.csv",
```

```
sep=";", dec=".", header=TRUE)
```

Data can be loaded from a file located on a hard drive, or downloaded directly from the Internet. In this case, we need to specify the following arguments:

- `file="http://biecek.pl/R/cats_birds.csv"`, i.e. text file path,
- `sep=";"`, which defines the separator character,
- `dec="."`, which defines the decimal point; usually a dot, but can also be a comma, as shown in this file,
- `header=TRUE`, which informs that the first row contains headers and not values.

To find out whether data has been loaded correctly, we can again use the `head()` function. Alternatively, you can also use the **Environment** tab in RStudio, as shown in Figure @ref(fig:wy218).

```
head(cats_birds)
```

```
##   species weight length velocity habitat lifespan team
## 1   Tiger   300    2.5      60    Asia      25   Cat
## 2    Lion   200    2.0      80   Africa      29   Cat
## 3  Jaguar   100    1.7      90 America      15   Cat
## 4    Puma    80    1.7      70 America      13   Cat
## 5 Panthera   70    1.4      85    Asia      21   Cat
## 6 Cheetah   60    1.4     115   Africa      12   Cat
```

	species	weight	length	velocity	habitat	lifespan
1	Tiger	300	2.5	60	Asia	25
2	Lion	200	2	80	Africa	29
3	Jaguar	100	1.7	90	America	15
4	Puma	80	1.7	70	America	13
5	Panthera	70	1.4	85	Asia	21
6	Cheetah	60	1.4	115	Africa	12
7	Snow leopard	50	1.3	65	Asia	18
8	Swift	0.05	0.2	170	Eurasia	20
9	Ostrich	150	2.5	70	Africa	45
10	Golden eagle	5	0.9	160	North	20
11	Peregrine falcon	0.7	0.5	110	North	15
12	Gyr Falcon	2	0.7	100	North	20
13	Albatross	4	0.8	120	South	50

Figure 3: Table preview with data in RStudio.

Loading big files with `read.table()` can take a lot of time. In such cases, you may consider using the `data.table::fread()` function, which is much faster and has a similar number of parameters.

`read.table()` in detail

Unfortunately, there is no single universal format to store data in text files. Separators, decimal points, comment lines or strings can be defined with various characters.

The `read.table()` function is able to load data written in virtually any format. It is, however, only possible with a major number of additional options. The declaration of `read.table()` is as follows:

```
read.table(file, header=FALSE, sep="", quote="\'", dec=".",
           row.names, col.names, as.is=!stringsAsFactors,
           na.strings="NA", colClasses=NA, nrows=-1, skip=0,
           check.names=T, fill=!blank.lines.skip, strip.white=F,
           blank.lines.skip=T, comment.char="#", allowEscapes=F,
```

```
flush=FALSE, stringsAsFactors, encoding = "unknown")
```

In the `utils` package, there are four other functions that work in the same way, but have other default values.

