



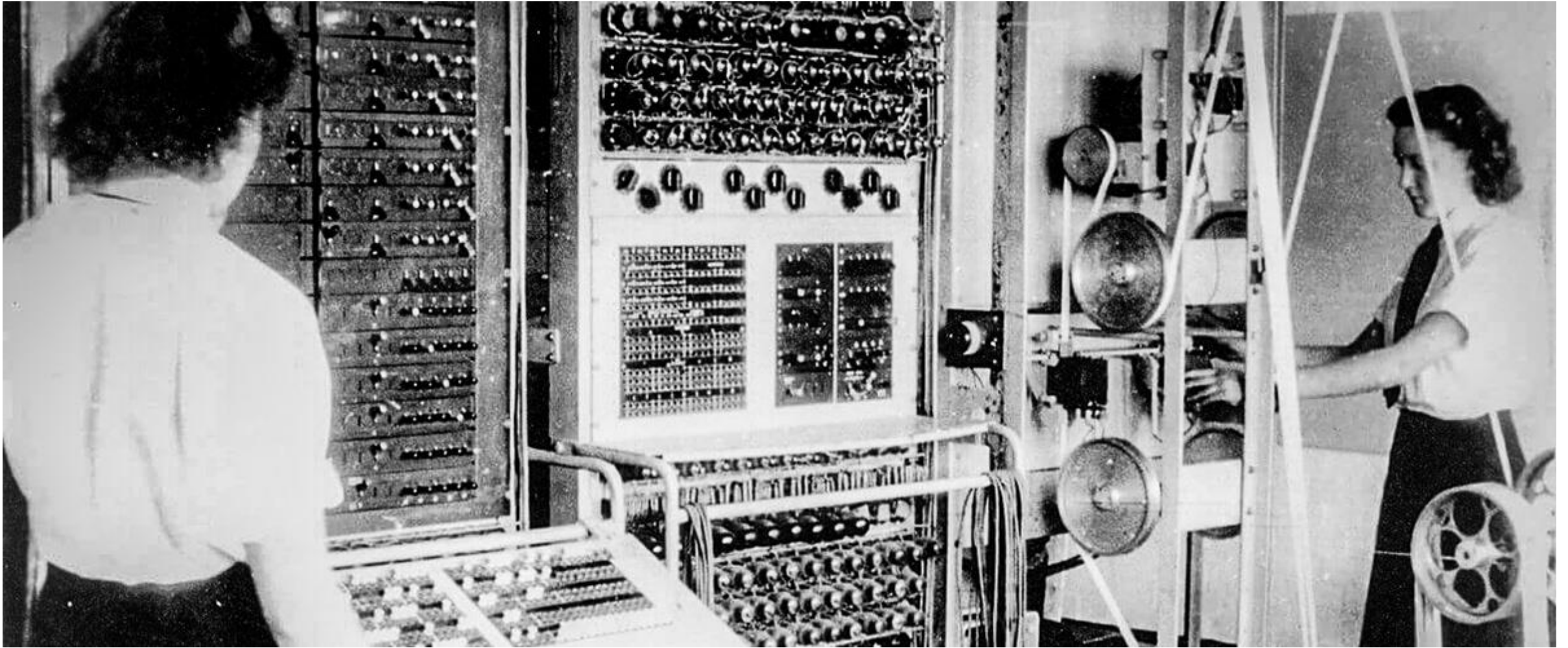
# Introduction to R programming

**NEUROSYSM930**

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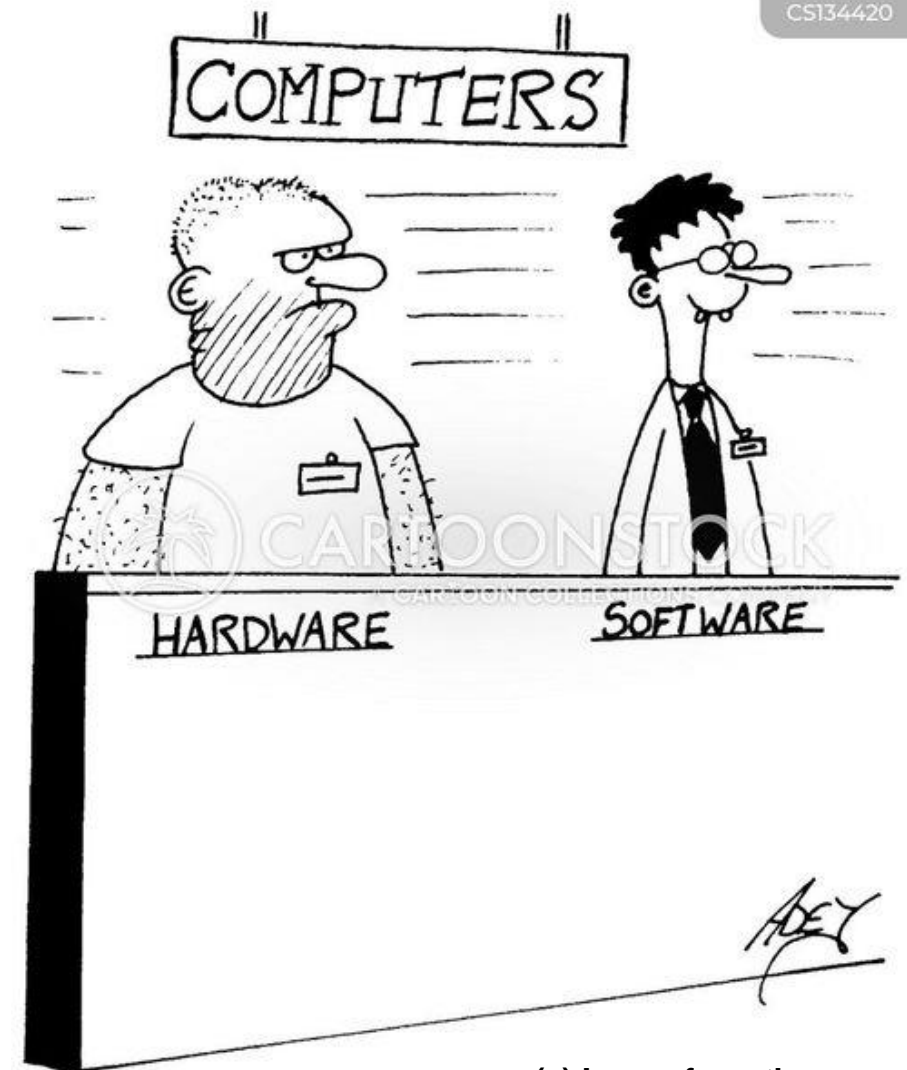
# Computers and Programming



