

# Basic skills in R: part II

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# Learning objectives

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- **Importing data from excel into the R environment**
- **Basic data management**
  - Management of numerical variables
  - Manipulation of categorical variables

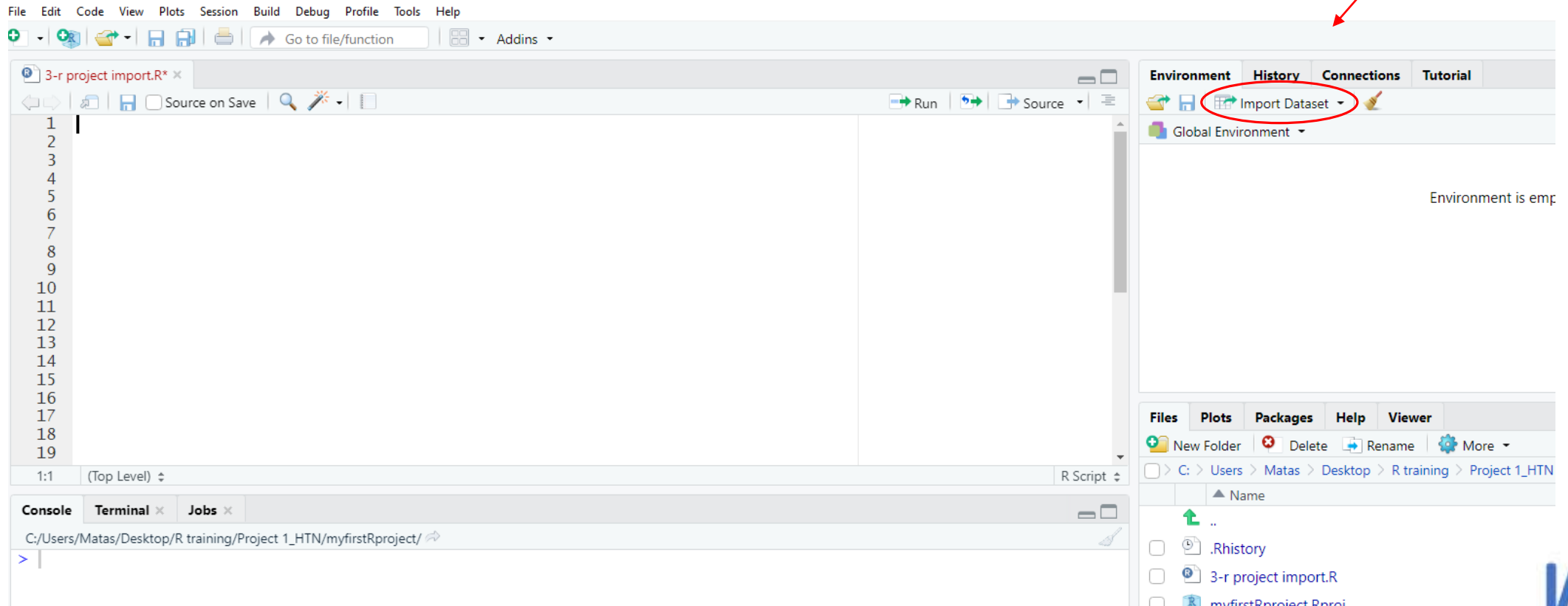
