

# **Espectroscopia em herbário com MicroNIR: Teoria e prática**

## **Aula 2 (tarde)**

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Manaus, 03 de abril de 2025

# Conteúdo

- Reunindo as leituras obtidas por cada grupo
- Aula expositiva sobre R:
  - Boas práticas no R
  - Executando um script R:
    - Instalação das bibliotecas necessárias
    - Importação e preparação das leituras espectrais
    - Inspeção visual dos espectros
    - Preparação dos conjuntos de dados para análise
- Mãos à obra!

# Reunião das leituras obtidas por cada grupo

- Os grupos devem repassar as leituras feitas
- Lembrando que as leituras estão dentro da pasta de trabalho criada por vocês em:
  - Documentos/InnoSpectra/Scan Results/

# Quais arquivos vamos precisar?

- Leituras espectrais (vários arquivos **.csv**)
- Script (**01\_Importando e preparando dados espectrais.R**)
- Banco de dados de *Handroanthus* do herbário INPA (**Handroanthus.xlsx**)

# Boas práticas no R

- Considere instalar numa pasta segura (por exemplo, C:\R):
  - R tools - <https://cran.r-project.org/bin/windows/Rtools/>

## RTools: Toolchains for building R and R packages from source on Windows

Choose your version of Rtools:

<a href="#">RTools 4.5</a>	for R versions from 4.5.0 (R-prerelease and R-devel)
<a href="#">RTools 4.4</a>	for R versions 4.4.x (R-release)
<a href="#">RTools 4.3</a>	for R versions 4.3.x (R-oldrelease)
<a href="#">RTools 4.2</a>	for R versions 4.2.x
<a href="#">RTools 4.0</a>	for R from version 4.0.0 to 4.1.3
<a href="#">old versions of RTools</a>	for R versions prior to 4.0.0



Rtools45 consists of Msys2 build tools, GCC 14/MinGW-w64 compiler toolchain, libraries built using the toolchain, and OPDF. Rtools45 supports 64-bit Windows and UCRT as the C runtime.

Compared to Rtools44, Rtools45 for 64-bit Intel machines has newer versions of two core components: GCC and binutils. It is recommended to re-compile all code with the new toolchain to avoid problems.

Rtools45 is also available for 64-bit ARM machines (aarch64): it includes Msys2 build tools (64-bit Intel builds running via emulation) and aarch64 builds of LLVM 19/MinGW-w64 compiler toolchain, libraries built using the toolchain, and again QPDF. The 64-bit ARM version of Rtools45 is experimental: a number of CRAN packages don't work with it and the Fortran compiler (flang-new) is not yet able to compile Fortran code of all CRAN packages. A number of CRAN packages don't work because they require not-yet-available 64-bit ARM versions of external software.

## Installing Rtools45

Rtools is only needed for installation of R packages from source (those that need compilation of C/C++ or Fortran code) or building R from source. R can be installed from the R binary installer and by default will install binary versions of CRAN packages, which does not require Rtools45.

Moreover, online build services are available to check and build R packages for Windows, for which again one does not need to install Rtools<sup>45</sup> locally. The [Winbuilder](#) check service uses identical setup as the CRAN incoming packages checks and has already all CRAN and Bioconductor packages pre-installed.

Rtools45 may be installed from the [Rtools45 installer](#) or [64-bit ARM Rtools45 installer](#). It is recommended to use the defaults, including the default installation location of C:\rtools45.

When using R installed by the installer, no further setup is needed after installing Rtools45 to build R packages from source. When using the default installation location, R and Rtools45 may be installed in any order and Rtools45 may be installed when R is already running.

On ARM, binary versions of packages are currently not available from CRAN, so Rtools<sup>45</sup> is required to install any package that needs compilation.

### Additional information

A detailed tutorial on how to build R and packages using Rtools<sup>45</sup> for R package authors and R developers is available for [R 4.5.x](#) and [R-devel](#).

From the user perspective, Rtools45 is the same as Rtools42-44. However, it uses newer versions of the compiler toolchain and libraries and some libraries have been added, hence some package authors will have to adapt their make files. Maintainers of CRAN and Bioconductor packages may use [these patches](#) for reference or re-use them in their code.