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# Computers and Programming

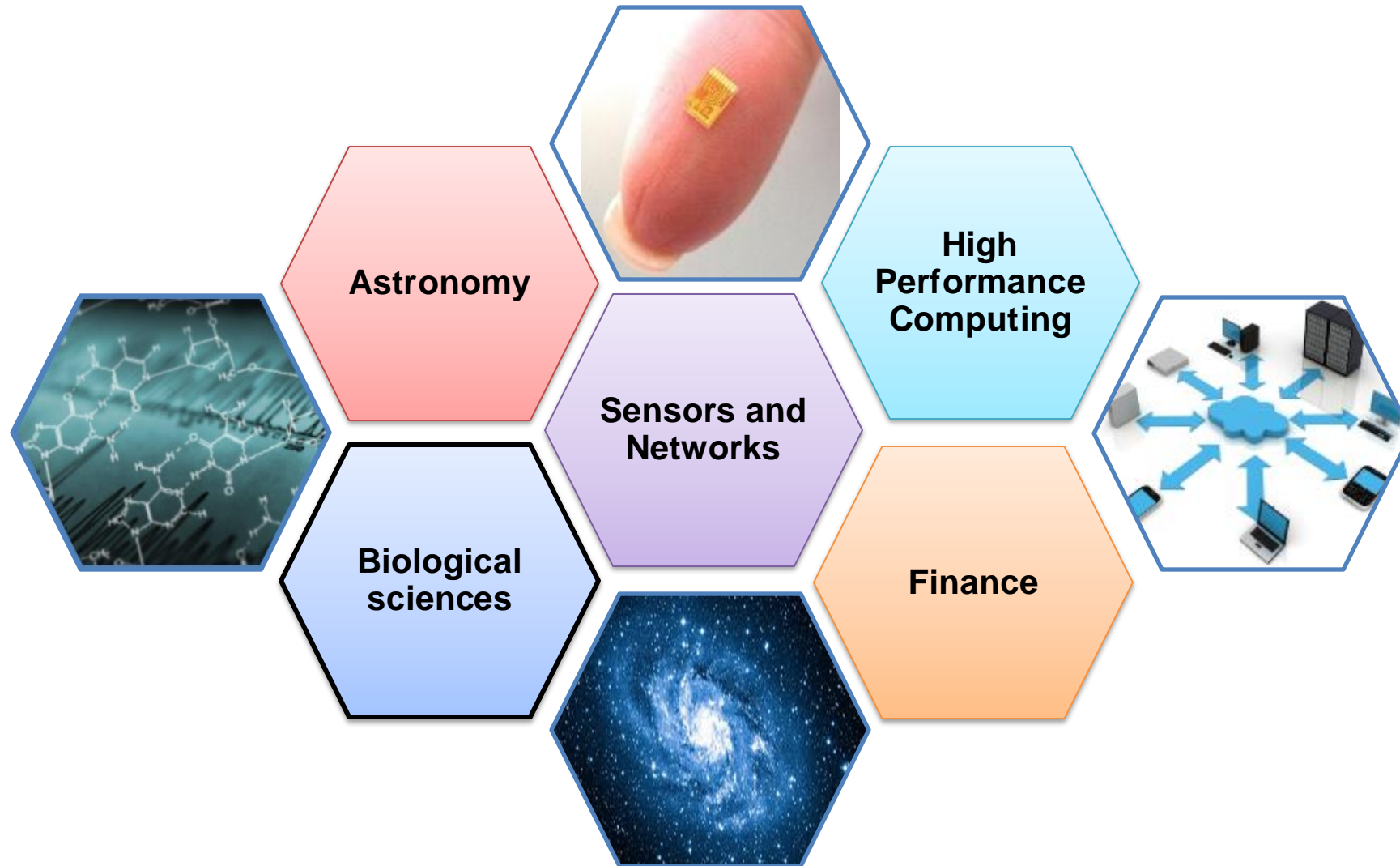
- ❖ A computer is just machine (**hardware**) for executing programs (**software**)
- ❖ The process of creating a program is called **programming**, and it is the focus of this course
- ❖ **Anyone can learn programming !**



