

# User Documentation

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## 1 Software and Hardware Requirements to Run the Project

To run this software, you must first have R installed along with any IDE that allows handling R files. In this case, RStudio has been used, but any other IDE is viable (Visual Studio Code, TinnR, Eclipse, Rbase, etc.). The project has been developed on Windows, so the installation instructions are specifically for this operating system.

Separately, we must also install the MongoDB driver, MongoDB Compass, so the system can create patient records locally. How to install all these programs is detailed in the next section.

## 2 Installation / Setup

### 2.1 Installing R

To install R on Windows, you can download it from the CRAN website [CRAN, 2024]. Once on the website, click on the "*Download*" section, as shown in Figure 1. This will download an executable (.exe) file that will be found in the Downloads folder.

When executing it, you must select the installation language 2. Once selected, the license agreements will appear 3, and you will be given the option to choose the installation directory 4; in this case, the default folder will be used. Then, you will select the components to install and specify if you want to use configuration options, as shown in Figures 5 and 6. Finally, before starting the installation, you will choose a folder to store the program's shortcuts 7 and define additional tasks to be performed during installation 8.

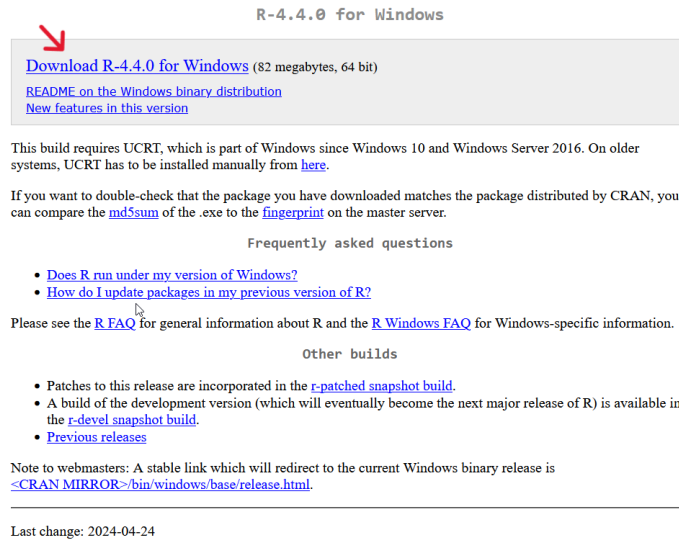


Figure 1: Download R

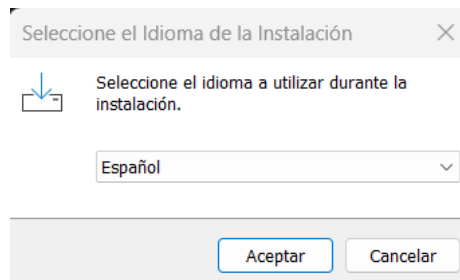