A change log for Rtools45 vs Rtools44 and of individual revisions of Rtools45 is available [here](#)

Rtools45 is also available in base and full toolchain tarballs suitable for users who have their own installation of Msys2. The base toolchain tarball is smaller and includes only what is needed to build R and the recommended packages. All Rtools files are available [here](#)

# Boas práticas no R

- Considere instalar numa pasta segura (por exemplo, C:\R):
  - R tools - <https://cran.r-project.org/bin/windows/Rtools/>
  - R base - <https://cran.r-project.org/bin/windows/base/>



[Download R-4.4.3 for Windows](#) (85 megabytes, 64 bit)

[README on the Windows binary distribution](#)

[New features in this version](#)

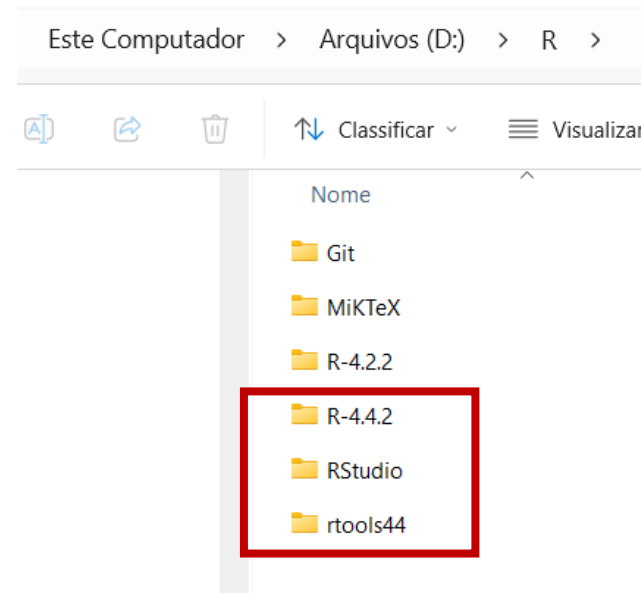
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## Frequently asked questions

- Other builds

# Boas práticas no R

- Considere instalar numa pasta segura (por exemplo, C:\R):
  - R tools - <https://cran.r-project.org/bin/windows/Rtools/>
  - R base - <https://cran.r-project.org/bin/windows/base/>
  - R Studio (IDE) - <https://posit.co/download/rstudio-desktop/>





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## DOWNLOAD

# RStudio Desktop

Used by millions of people weekly, the RStudio integrated development environment (IDE) is a set of tools built to help you be more productive with R and Python.

Don't want to download or install anything? Get started with RStudio on [Posit Cloud for free](#). If you're a professional data scientist looking to download RStudio and also need common enterprise features, don't hesitate to [book a call with us](#).

Want to learn about core or advanced workflows in RStudio? Explore the [RStudio User Guide](#) or the [Getting Started](#) section.

## 1: Install R

RStudio requires R 3.6.0+. Choose a version of R that matches your computer's operating system.

*R is not a Posit product. By clicking on the link below to download and install R, you are leaving the Posit website. Posit disclaims any obligations and all liability with respect*