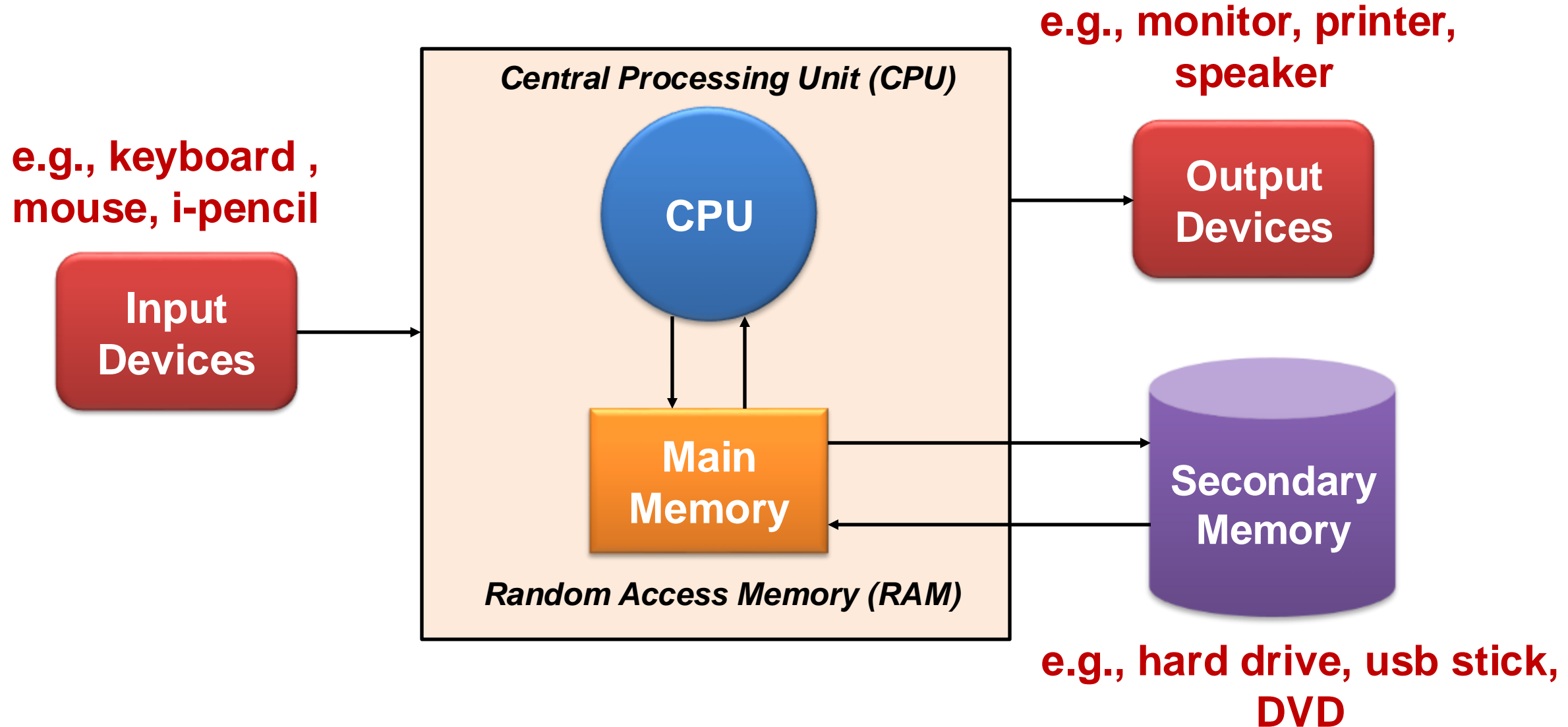
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# Computers and programming are part of our life



# But first we need to understand a computer

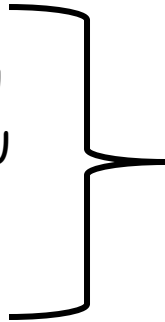


# Programming languages

- The hardware understands a **very low-level language** known as machine language which is translated to binary....

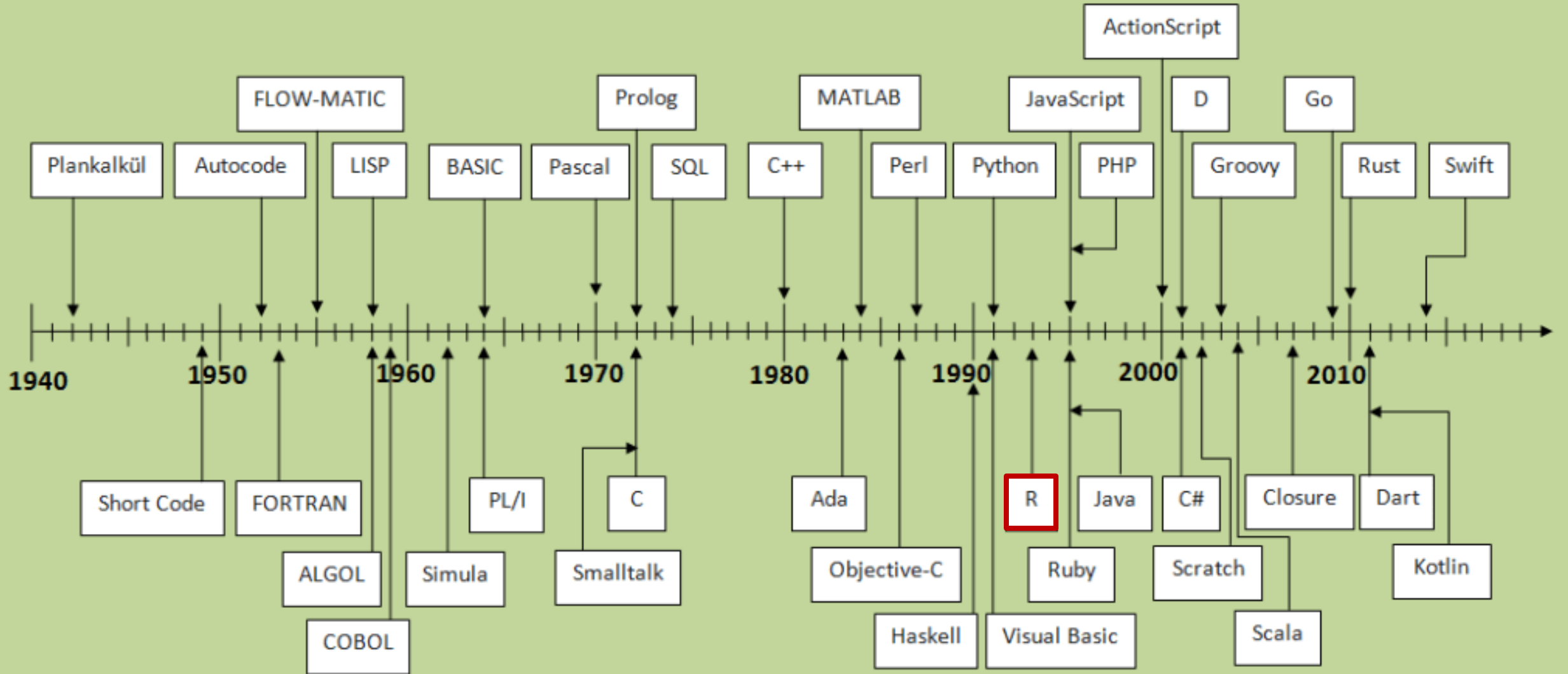
For example:

1. Load the number from memory location 2001 into the CPU
2. Load the number from memory location 2002 into the CPU
3. Add the two numbers in the CPU
4. Store the result into location 2003


$$A + B = C$$

- We need to use a **higher-level language and convert it** into a machine language that the hardware can execute...
  - **Compiled languages** → use a complex program called compiler
  - **Interpreted languages** → executes the code on the fly

# Timeline of Programming Languages



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# The brief history of R



***“R is a programming language for statistical computing and data visualization. It has been adopted in the fields of data mining, bioinformatics, and data analysis” (wiki)***

- R was developed in 1993 by **R**oss Ihaka and **R**obert Gentleman at the University of Auckland
- It was based on the S statistical programming language
- R software is **open-source and free** software licensed by the GNU Project
- In 1998, the Comprehensive R Archive Network **CRAN** was established
  - ✓ Now CRAN has more than 21,500 packages



# Explore the R architecture

**R is the engine**



**RStudio is the dashboard**



# Use R from the terminal

```
(base) c02z46udlvdc:NEUROSYSMED930_R_workshop kleftogi$ R

R version 4.4.1 (2024-06-14) -- "Race for Your Life"
Copyright (C) 2024 The R Foundation for Statistical Computing
Platform: x86_64-apple-darwin20

R is free software and comes with ABSOLUTELY NO WARRANTY.
You are welcome to redistribute it under certain conditions.
Type 'license()' or 'licence()' for distribution details.

R is a collaborative project with many contributors.
Type 'contributors()' for more information and
'citation()' on how to cite R or R packages in publications.

Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

> |
```

```
R Console

R version 4.4.1 (2024-06-14) -- "Race for Your Life"
Copyright (C) 2024 The R Foundation for Statistical Computing
Platform: x86_64-apple-darwin20

R is free software and comes with ABSOLUTELY NO WARRANTY.
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Natural language support but running in an English locale

R is a collaborative project with many contributors.
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Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

During startup - Warning messages:
1: Setting LC_CTYPE failed, using "C"
2: Setting LC_COLLATE failed, using "C"
3: Setting LC_TIME failed, using "C"
4: Setting LC_MESSAGES failed, using "C"
5: Setting LC_MONETARY failed, using "C"
[R.app GUI 1.80 (8416) x86_64-apple-darwin20]

WARNING: You're using a non-UTF8 locale, therefore only ASCII characters will work.
Please read R for Mac OS X FAQ (see Help) section 9 and adjust your system preferences accordingly.
[Workspace restored from /Users/kleftogi/.RData]
[History restored from /Users/kleftogi/.Rapp.history]

2024-11-18 10:20:30.006 R[48455:10555424] +[IMKClient subclass]: chose IMKClient_Modern
2024-11-18 10:20:30.823 R[48455:10555424] +[IMKInputSession subclass]: chose IMKInputSession_Modern
> |
```

## Use the R console

# The R studio environment

The screenshot shows the R Studio interface with several key components highlighted by red boxes and arrows:

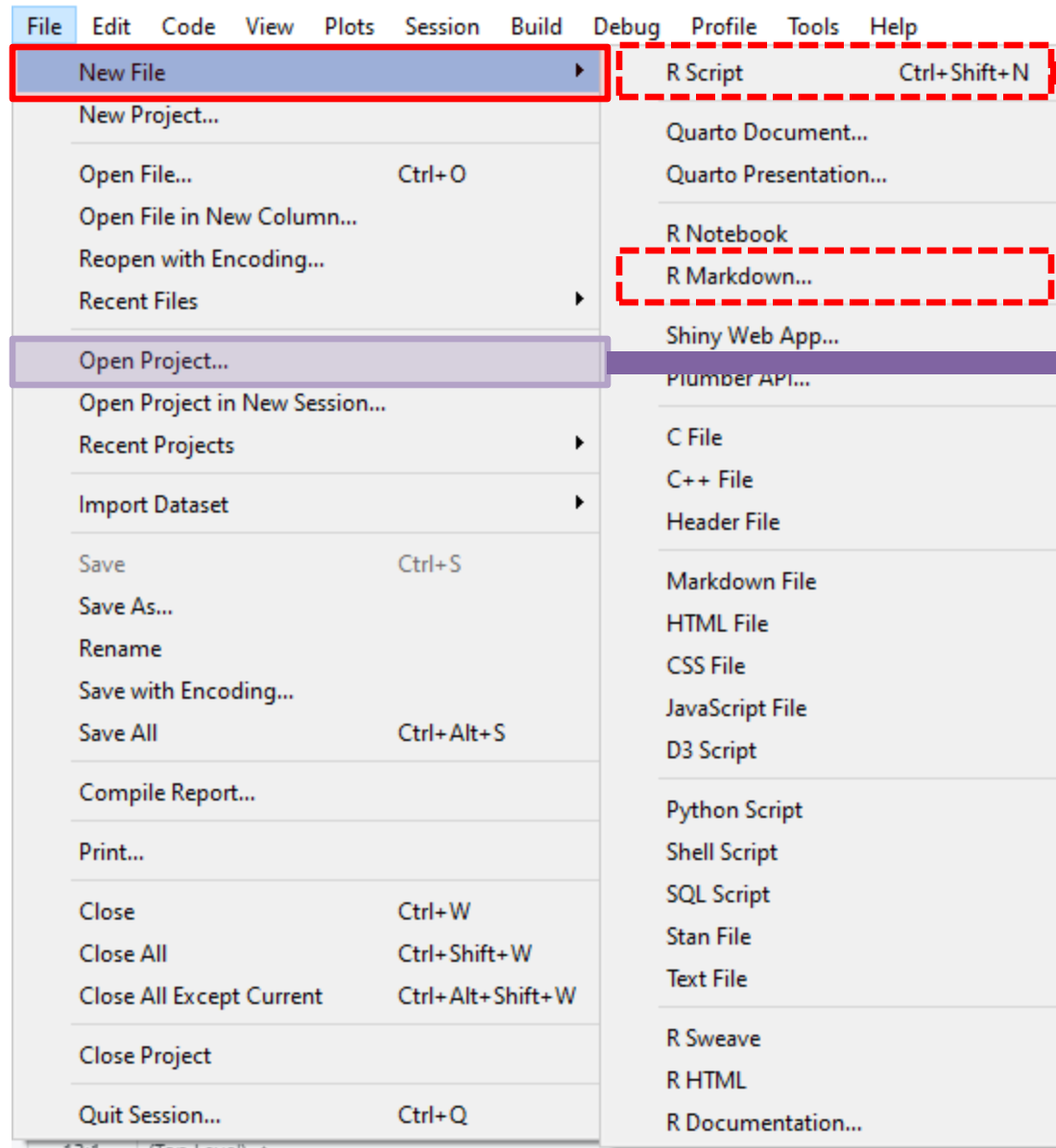
- Run Button:** A red box labeled "To run code, click on 'Run'" points to the "Run" button (a green arrow) in the top toolbar.
- Code Editor:** A red box labeled "It is just a text editor. Write code here." points to the main editor area where code is written.
- Environment Panel:** A red box labeled "Object will occur here:" with a bulleted list points to the Environment panel on the right. The list includes:
  - Data
  - Functions, etc.
- Console:** A red box labeled "R console Code will be executed here." points to the Console panel at the bottom left, which shows the R prompt and output.
- Bottom Panel:** A red box labeled with a bulleted list points to the bottom panel, which contains:
  - File explorer (working directory)
  - Plots
  - Packages to be downloaded
  - Help

The R Studio interface includes a menu bar (File, Edit, Code, View, Plots, Session, Build, Debug, Profile, Tools, Help), a toolbar with icons for file operations and running code, and a status bar at the bottom showing the current file and session information.

# What shall I do now ?

- As any other programming language R has its own syntax and semantics
- No buttons and clicking to perform computations...
- **But you need to learn how to program and you must invest time on that !**
  - Because...**
    - ✓ You can do basically everything you need in your daily data wrangling life with only one software solution...
    - ✓ Statistics and machine learning
    - ✓ Small and big data exploration
    - ✓ Great visualization
    - ✓ Develop web-apps even databases
- R packages are essentially free R programs that provide extra functionalities
  - ✓ Need to be downloaded and installed