# Scenario

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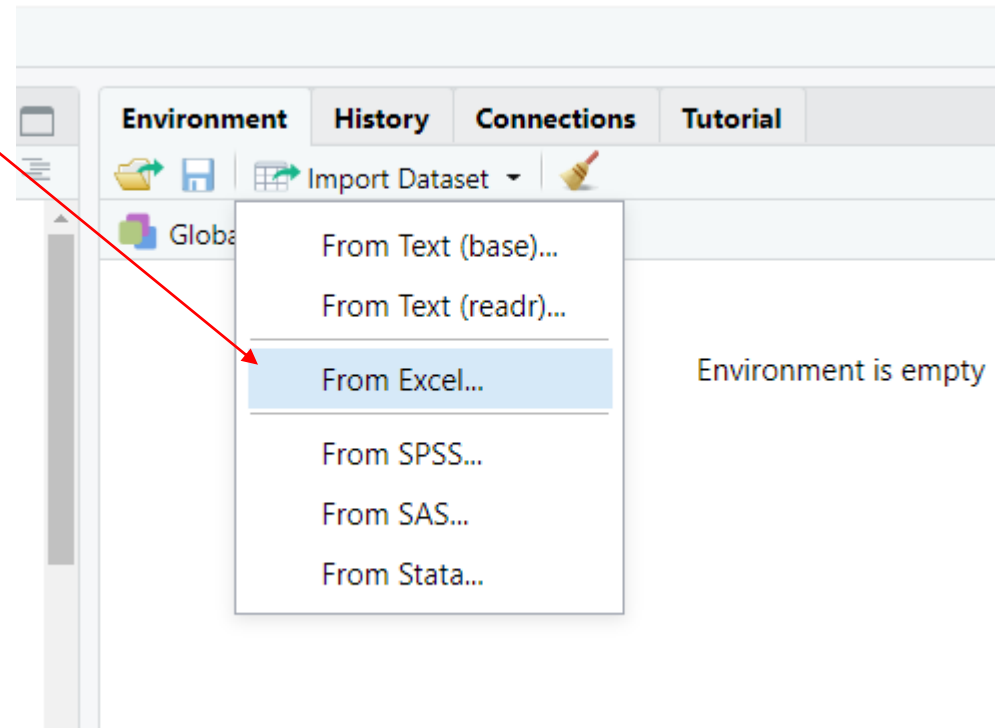
- **Imagine that your colleague sends you a patient dataset in excel format and asks you to perform exploratory data analysis.**
- **Let's go through how to upload a file in RStudio**

# How to upload an excel file in R

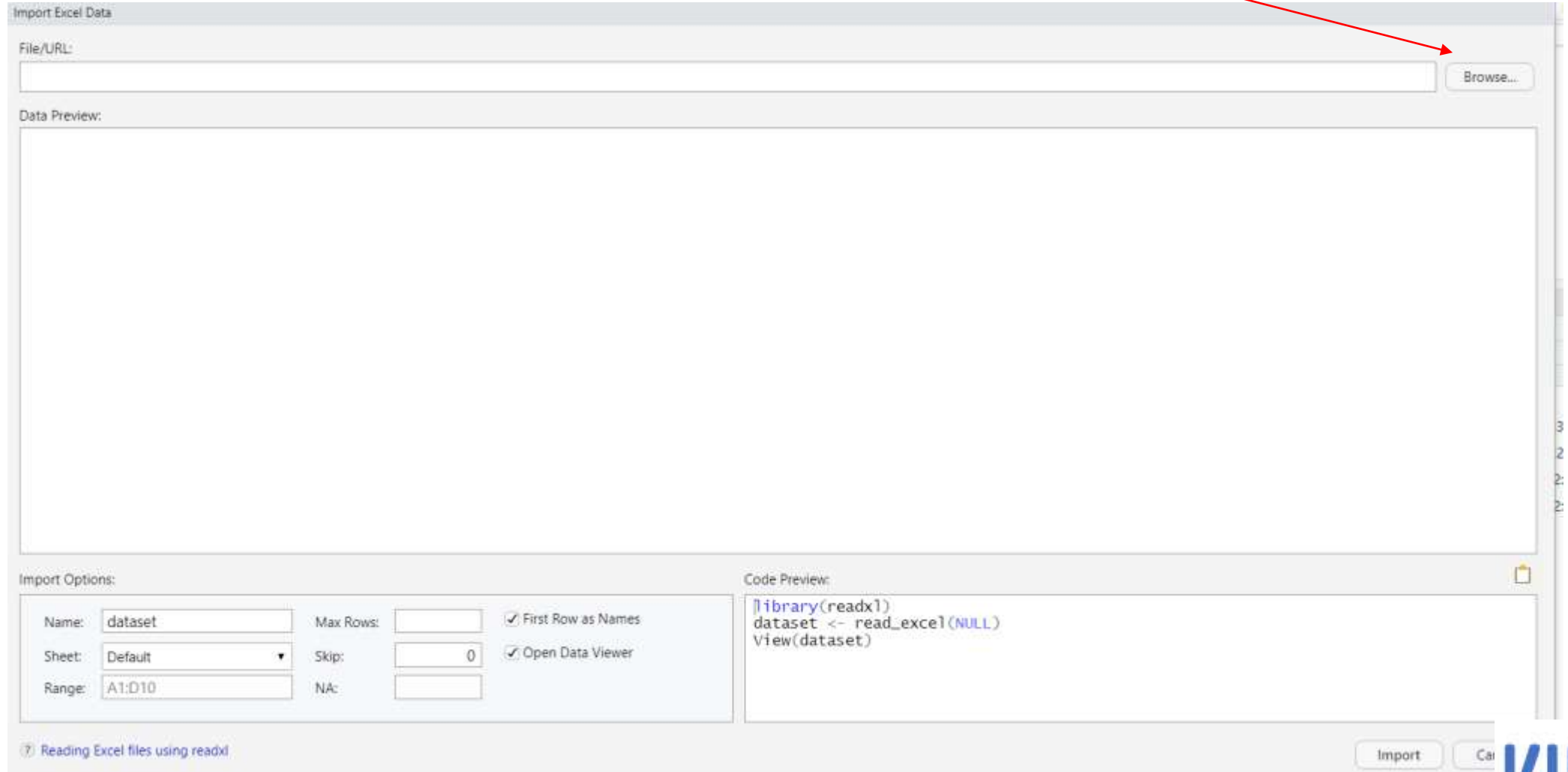
The easiest way is to click “import dataset”



