

R - Packages

R packages are a collection of R functions, compiled code and sample data. They are stored under a directory called "**library**" in the R environment. By default, R installs a set of packages during installation. More packages are added later, when they are needed for some specific purpose. When we start the R console, only the default packages are available by default. Other packages which are already installed have to be loaded explicitly to be used by the R program that is going to use them.

All the packages available in R language are listed at R Packages.

Below is a list of commands to be used to check, verify and use the R packages.

Check Available R Packages

Example: Get library locations containing R packages

```
> .libPaths()  
[1] "C:/Users/hp/AppData/Local/R/win-library/4.4"  
[2] "C:/Program Files/R/R-4.4.0/library"
```

Example: Get the list of all the packages installed:

```
> library()
```

When we execute the above code, it produces the following result. It may vary depending on the local settings of your pc. In our case, the results will be:

```
Packages in library 'C:/Users/hp/AppData/Local/R/win-library/4.4':
```

abind	Combine Multidimensional Arrays
AnnotationDbi	Manipulation of SQLite-based annotations in Bioconductor
askpass	Password Entry Utilities for R, Git, and SSH
base64enc	Tools for base64 encoding
BH	Boost C++ Header Files
Biobase	Biobase: Base functions for Bioconductor
BiocGenerics	S4 generic functions used in Bioconductor
BiocIO	Standard Input and Output for

Get all packages currently loaded in the R environment

```
> search()
[1] ".GlobalEnv"          "tools:rstudio"       "package:stats"
[4] "package:graphics"    "package:grDevices"   "package:utils"
[7] "package:datasets"    "package:methods"     "AutoLoads"
[10] "package:base"
```

When we execute the above code, it produces the above result. It may vary depending on the local settings of your pc.

Install a New Package

There are two ways to add new R packages. One is installing directly from the CRAN (Comprehensive R Archive Network) directory and another is downloading the package to your local system and installing it manually.

Install directly from CRAN

The following command gets the packages directly from CRAN webpage and installs the package in the R environment. You may be prompted to choose a nearest mirror. Choose the one appropriate to your location.

install.packages("Package Name")

```
> # Install the package named "XML".
> install.packages("XML")
```

WARNING: Rtools is required to build R packages but is not currently installed. Please download and install the appropriate version of Rtools before proceeding:

<https://cran.rstudio.com/bin/windows/Rtools/>
Installing package into 'C:/Users/hp/AppData/Local/R/win-library/4.4'
(as 'lib' is unspecified)

```
There is a binary version available but the source version
is later:
      binary      source needs_compilation
XML 3.99-0.16.1 3.99-0.17                TRUE
```

```
Binaries will be installed
trying URL 'https://cran.rstudio.com/bin/windows/contrib/4.4/XML_3.99-0.16.1.zip'
Content type 'application/zip' length 3103340 bytes (3.0 MB)
```

downloaded 3.0 MB

package 'XML' successfully unpacked and MD5 sums checked

The downloaded binary packages are in
C:\Users\Public\Documents\iSkysoft\CreatorTemp\RtmpweH194\
downloaded_packages

Install package manually

Go to the link R Packages to download the package needed. Save the package as a **.zip** file in a suitable location in the local system.

Now you can run the following command to install this package in the R environment.

```
install.packages(file_name_with_path, repos = NULL, type = "source")
```

Example:

```
> # Install the package named "XML"  
> install.packages("E:/XML_3.98-1.3.zip", repos = NULL, type = "source")
```

Load Package or Library

Before a package can be used in the code, it must be loaded to the current R environment. You also need to load a package that is already installed previously but not available in the current environment. A package is loaded using the following command:

```
> # Loading library called "xlsx"  
> library("xlsx")
```

Viewing the contents of loaded library

In R, you can view the contents of a library (also known as a package) using the `ls()` function or by exploring the package's documentation. Here's how you can do it:

```
> ls("package:xlsx")
```

[1]	"addAutoFilter"	"addDataFrame"
[3]	"addHyperlink"	"addMergedRegion"
[5]	"addPicture"	"Alignment"
[7]	"autoSizeColumn"	"Border"
[9]	"BORDER_STYLES_"	"CB.setBorder"
[11]	"CB.setColData"	"CB.setFill"
[13]	"CB.setFont"	"CB.setMatrixData"
[15]	"CB.setRowData"	"CELL_STYLES_"
[17]	"CellBlock"	"CellProtection"
[19]	"CellStyle"	"createCell"
[21]	"createCellComment"	"createFreezePane"
[23]	"createRange"	"createRow"
[25]	"createSheet"	"createSplitPane"
[27]	"createWorkbook"	"DataFormat"
[29]	"Fill"	"FILL_STYLES_"
[31]	"Font"	"forceFormulaRefresh"
[33]	"forcePivotTableRefresh"	"get_java_tmp_dir"
[35]	"getCellComment"	"getCells"
[37]	"getCellStyle"	"getCellValue"
[39]	"getRanges"	"getRows"
[41]	"getSheets"	"HALIGN_STYLES_"
[43]	"INDEXED_COLORS_"	"is.Alignment"
[45]	"is.Border"	"is.CellBlock"
[47]	"is.CellProtection"	"is.CellStyle"
[49]	"is.DataFormat"	"is.Fill"
[51]	"is.Font"	"loadWorkbook"
[53]	"printSetup"	"read.xlsx"
[55]	"read.xlsx2"	"readColumns"
[57]	"readRange"	"readRows"
[59]	"removeCellComment"	"removeMergedRegion"
[61]	"removeRow"	"removeSheet"
[63]	"saveWorkbook"	"set_java_tmp_dir"
[65]	"setCellStyle"	"setCellValue"
[67]	"setColumnWidth"	"setPrintArea"
[69]	"setRowHeight"	"setZoom"
[71]	"VALIGN_STYLES_"	"write.xlsx"
[73]	"write.xlsx2"	