# Different ways to create code



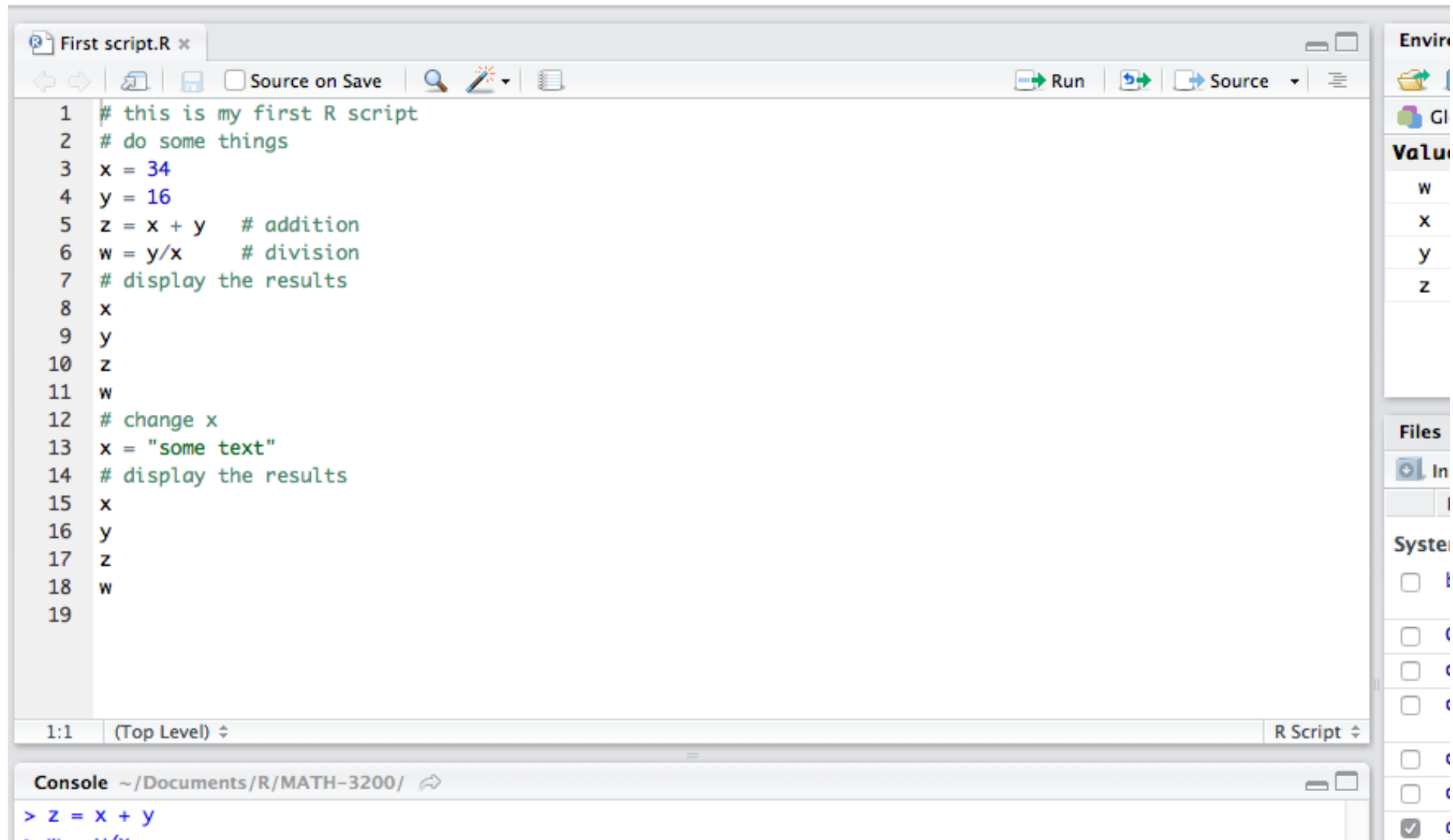
**New text editor (to write code)**

**Another type of text editor great for data presentation**

**Includes several files and generates a nice working space that:**

- ✓ You can copy the folder anywhere
- ✓ You can share the whole folder with colleagues

# R Scripts vs. R Markdowns



The screenshot displays the RStudio environment. The main editor window shows a script file named "First script.R" with the following R code:

```
1 # this is my first R script
2 # do some things
3 x = 34
4 y = 16
5 z = x + y # addition
6 w = y/x   # division
7 # display the results
8 x
9 y
10 z
11 w
12 # change x
13 x = "some text"
14 # display the results
15 x
16 y
17 z
18 w
19
```

The console window at the bottom shows the execution of the first two lines of the script:

```
> z = x + y
~ w = y/x
```

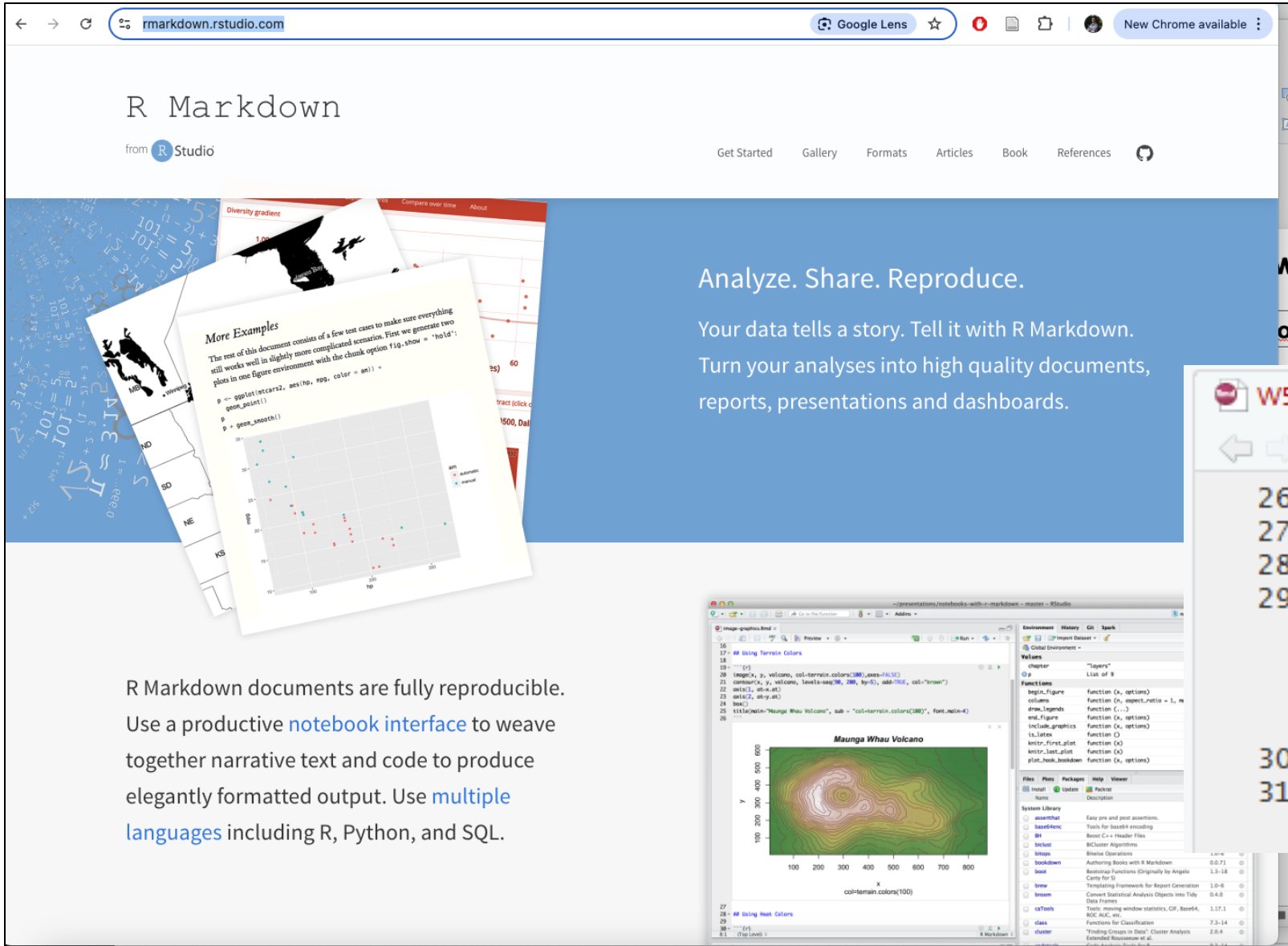
The right sidebar shows the "Environment" pane with a table of variables:

Value
w
x
y
z

The "Files" pane below it shows a folder icon and the text "In". The "System" pane at the bottom right shows a list of system variables with checkboxes.



# R Scripts vs. R Markdowns



R Markdown

from R Studio

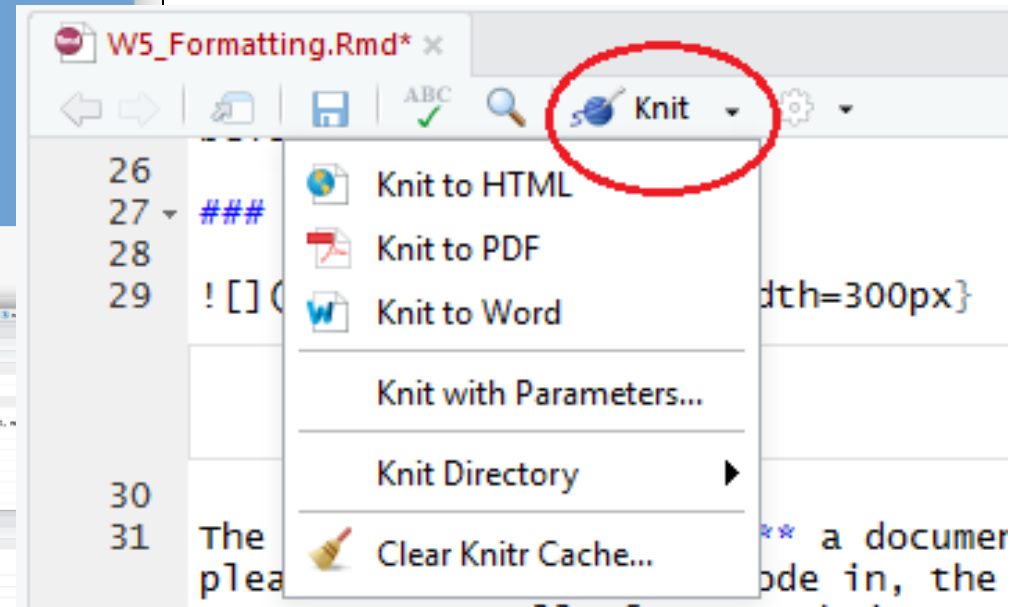
Get Started Gallery Formats Articles Book References

Analyze. Share. Reproduce.

Your data tells a story. Tell it with R Markdown.

Turn your analyses into high quality documents, reports, presentations and dashboards.

R Markdown documents are fully reproducible. Use a productive [notebook interface](#) to weave together narrative text and code to produce elegantly formatted output. Use [multiple languages](#) including R, Python, and SQL.



# Basic R concepts and terminology

- **Console** → where you enter in commands
- **Run** → the act of telling R to perform a computation by giving it commands in the console
- **Objects** → where values are saved in R, check the **Environment panel**
- **Data types** → integers, doubles/numerics, logicals, and characters.



# Errors, warnings and messages

```
> dat <- read.csv("data.csv")
```

```
Error in file
```

```
In addition
```

```
In file(fil
```

```
cannot op
```

```
> |
```

1:1 (Top Level) R Script

Console Terminal x Jobs x

~/

```
> library("dplyr")
```

Attaching p

The followi

between

The followi

filter,

The followi

interse

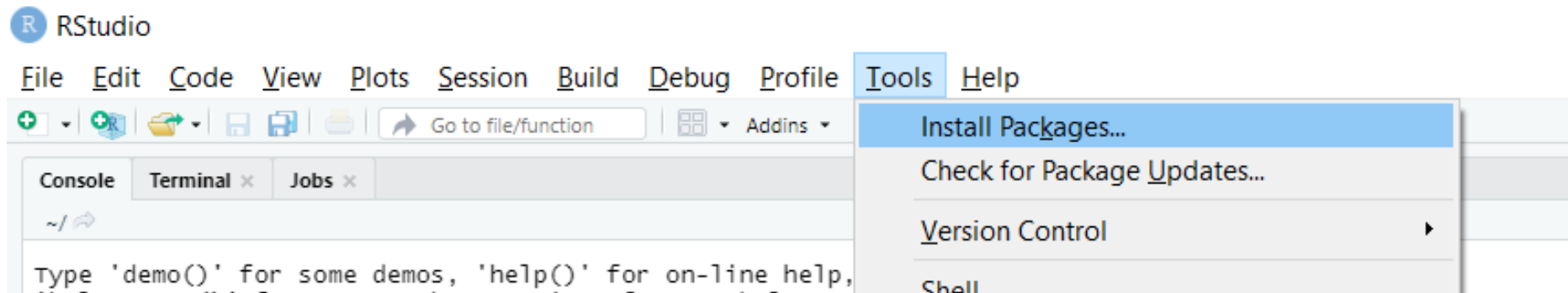
Warning message:  
package 'dplyr' was built under R version 3.6.2



oup"