## 2: Install RStudio

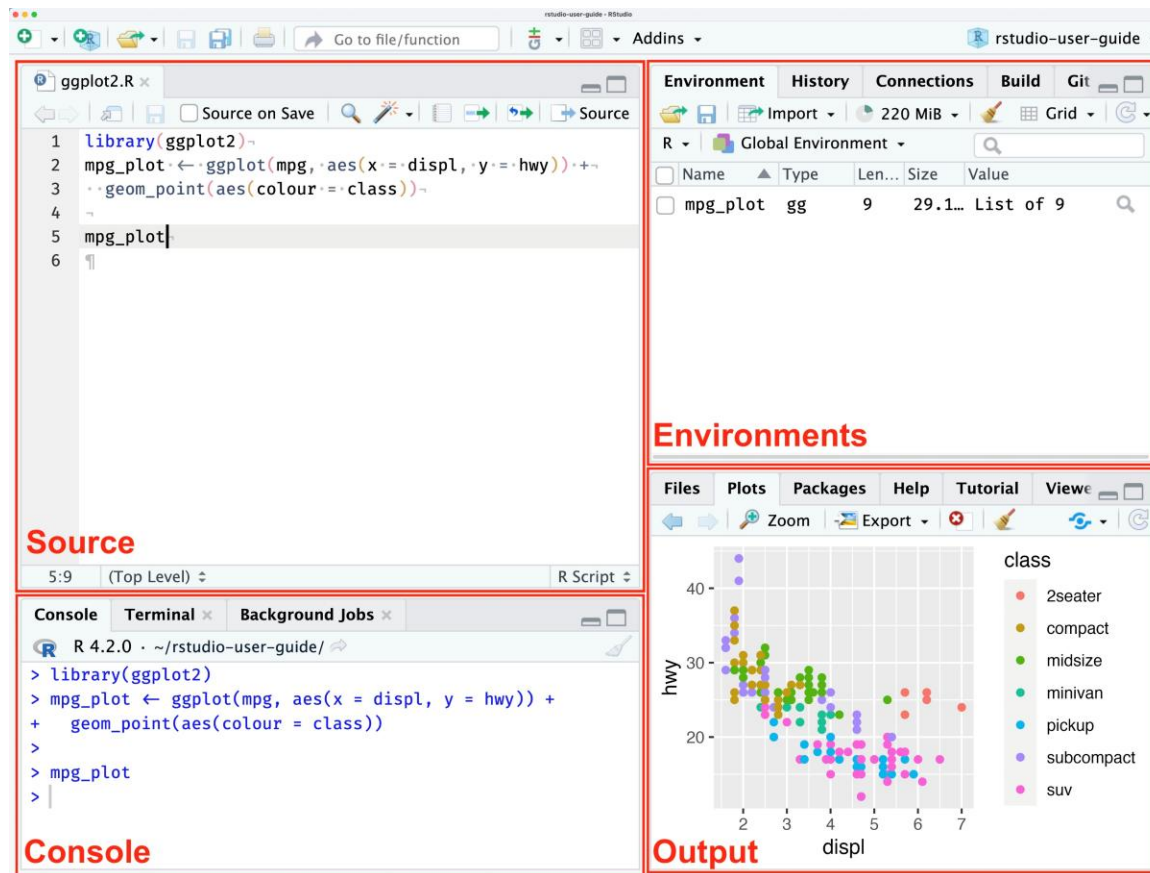
DOWNLOAD RSTUDIO DESKTOP FOR WINDOWS

Size: 265.28 MB | [SHA-256: BB369743](#) | Version: 2024.12.1+563  
Released: 2025-02-13