Figure 2: Language Selection

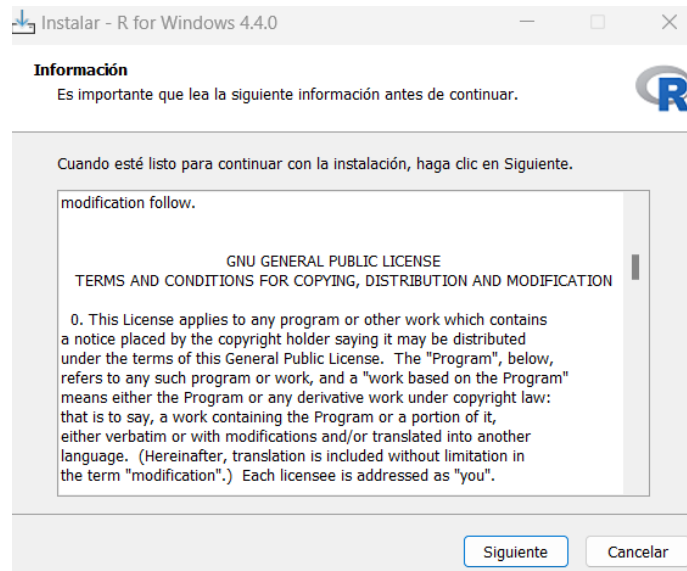


Figure 3: Licenses

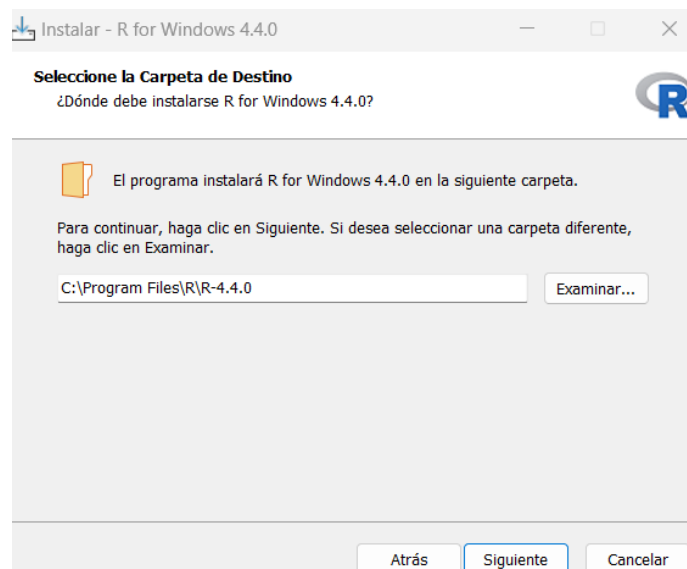


Figure 4: Destination Folder Selection

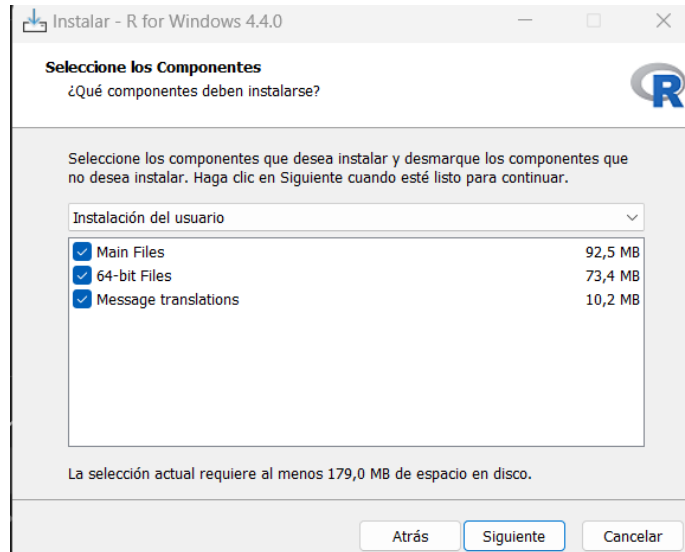


Figure 5: Component Selection

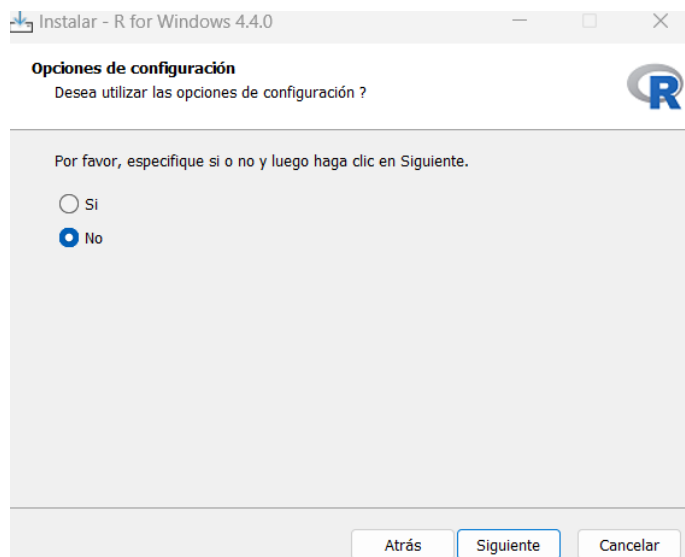


Figure 6: Configuration Options

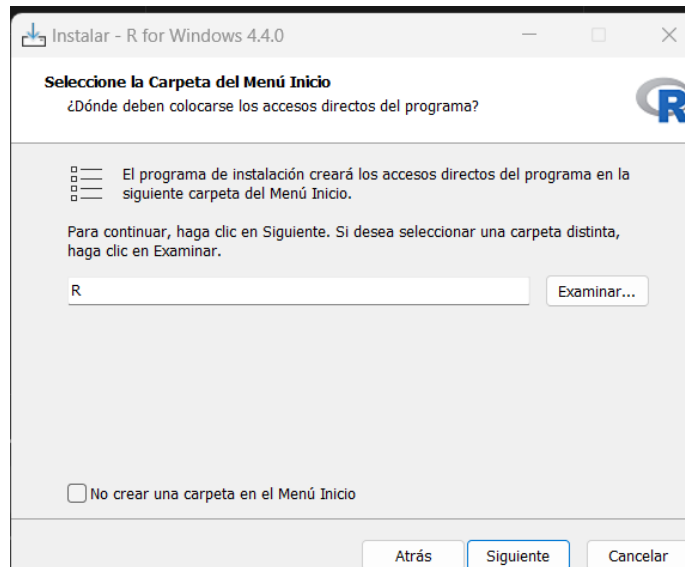


Figure 7: Shortcut Folder Selection

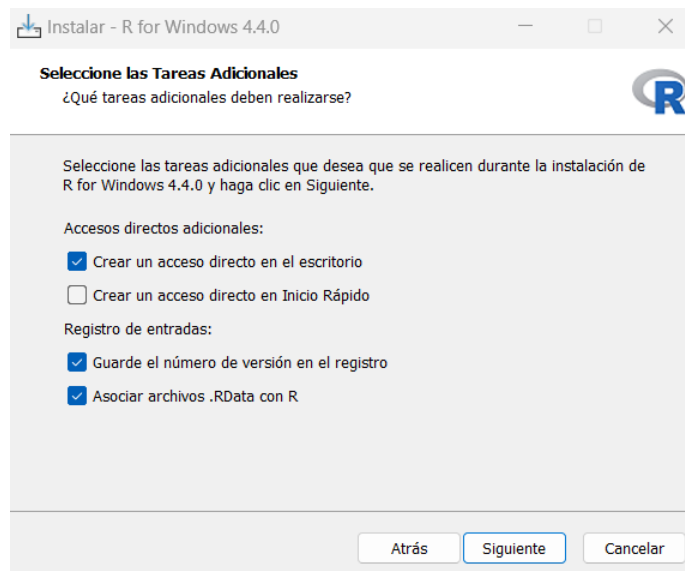


Figure 8: Additional Tasks

## 2.2 Installing RStudio

The installation of RStudio is similar to that of R. To download the installation file 9, go to the Posit website [Posit, 2024].

When executing the file, you will be asked to indicate the destination directory 4 and the folder for RStudio shortcuts 11. Once this information is completed, the installation will begin.