```
read.csv(file, header=TRUE, sep=",", quote="\"", dec=".",
         fill=TRUE, comment.char="", ...)
read.csv2(file, header=TRUE, sep=";", quote="\"", dec=".",
          fill=TRUE, comment.char="", ...)
read.delim(file, header=TRUE, sep="\t", quote="\"", dec=".",
           fill=TRUE, comment.char="", ...)
read.delim2(file, header=TRUE, sep="\t", quote="\"", dec=".",
            fill=TRUE, comment.char="", ...)
```

If we saved the data in the CSV format using the Polish version of Excel, semicolons will be used as separators, and commas as decimal points. In such cases, the `read.csv2()` function is most suitable. If we use the English version of Excel, dots will be used as decimal points, and the `read.csv()` function will be a better solution.

Table @ref(tab:tab02) presents the descriptions of `read.table()` arguments. The English version of these descriptions can be found in the help file: `?read.table`. As can be seen, there are many arguments, and we suggest you read about each of them at least once.

Table 1: Arguments of ‘`read.table()`’

Function	Description
file	String value. It is the only obligatory argument which must be explicitly provided. Its value is the file path where the data is stored. We can provide a local file, or a file from the Internet by providing the URL that specifies the server and directory of the file. This field can also specify a connection or a socket. If we provide "clipboard" instead of a file name, data will be read from the system clipboard, which is a convenient way of loading data from various MS Office programs.
header	Logical value. If it is <code>TRUE</code> , the first argument will be treated as a list of columns. If this argument is not provided, and the first line contains one field less than other lines, R will automatically treat this line as the header.
sep	String value. The given string will be treated as the field separator. The default "" value treats each whitespace (single space, multiple spaces, tabulator, new line) as a separator.
quote	String value. It defines the character that represents the decimal point. If data is saved according to Polish standards, the comma sign "," is the decimal point.
row.names	A vector of strings or a single number. If a vector of strings is provided, it will be treated as the names of subsequent rows. If this argument is a number, it will be treated as the ordinal number of the column in the loaded file that contains row names. If the data contains a header (i.e. <code>header=TRUE</code>) and the first line has one field less than other lines, then the first line is automatically treated as a vector with row names.
col.names	Vector of strings. Column names can be specified with this argument. If it is not provided, and the file does not contain a header, then subsequent column names start with the letter "V", followed by increasing integer numbers.
nrows	Numerical field. Specifies the maximal number of rows to be read.
skip	Numerical value. Specifies the number of initial lines that should be ignored when reading the file.
check.names	Logical value. Specifies whether column names should be checked for correctness (i.e. without illegal characters or repetitions).

Table 1: Arguments of ‘read.table()’ (*continued*)

Function	Description
as.is	Vector of logical or numerical variables. By default, each field which is not converted into a real number or logical value is converted into a factor type variable. If the as.is parameter is a vector of logical values, then it indicates which columns can be converted into the factor type, and which cannot. A vector of numbers, instead, is treated as indexes of columns which should not be converted.
na.strings	Vector of strings. Indicates which values should be treated as NA, i.e. which values are used for missing observations. Additionally, empty fields in columns with numerical or logical values are marked as missing observations.
colClasses	Vector of characters. Each character specifies the type of a single variable (column). Possible values are: (1) NA - default value, signifies automatic conversion, (2) NULL - a column will be skipped, (3) one of atomic classes (logical, integer, numeric, complex, character, raw, factor, Date or POSIXct).
fill	Logical value. If subsequent lines have different numbers of values, and fill=FALSE (default value), then read.table() will throw an error. If fill=TRUE , shorter lines will be complemented with NA values to get the same line length.
blank.lines.s	Logical value. When TRUE , empty lines will be skipped while reading the file.
comment.char	Character value. This character is treated as a comment character. All remaining characters in a given line after this character will be ignored.
allowEscapes	Logical value. When TRUE , special escape characters are recognised. For instance, or are changed into tabulators or new line characters, respectively. FALSE (default value) means the text will be read literally, without any escape characters.
stringsAsFactors	Logical value. If TRUE , strings will be converted into factor types. If FALSE , they will be represented as strings. The default value is TRUE .

Saving data into text files

write.table() function

The **write.table()** function saves the values of a variable (atomic variable, vector, matrix or data frame) into a text file with a given formatting style. Below, we present the full declaration of this function:

```
write.table(x, file="", append=FALSE, quote=TRUE, sep=" ",
           eol="\n", na="NA", dec=".", row.names=TRUE,
           col.names=TRUE, qmethod=c("escape","double"))
```

Similarly to **read.table()**, the **utils** package features two functions that work in the same way as **write.table()**, but use other parameterisation of default arguments. Both functions are used to save data in the csv format.