# Packages

- Different ways to install packages exist



## Install Bioconductor Packages

To install core packages, type the following in an R command window:

```
if (!require("BiocManager", quietly = TRUE))
  install.packages("BiocManager")
BiocManager::install(version = "3.20")
```

To install core packages, type the following in an R command window:

```
BiocManager::install(c("GenomicFeatures", "AnnotationDbi"))
```

The `install.packages()` function (in the `BiocManager` package) has arguments that change its default behavior type `BiocManager::install` for further help.

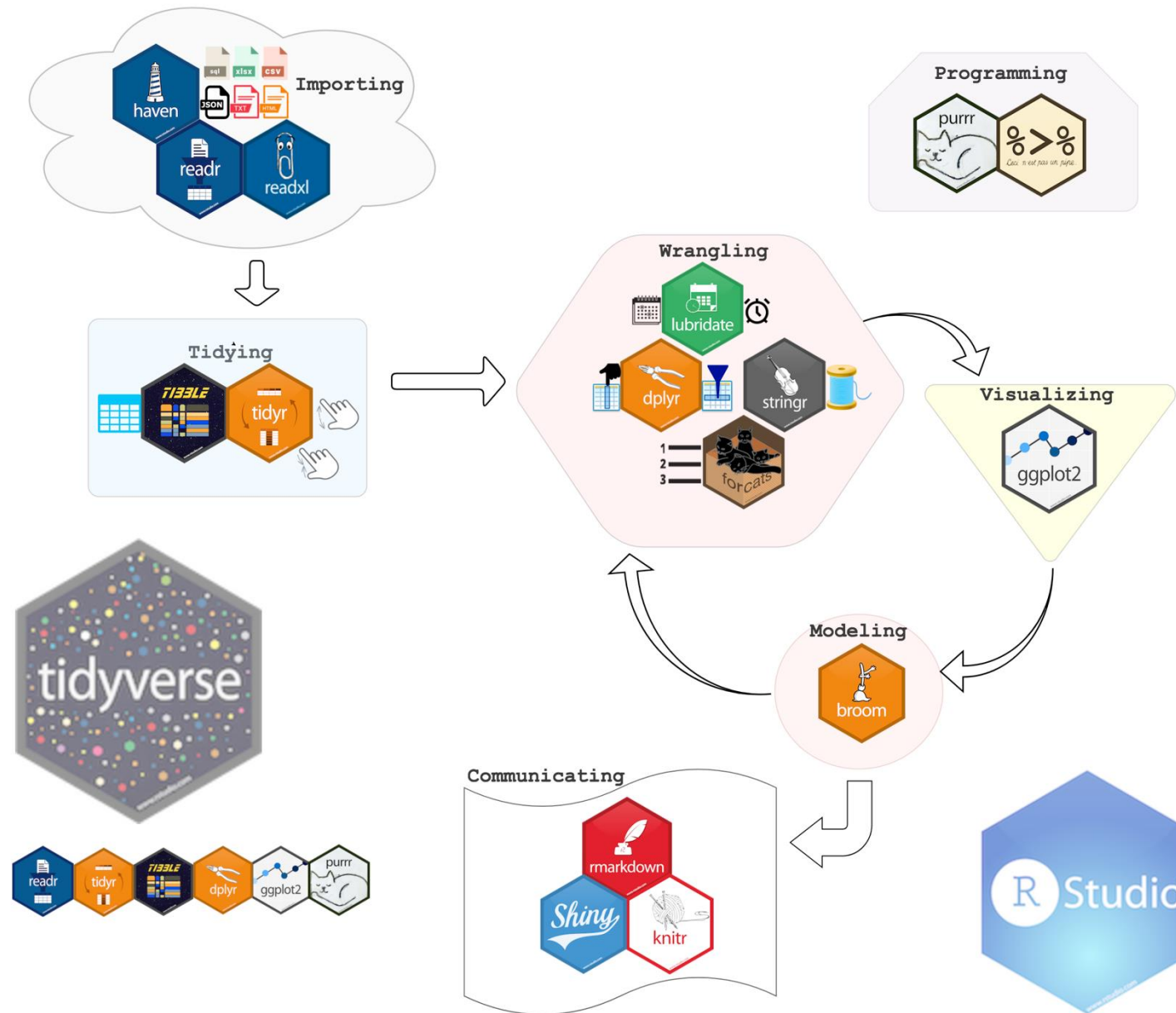
# Packages

- Never forget to load the installed packages....

A screenshot of an R console window. The window has a title bar with a green 'R' icon. Below the title bar, there are three circular icons: a copy icon, a paste icon, and a moon icon. The main area of the console is a dark gray bar containing the text `library("ggplot2")` in a light blue font.

**And never forget that R is distributed with fifteen "base packages":** *base, compiler, datasets, grDevices, graphics, grid, methods, parallel, splines, stats, stats4, tcltk, tools, translations, and utils*

# By combining R packages, we create workflows





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# Thanks for your attention !

## Q&A

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