You can also explore the documentation of a package to see its contents by Using **help()** or **?** to Explore Documentation in the help tab (down right window of RStudio).

```
> # Explore the documentation of package in help tab
> help(package = "xlsx")
>
> # Using ? to get help (this is used to get help of everyt
hing)
> ? "xlsx"
```

Detaching a Package

you can **detach** a package from the search path, which effectively removes its functions and datasets from your current R session. Use the **detach()** function to remove a package from the search path:

```
> # viewing current loaded packages
>
> search()
[1] ".GlobalEnv"          "package:xlsx"          "tools:rstudio"
[4] "package:stats"        "package:graphics"      "package:grDevices"
[7] "package:utils"        "package:datasets"      "package:methods"
[10] "AutoLoads"           "package:base"
>
> # Detach "xlsx" package
> detach("package:xlsx", unload = TRUE)
>
> # viewing loaded packages after detaching "xlsx" package
>
> search()
[1] ".GlobalEnv"          "tools:rstudio"         "package:stats"
[4] "package:graphics"    "package:grDevices"     "package:utils"
[7] "package:datasets"    "package:methods"       "AutoLoads"
[10] "package:base"
```

- "package:dplyr": Specifies the package to detach.
- unload = TRUE: Ensures the package is unloaded from memory (optional but recommended).

R Data Interferences

CSV files

In R, we can read data from files stored outside the R environment. We can also write data into files which will be stored and accessed by the operating system. R can read and write into various file formats like csv, excel, xml etc.

In this section we will learn to read data from a csv file and then write data into a csv file. **The file should be present in current working directory so**

that R can read it, elsewhere you have to specify path . Of course we can also set our own directory and read files from there.

Getting and Setting the Working Directory

You can check which directory the R workspace is pointing to using the **getwd()** function. You can also set a new working directory using **setwd()** function.

Example:

```
> # Get and print current working directory.  
> print(getwd())
```

```
"C:/Users/hp/R"
```

```
> # Set current working directory.  
> setwd("C:/Users/hp/OneDrive/Documents")  
>  
> # Get and print current working directory.  
> print(getwd())  
"C:/Users/hp/R"
```

This result depends on your OS and your current directory where you are working.

Input as CSV File

The csv file is a text file in which the values in the columns are separated by a comma. Let's consider the following data present in the file named **input.csv**.

You can create this file using windows notepad by copying and pasting this data. Save the file as **input.csv** using the save As All files(*.*) option in notepad.

```
id,name,salary,start_date,dept  
1,Rick,623.3,2012-01-01,IT  
2,Dan,515.2,2013-09-23,Operations  
3,Michelle,611,2014-11-15,IT  
4,Ryan,729,2014-05-11,HR  
5,Gary,843.25,2015-03-27,Finance
```

```
6,Nina,578,2013-05-21,IT
7,Simon,632.8,2013-07-30,Operations
8,Guru,722.5,2014-06-17,Finance
```

Reading a CSV File

Following is a simple example of **read.csv()** function to read a CSV file available in your current working directory:

```
> data <- read.csv("input.csv")
> print(data)
```

	id	name	salary	start_date	dept
1	1	Rick	623.30	2012-01-01	IT
2	2	Dan	515.20	2013-09-23	Operations
3	3	Michelle	611.00	2014-11-15	IT
4	4	Ryan	729.00	2014-05-11	HR
5	5	Gary	843.25	2015-03-27	Finance
6	6	Nina	578.00	2013-05-21	IT
7	7	Simon	632.80	2013-07-30	Operations
8	8	Guru	722.50	2014-06-17	Finance

Analyzing the CSV File

By default the **read.csv()** function gives the output as a data frame. This can be easily checked as follows. Also we can check the number of columns and rows.

Example:

```
> data <- read.csv("input.csv")
>
> print(is.data.frame(data))
[1] TRUE
>
> print(ncol(data))
[1] 5
>
> print(nrow(data))
[1] 8
```

Once we read data in a data frame, we can apply all the functions applicable to data frames as explained in subsequent section.