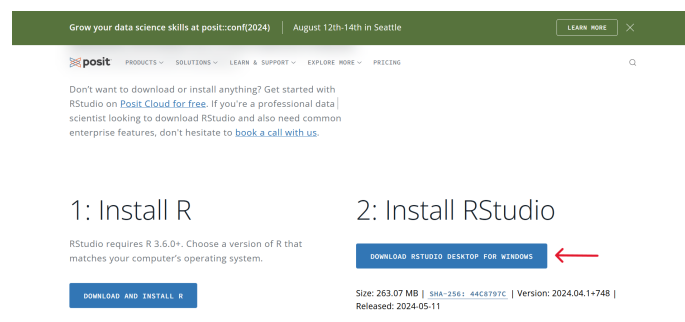


Figure 9: Download Installation File

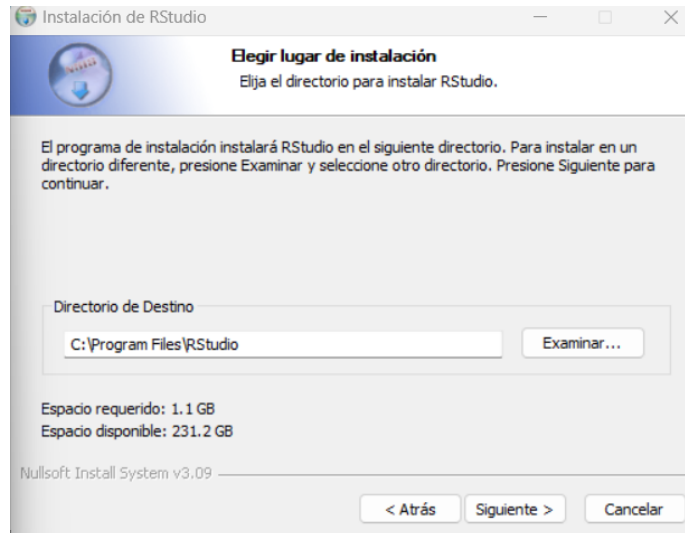


Figure 10: Destination Folder Selection

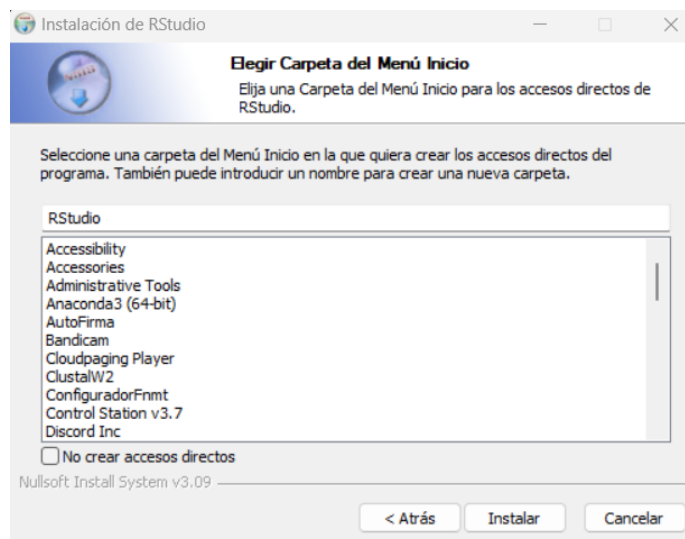


Figure 11: Shortcut Folder Selections

## 2.3 Installing MongoDB Compass

On the official MongoDB website [MongoDB, 2024], you can find the MongoDB Compass installer 12. In this case, no configuration is required; the application will install automatically upon executing the downloaded file.

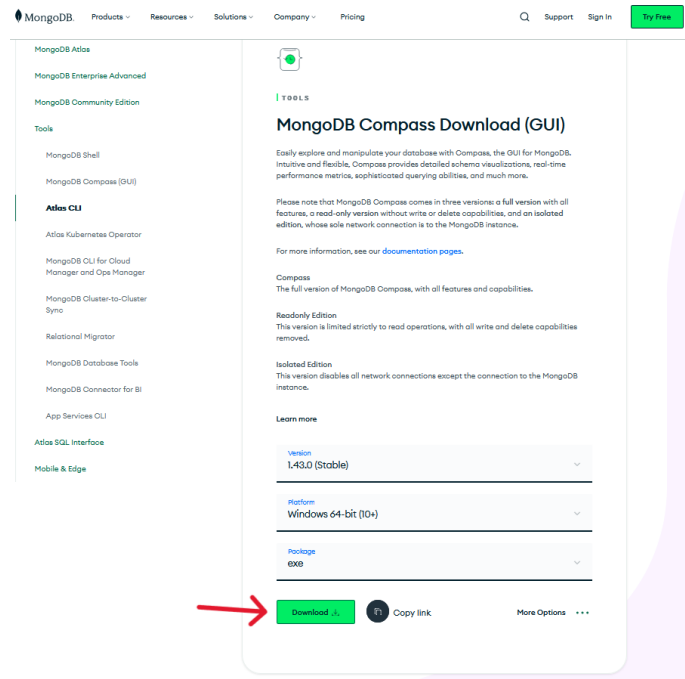


Figure 12: Download MongoDB Compass Installer

Finally, to launch the Shiny application, you only need to clone the GitHub repository, open the PDF-Scrapping.R file, and execute it.