Select "from Excel"



Select “browse” and search for your excel file. Make sure it’s saved on your computer!



Import Excel Data

File/URL:

Data Preview:

Import Options:

Name: <input type="text" value="dataset"/>	Max Rows: <input type="text"/>	<input checked="" type="checkbox"/> First Row as Names
Sheet: <input type="text" value="Default"/>	Skip: <input type="text" value="0"/>	<input checked="" type="checkbox"/> Open Data Viewer
Range: <input type="text" value="A1:D10"/>	NA: <input type="text"/>	

Code Preview:

```
library(readxl)
dataset <- read_excel(NULL)
View(dataset)
```

[\(?\) Reading Excel files using readxl](#)

Once you select your excel file, you will see the preview of the data. Next, select “import”.

Import Excel Data

File/URL:  
C:/Users/User/Box/\_Courses/R Class/R data management/Data sets/pat\_info.xlsx Update

Data Preview:

ID <small>(double)</small>	Age <small>(double)</small>	Sex <small>(character)</small>	HTN_Med <small>(double)</small>	Race <small>(double)</small>
1	50	M	0	1
2	67	M	1	3
3	75	F	1	2
4	31	F	0	1
5	29	F	0	1
6	74	M	0	4
7	58	F	1	2
8	41	M	0	3
9	86	M	1	4
10	22	M	1	1

Previewing first 50 entries.

Import Options:

Name:  Max Rows:   
Sheet:  Skip:   
Range:  NA:

First Row as Names  
 Open Data Viewer

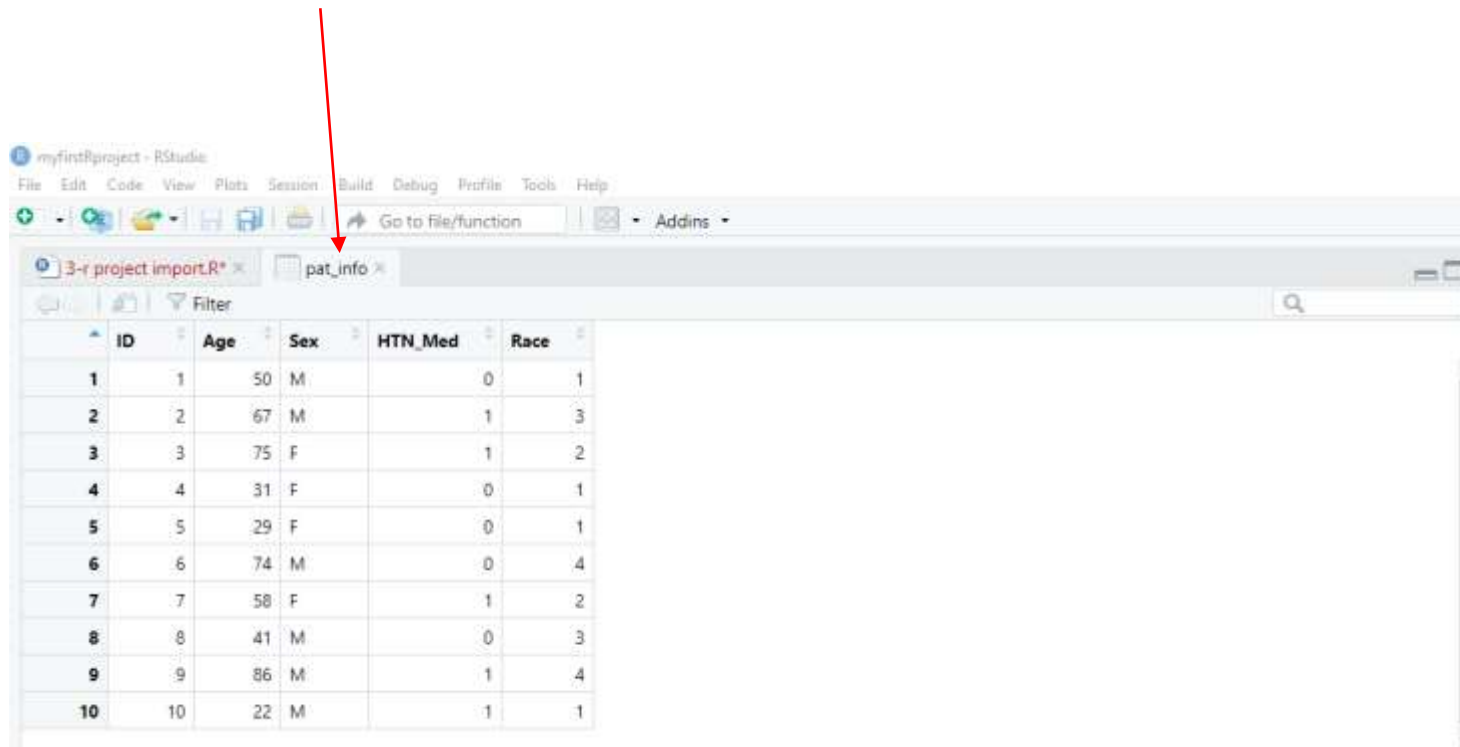
Code Preview:

```
library(readxl)
pat_info <- read_excel("C:/Users/Matas/Box/_Courses/R Class/R data management/Data
sets/pat_info.xlsx")
View(pat_info)
```