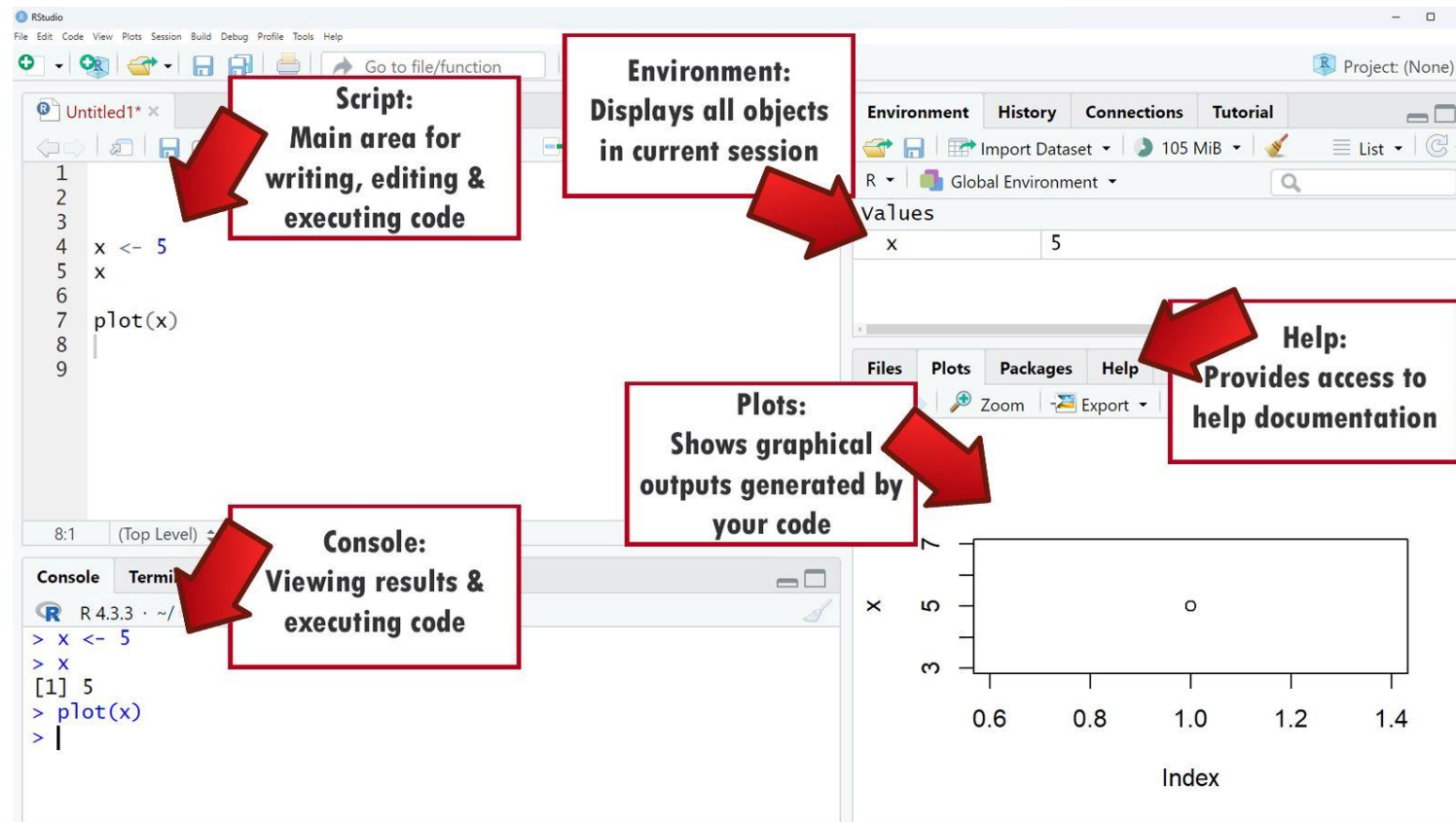
# Boas práticas no R

- Área de trabalho do R Studio (painéis)