```
write.csv(x, file = "", sep = ",", dec=".", ...)
write.csv2(x, file = "", sep = ";", dec=".", ...)
```

Table @ref(tab:tab03) describes all parameters of **write.table()** based on the help file: **?write.table**.

Table 2: Arguments of ‘write.table()’

Function	Description
x	Object to save. Obligatory field. This object is usually a matrix or a data frame.
file	String value. Represents the path to the file in which the data should be written. This field can also represent a connection or a socket, just like <code>read.table()</code> . Similarly, if we provide "clipboard" instead, the data will be saved into the clipboard. The " " value will print the object to the console.
qmethod	String value. Indicates how special characters should be marked. "escape" value is a C-style marking (with a backslash sign), while "double" value means a double backslash should be used instead.
sep	String value. This string will be treated as the separator of subsequent fields (the separator does not need to be a single character).
eol	String value. This string will be used as a line ending string. In Windows, the default value is <code>eol=" "</code> .
quote	Logical value. TRUE means that factor types and strings will be surrounded with quotes ‘”’.
dec	String value. This field defines the decimal point. By default, it is a "." (dot). If the data is written in accordance with Polish standards, then the decimal point should be a "," (comma).
row.names	Vector of strings or logical value. If a vector of strings is provided, it will be used to name subsequent rows. If a logical value is provided, it will specify whether row names should be written or not.
col.names	Vector of strings or logical value. If a vector of strings is provided, it will be used to name subsequent columns. If a logical value is provided, it will specify whether column names should be written or not.

Below, we show a few examples of how to save a vector, number or a data frame into a file.

This instruction prints a vector of numbers onto the screen instead of saving it into a file:

```
write.table(runif(100),"")
```

This instruction will write a vector into the clipboard:

```
write.table(1:100,"clipboard",col.names=FALSE, row.names=FALSE)
```

This instruction will write a string into a text file:

```
write.table("Monday","c:/worstWeekday.txt")
```

This instruction will save data in the CSV format into a file:

```
write.csv2(iris, file="forExcel.csv")
```

Writing big objects of **data.frame** type into text files may be time-consuming. This happens due to the fact that formatting is switched between various columns that can present data of various types. R checks the type of each column and row, and formats the values based on their types. To speed up the process, it is sometimes better to convert the data into a matrix using `as.matrix()`.

Loading and saving JSON data

Previously, we mainly discussed tabular data, but not all data can be saved as tables.

A universal and very flexible format used to transport various data sets is JSON (*JavaScript Object Notation*). It is frequently found in solutions that use Internet communication, such as REST services.

JSON is a text format (which means it is human-readable), describes data with a hierarchical structure (which is a very flexible solution), and is lightweight (does not take much space).

Below, we present a sample JSON file. It describes a two-element list whose first value is a string, and the second value is a table.

```
{ "source": "Przewodnik",
  "indicators": [
    {"year": "2010", "value": 15.5},
    {"year": "2011", "value": 10.1},
    {"year": "2012", "value": 12.2}
  ] }
```

In R, there are a few packages to read or write JSON files. The three most popular ones are `RJSONIO`, `rjson` and `jsonlite`. Each of them contains a `toJSON()` function which converts data into JSON, and a `fromJSON()` function which loads data from JSON.

The most stable option is the `jsonlite` package. Below, we only present some examples for this package.

From JSON to R: The data we load is initially a list. The structure is simplified into a data frame or a vector, where possible.

```
library("jsonlite")
fromJSON('{ "source": "Przewodnik", "indicators": [{"year": "2010", "value": 15.5},
  {"year": "2011", "value": 10.1}, {"year": "2012", "value": 12.2}]}')
```

```
## $source
## [1] "Przewodnik"
##
## $indicators
##   year value
## 1 2010  15.5
## 2 2011  10.1
## 3 2012  12.2
```

From R to JSON: The example below shows how to convert the first two rows of the `cats_birds` data frame:

```
toJSON(cats_birds[1:2,])
```

```
## [{"species":"Tiger","weight":300,"length":2.5,"velocity":60,"habitat":"Asia","lifespan":25,"team":"C
```