The instructions to clone a repository [GitHub, 2024] are:

- Go to the GitHub repository page.
- Click the green button labeled "Code" 13.
- Select the cloning method (HTTPS, SSH, or GitHub CLI).
- Copy the displayed URL.
- Open Git Bash and navigate to the directory where you want to clone the repository.
- Type "git clone" followed by the copied URL.



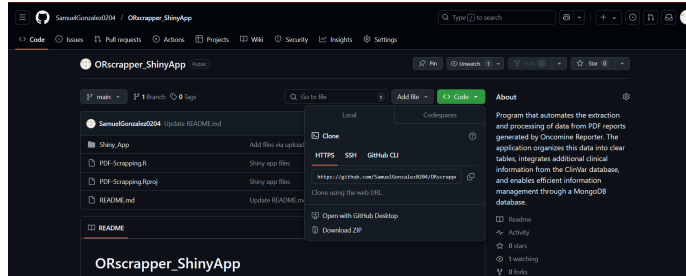


Figure 13: Repository Cloning

## 3 Manuals and/or Practical Demonstrations

Following the use case diagram in Annex F, demonstrations for each use will be shown.

### 3.1 Processing PDFs

Figure 14 shows how data has been extracted from three uploaded files.

Numero de estudio	Numero de muestra	NIC	Diagnostico	Fecha de informe	diagnostico	Sexo	Porcentaje tumoral	Calidad	Diagnostico	Numero
1	106.0	BU220001 A1	000	16 feb 2024	ADENOCARCINOMA GASTROINTESTINAL DE PULMON	MASC	50	ADENOCARCINOMA GASTROINTESTINAL DE PULMON	Centromex	10.1
2	140.0	BU220001 A1	100	27 mar 2024	ADENOCARCINOMA GASTROINTESTINAL DE PULMON	MASC	60	ADENOCARCINOMA GASTROINTESTINAL DE PULMON	Centromex	10.1
3	140.0	BU22012001 A2	00000	28 mar 2024	MELANOMA	FEM	75	ADENOCARCINOMA GASTROINTESTINAL DE PULMON	Centromex	10

Figure 14: Demonstration of PDF Processing

### 3.2 Querying Searches in Clinical Databases

The column Patogenicidad.Buscada in the table of Figure 15 corresponds to the searches for mutations in the ClinVar database.

Relaciones de mutaciones	Numero de mutaciones en el archivo PDF	Total del numero de mutaciones	Porcentaje de frecuencia en la poblacion	Frases de ID	Patogenicidad	Patogenicidad Buscada	Ensayos clínicos	SNP array	Frecuencia de mutaciones	SNP array
ALK/TP53	4.25	2	10.75.36.42			SNP array	0	0	0	0
BRCA1/BRCA2	10.12	2	22.91.31.30		Patogenicidad	Patogenicidad	10	1	7	1

Figure 15: Demonstration of querying clinical databases

### 3.3 Modifying Tables

Double-clicking on any cell in the tables will open an editor where we can modify the value of that cell.

Numero de chip	Numero de Hospital	NIC	Estado actual	Fecha de ingreso	Diagnostico	Sexo	Porcentaje tumoral	Celulas	Diagnostico	Nombre
1 108.0	BUZOS001A1	000	1	16-04-2024	ADENOCARCINOMA EPIDERMICO DE PULMON	MASC	55	ADENOCARCINOMA EPIDERMICO DE PULMON	Centinela pulmonar no metastatizante	15.1
2 148.0	BUZOS001A1 - externo	17	1	27-mar-2024	ADENOCARCINOMA DE PULMON	MASC	60	ADENOCARCINOMA DE PULMON	Centinela pulmonar no metastatizante	15.1
3 148.0	BUZOS001A2	00000	1	26-mar-2024	MELANOBALIV	FEM	75	ADENOCARCINOMA DE PULMON	Metastasis	15

Figure 16: Demonstration of table modification

### 3.3.1 Sorting Values

In this example 17, the column Porcentaje\_tumoral has been sorted in descending order.