Get the maximum salary:

```
> # Create a data frame.
> data <- read.csv("input1.csv")
>
> # Get the max salary from data frame.
> sal <- max(data$salary)
> print(sal)

[1] 843.25
```

Writing into a CSV File

R can create csv file from existing data frame. The **write.csv()** function is used to create the csv file. This file gets created in the working directory.

```
> # Create a data frame.
> data <- read.csv("input.csv")
> retval <- subset(data, start_date >= "2014-01-01")
>
> # write filtered data into a new file.
> write.csv(retval, "output.csv")
> newdata <- read.csv("output.csv")
> print(newdata)
```

	X	id	name	salary	start_date	dept
1	3	3	Michelle	611.00	2014-11-15	IT
2	4	4	Ryan	729.00	2014-05-11	HR
3	5	5	Gary	843.25	2015-03-27	Finance
4	8	8	Guru	722.50	2014-06-17	Finance

R Data Interferences**Excel files**

Microsoft Excel is the most widely used spreadsheet program which stores data in the .xls or .xlsx format. R can read directly from these files using some excel specific packages. The popular R packages for reading and writing Excel files are:

Reading Excel Files:

- **readxl:** This is a popular and user-friendly choice for reading data from both .xls and .xlsx Excel files into R data frames. It's known for its

simplicity and lack of external dependencies, making it work seamlessly across different operating systems.

Example: reading excel file using **read_excel()** function of readxl package

```
> # Loading the required library "readxl"
> library(readxl)
>
> input <- read_excel("input.xlsx", sheet = "sheet1")
> view(input) # this command optional, view the table in editing window (upper left part of R Studio)
> input
# A tibble: 8 × 5
   id name      salary start_date      dept
  <dbl> <chr>    <dbl> <dtm>    <chr>
1     1 Rick      623. 2002-01-01 00:00:00 012      IT
2     2 Dan       515. 2013-09-23 00:00:00 Operations
3     3 Michelle  611. 2014-11-15 00:00:00 IT
4     4 Ryan      729. 2014-05-11 00:00:00 HR
5     5 Gary      43.2 2015-03-27 00:00:00 Finance
6     6 Nina      578. 2013-05-21 00:00:00 IT
7     7 Simon     633. 2013-07-30 00:00:00 Operations
8     8 Guru      722. 2014-06-17 00:00:00 Finance
>
> # exploring specific field "name" of read table
> name <- input$name
> name
[1] "Rick"      "Dan"      "Michelle" "Ryan"      "Gary"
[6] "Nina"      "Simon"    "Guru"
> salary <- input$salary
> salary
[1] 623.30 515.20 611.00 729.00 43.25 578.00 632.80 722.50
>
> # If we want to get the sum of salaries
> Sum <- sum(input$salary)
> Sum
[1] 4455.05
> # we can convert the loaded data to a data frame
> df <- data.frame(input)
> df
   id      name salary start_date      dept
1   1      Rick 623.30 2002-01-01 012      IT
2   2        Dan 515.20 2013-09-23      Operations
3   3  Michelle 611.00 2014-11-15      IT
4   4      Ryan 729.00 2014-05-11      HR
5   5      Gary  43.25 2015-03-27      Finance
6   6      Nina 578.00 2013-05-21      IT
7   7     Simon 632.80 2013-07-30      Operations
8   8      Guru 722.50 2014-06-17      Finance
>
```

```
> #exploring selective fields
> df$name
[1] "Rick"      "Dan"      "Michelle" "Ryan"      "Gary"
[6] "Nina"      "Simon"    "Guru"
>
> # Summarize data
> summary(df)
      id          name          salary
Min.   :1.00   Length:8   Min.    : 43.25
1st Qu.:2.75   Class :character 1st Qu.:562.30
Median :4.50   Mode  :character Median :617.15
Mean   :4.50                Mean  :556.88
3rd Qu.:6.25                3rd Qu.:655.23
Max.   :8.00                Max.   :729.00
 start_date          dept
Min.   :2002-01-01 00:00:00 Length:8
1st Qu.:2013-07-12 12:00:00 Class :character
Median :2014-01-16 00:00:00 Mode  :character
Mean   :2012-09-13 06:00:00
3rd Qu.:2014-07-24 18:00:00
Max.   :2015-03-27 00:00:00
```

We can also, import excel file

Writing Excel Files:

- **writexl**: This package complements readxl and excels (pun intended) at writing R data frames to new .xlsx Excel files. It offers a straightforward approach without requiring additional dependencies.

Example: using **write_excel()** function of “writexl” package to generate excel file in working directory (if not you have to specify the path)

```
> # Load the “writexl” package
> library(writexl)
>
> # Create a sample data frame
> df <- data.frame(
+   Name = c("Alice", "Bob", "Charlie"),
+   Age = c(25, 30, 35),
+   Salary = c(50000, 60000, 70000)
+ )
>
> # write the data frame to an Excel file
> write_excel(df, path = "example.xlsx")
```