Import

Reading Excel files using readxl

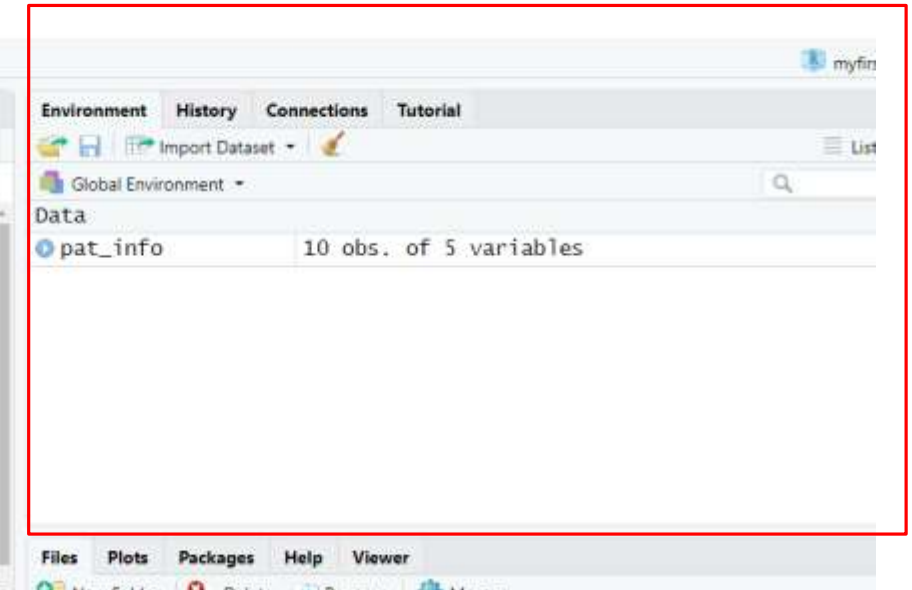
You see a new tab that has opened your dataset in R.



The screenshot shows the RStudio interface with a new tab titled 'pat\_info' active. A red arrow points to this tab. The main window displays a data table with the following columns: ID, Age, Sex, HTN\_Med, and Race. The data is as follows:

ID	Age	Sex	HTN_Med	Race
1	1	50 M	0	1
2	2	67 M	1	3
3	3	75 F	1	2
4	4	31 F	0	1
5	5	29 F	0	1
6	6	74 M	0	4
7	7	58 F	1	2
8	8	41 M	0	3
9	9	86 M	1	4
10	10	22 M	1	1

The dataset is officially in the R environment as shown here. Name of dataset is "pat\_info"



The screenshot shows the RStudio Environment pane. A red arrow points to the 'pat\_info' dataset listed under the 'Data' section. The dataset is described as having 10 observations and 5 variables.

Environment	History	Connections	Tutorial
Global Environment			
Data			
pat_info			



The screenshot shows the RStudio interface. At the top, there is a menu bar with options: File, Edit, Code, View, Plots, Session, Build, Debug, Profile, Tools, Help. Below the menu bar is a toolbar with icons for file operations and a search bar labeled 'Go to file/function'. The main workspace is divided into two panes. The left pane shows a list of files, with '3-r project import.R\*' selected and highlighted in a blue box. A blue arrow points from the text 'Click the R script tab to begin typing your code' to the '3-r project import.R\*' tab. The right pane is empty. At the bottom, there is a status bar with '1:1 (Top Level)' and 'R Script'. Below the status bar are tabs for 'Console', 'Terminal', and 'Jobs'.

3-r project import.R\* x pat\_info x

Source on Save Run Source

1  
2  
3  
4  
5  
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17  
18  
19