Numero de chip	Numero de Hospital	NIC	Estado actual	Fecha de ingreso	Diagnostico	Sexo	Porcentaje tumoral	Celulas	Diagnostico	Nombre
3 148.0	BUZOS001A2	00000	1	26-mar-2024	MELANOBALIV	FEM	75	ADENOCARCINOMA DE PULMON	Metastasis	15
2 148.0	BUZOS001A1 - externo	17	1	27-mar-2024	ADENOCARCINOMA DE PULMON	MASC	60	ADENOCARCINOMA DE PULMON	Centinela pulmonar no metastatizante	15.1
1 108.0	BUZOS001A1	000	1	16-04-2024	ADENOCARCINOMA EPIDERMICO DE PULMON	MASC	55	ADENOCARCINOMA EPIDERMICO DE PULMON	Centinela pulmonar no metastatizante	15.1

Figure 17: Demonstration of sorting values

### 3.3.2 Filtering Rows

In Figure 18, only records corresponding to male patients have been filtered.

Numero de chip	Numero de Hospital	NIC	Estado actual	Fecha de ingreso	Diagnostico	Sexo	Porcentaje tumoral	Celulas	Diagnostico	Nombre
2 148.0	BUZOS001A1 - externo	17	1	27-mar-2024	ADENOCARCINOMA DE PULMON	MASC	60	ADENOCARCINOMA DE PULMON	Centinela pulmonar no metastatizante	15.1
1 108.0	BUZOS001A1	000	1	16-04-2024	ADENOCARCINOMA EPIDERMICO DE PULMON	MASC	55	ADENOCARCINOMA EPIDERMICO DE PULMON	Centinela pulmonar no metastatizante	15.1

Figure 18: Demonstration of filtering rows

### 3.4 Exporting Data

In this example, the data has been extracted into a CSV document 19, and the visualization of the downloaded file is shown in 20.

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Figure 19: Demonstration of exporting data

Id	Nombre	Sexo	Fecha de diagnóstico	Diagnóstico	Sexo	Porcentaje	Número	Diagnóstico
1	100.0	BL000000-A1	17	1	10 mar 2024	ADENOCARCINOMA EPIDIDIMO DE PULMON	MASC	65
2	100.0	BL000000-A1	17	1	27 mar 2024	ADENOCARCINOMA EPIDIDIMO DE PULMON	MASC	65
3	100.0	BL000000-A2	17	1	28 mar 2024	ADENOCARCINOMA EPIDIDIMO DE PULMON	FEM	75

Figure 20: Demonstration of exporting data

### 3.5 Creating Patient History

In Figure 21, the records of the analyzed documents are shown in the MongoDB Compass application.

Id	Nombre	Sexo	Fecha de diagnóstico	Diagnóstico	Sexo	Porcentaje	Número	Diagnóstico
1	100.0	BL000000-A1	17	1	10 mar 2024	ADENOCARCINOMA EPIDIDIMO DE PULMON	MASC	65
2	100.0	BL000000-A1	17	1	27 mar 2024	ADENOCARCINOMA EPIDIDIMO DE PULMON	MASC	65
3	100.0	BL000000-A2	17	1	28 mar 2024	ADENOCARCINOMA EPIDIDIMO DE PULMON	FEM	75

Figure 21: Demonstration of creating patient history

## References

- [CRAN, 2024] CRAN (2024). Download r-4.4.0 for windows. the r-project for statistical computing. <https://cran.r-project.org/bin/windows/base/>. Acceso realizado el 5 de junio de 2024.
- [GitHub, 2024] GitHub (2024). Clonar un repositorio - documentación de github. <https://docs.github.com/es/repositories/creating-and-managing-repositories/cloning-a-repository>. Acceso realizado el 11 de junio de 2024.
- [MongoDB, 2024] MongoDB (2024). Download mongodb atlas cli — mongodb. <https://www.mongodb.com/try/download/atlascli>. Acceso realizado el 5 de junio de 2024.
- [Posit, 2024] Posit (2024). Rstudio desktop - posit. <https://posit.co/download/rstudio-desktop/>. Acceso realizado el 5 de junio de 2024.