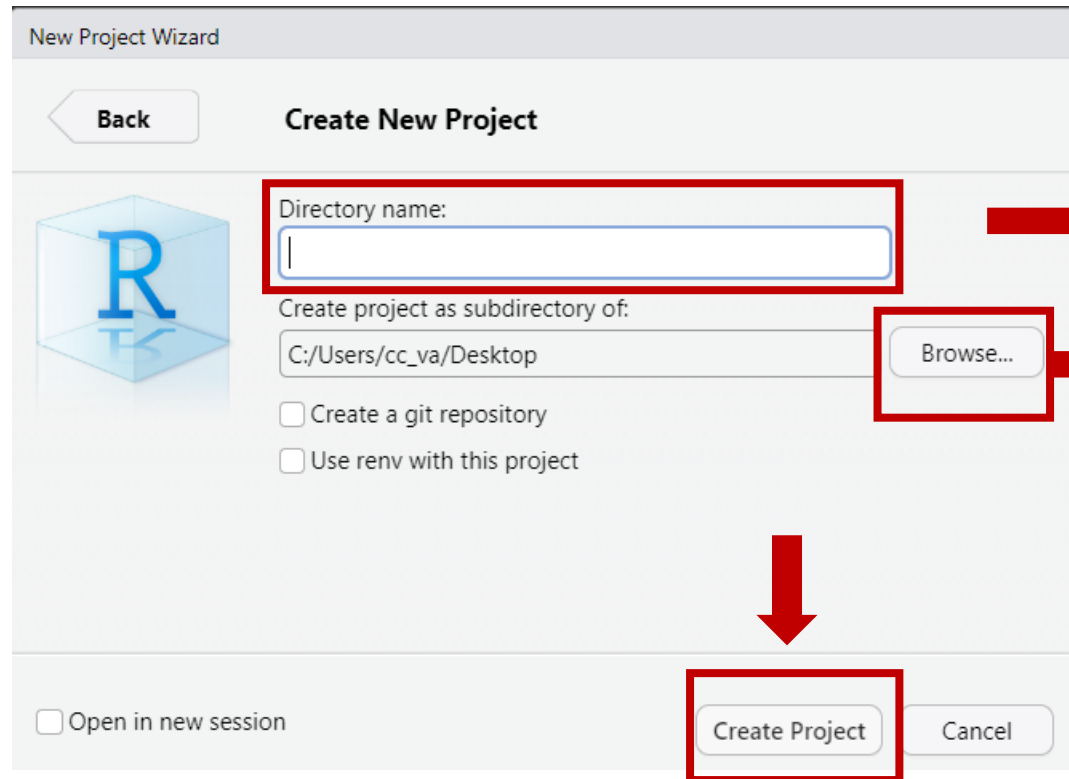
# Boas práticas no R

- Área de trabalho do R Studio (o que você precisa saber!)



# Boas práticas no R

- Criando um projeto no R Studio (.Rproj)
- File > New Project > New Directory > New Project



**Vamos usar "NIR\_spp\_CITES"**

**Define o local onde a pasta deve ficar. No meu caso eu deixei no Desktop**

R NIR\_spp\_CITES - RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

+ - + - Go to file/function Addins

Console Terminal Background Jobs

R 4.4.2 · C:/Users/cc\_va/Desktop/NIR\_spp\_CITES/

R version 4.4.2 (2024-10-15) "The R Project for Statistical Computing"  
Copyright (C) 2024 The R Foundation for Statistical Computing  
Platform: x86\_64-w64-mingw32 x86\_64-pc-windows-gnu

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'citation()' para saber como citar o R ou pacotes do R em publicações.

Digite 'demo()' para demonstrações, 'help()' para o sistema on-line de ajuda,  
ou 'help.start()' para abrir o sistema de ajuda em HTML no seu navegador.  
Digite 'q()' para sair do R.

> |

Environment History Connections Tutorial

Import Dataset 92 MiB

R Global Environment

Environment is empty

Files Plots Packages Help Viewer Presentation

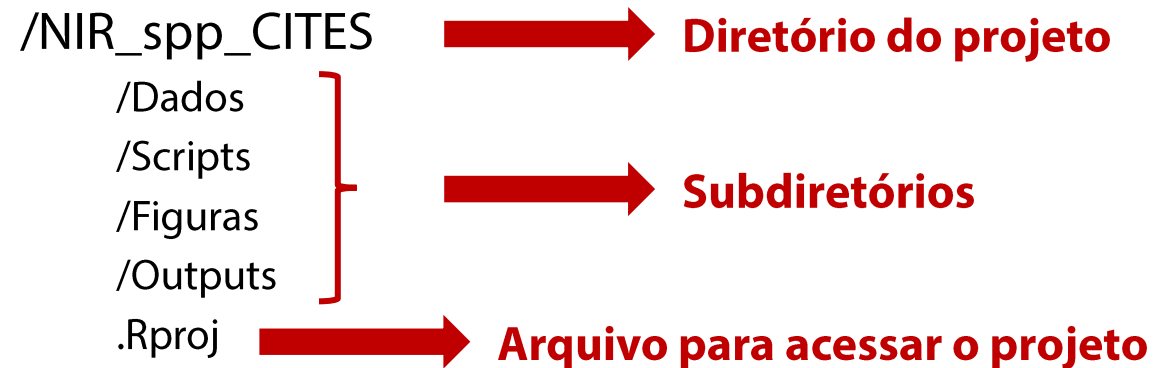
C: > Users > cc\_va > Desktop > NIR\_spp\_CITES

	Name	Size	Modified
↑	..		
📁	NIR_spp_CITES.Rproj	267 B	Apr 3, 2025, 2:34

**Projeto criado!**

# Boas práticas no R

- Estruturas de pastas



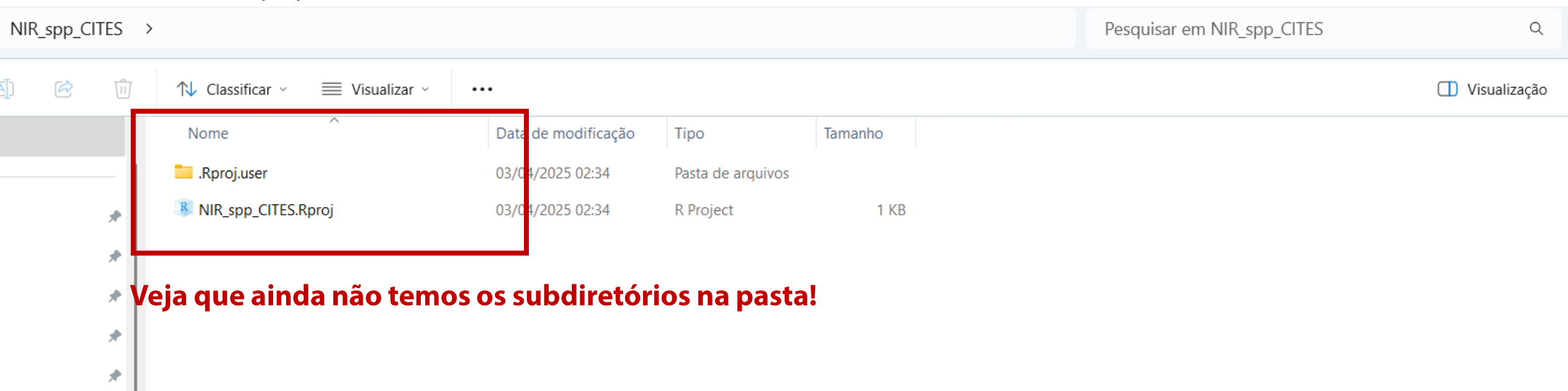
- Agora vamos, criar esses subdiretórios!



# Boas práticas no R

- Estruturas de pastas

/NIR\_spp\_CITES  
.Rproj

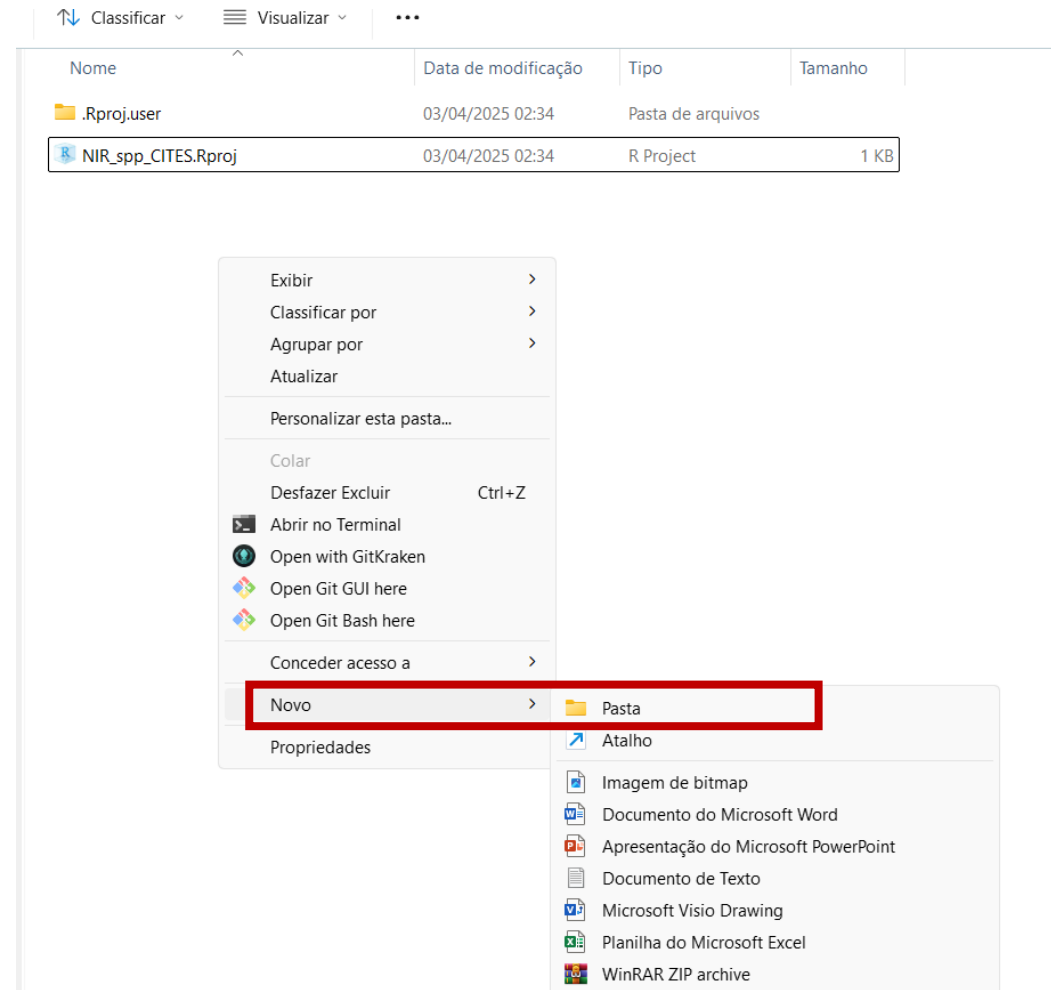


# Boas práticas no R

**Você pode fazer  
de forma manual**

**Ou**

**Através do R  
Studio (próximos  
slides)**



Faça isso para cada:

**/Dados  
/Scripts  
/Figuras  
/Outputs**

# Boas práticas no R

- Código para copiar e colar no **console** do R Studio:

```
subdirs <- c("Scripts", "Dados", "Figuras", "Outputs")

for (d in subdirs) {
  if (!dir.exists(d)) {
    dir.create(d)
  }
}
```

NIR\_spp\_CITES - RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

+ - + -

Go to file/function

Addins

Console Terminal Background Jobs

R 4.4.2 · C:/Users/cc\_va/Desktop/NIR\_spp\_CITES/

R version 4.4.2 (2024-10-31 ucrt) -- "Pile of Leaves"  
Copyright (C) 2024 The R Foundation for Statistical Computing  
Platform: x86\_64-w64-mingw32/x64

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```
> subdirs <- c("Scripts", "Dados", "Figuras", "Outputs")

for (d in subdirs) {
  if (!dir.exists(d)) {
    dir.create(d)
  }
}
```

Environment History Connections Tutorial

Import Dataset 91 MiB

List

R Global Environment

Environment is empty

Files Plots Packages Help Viewer Presentation

C: > Users > cc\_va > Desktop > NIR\_spp\_CITES

	Name	Size	Modified
↑	..		
📄	.Rhistory	255 B	Apr 3, 2025, 3:04
📁	NIR_spp_CITES.Rproj	267 B	Apr 3, 2025, 3:04

Digite ou copie/cole este  
código no console e execute  
(Ctrl + R)

# Executando um script R

- Copie os arquivos das leituras espectrais para o subdiretório **“Dados”**
- Copie também o arquivo **“Handroanthus.xlsx”** para o subdiretório **“Dados”**
- Agora, copie o arquivo **“01\_Importando e preparando dados espectrais.R”** para o subdiretório **“Scripts”** na pasta do projeto
- Com o projeto aberto, abra o script no R Studio: File > Open File > Selecione a pasta **Scripts** > Selecione o arquivo do script **“01\_Importando e preparando dados espectrais.R”**

# Acompanhe a execução do script na tela do R Studio



**Agora é com vocês:**

**Mãos à obra!**

# Lista de presença (Dia 2/tarde)

<https://forms.gle/WnXsJYpzR1aw5kxT8>

ou