1:1 (Top Level) R Script

Console Terminal Jobs

Click the R script tab to begin typing your code

# Basic data exploration

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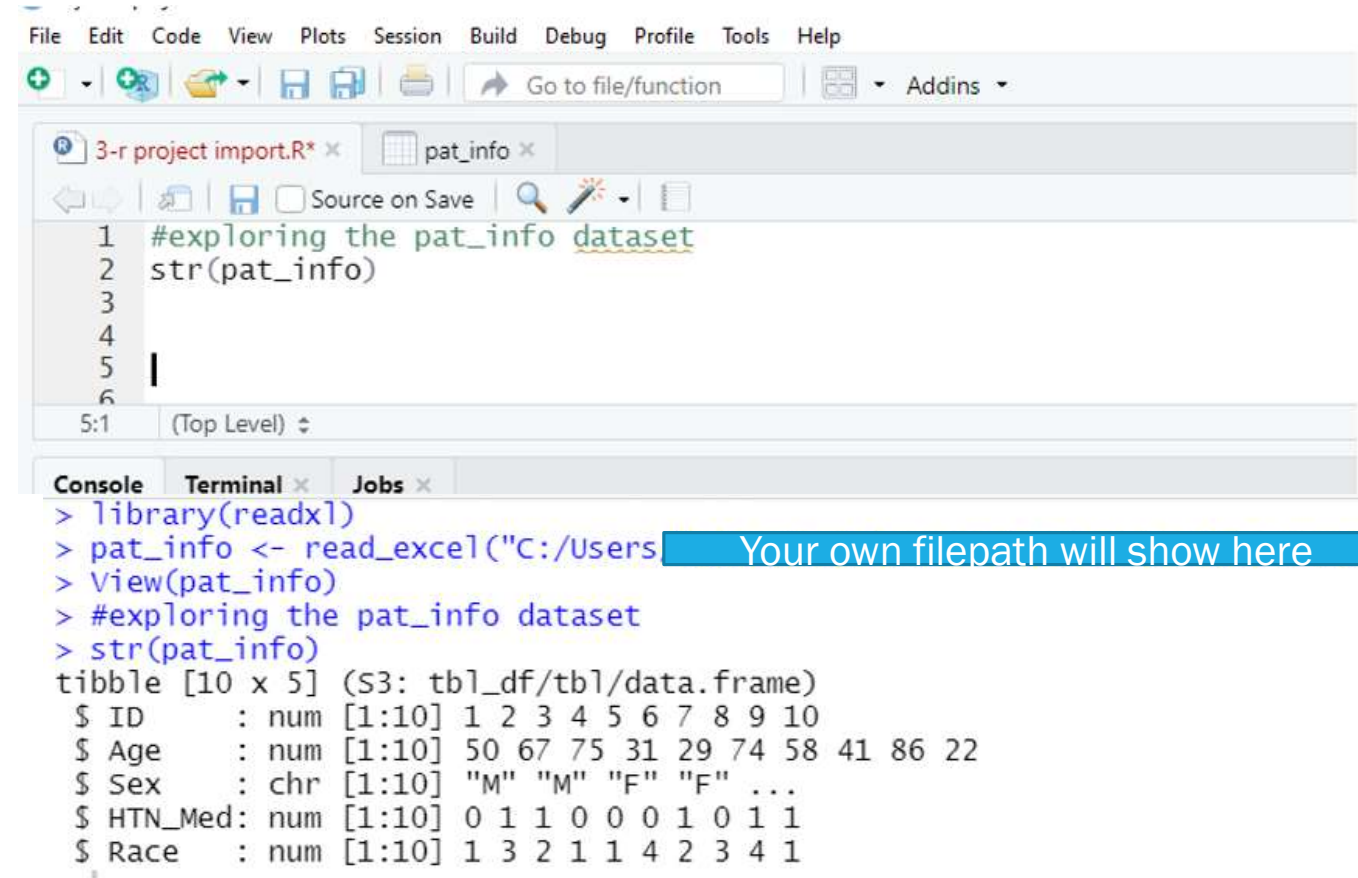
**NUMERICAL VARIABLES**

# Exploring the dataset: str function

str function: displays the structure of a R object (in this case a data frame).

Output is displayed here

Note: everything in R is case-sensitive




The screenshot shows the RStudio environment. The script editor contains the following code:

```
1 #exploring the pat_info dataset
2 str(pat_info)
3
4
5
6
```

The console shows the following output:

```
> library(readxl)
> pat_info <- read_excel("C:/Users/Your own filepath will show here")
> View(pat_info)
> #exploring the pat_info dataset
> str(pat_info)
tibble [10 x 5] (S3: tbl_df/tbl/data.frame)
 $ ID       : num [1:10] 1 2 3 4 5 6 7 8 9 10
 $ Age      : num [1:10] 50 67 75 31 29 74 58 41 86 22
 $ Sex      : chr [1:10] "M" "M" "F" "F" ...
 $ HTN_Med  : num [1:10] 0 1 1 0 0 0 1 0 1 1
 $ Race     : num [1:10] 1 3 2 1 1 4 2 3 4 1
```

Down here we see that the data frame weights contains 5 variables: 4 numerical and 1 character



The head function in R environment displays the first observations of a dataframe or variable. By specifying the “n” option, you control how many observations will be displayed.

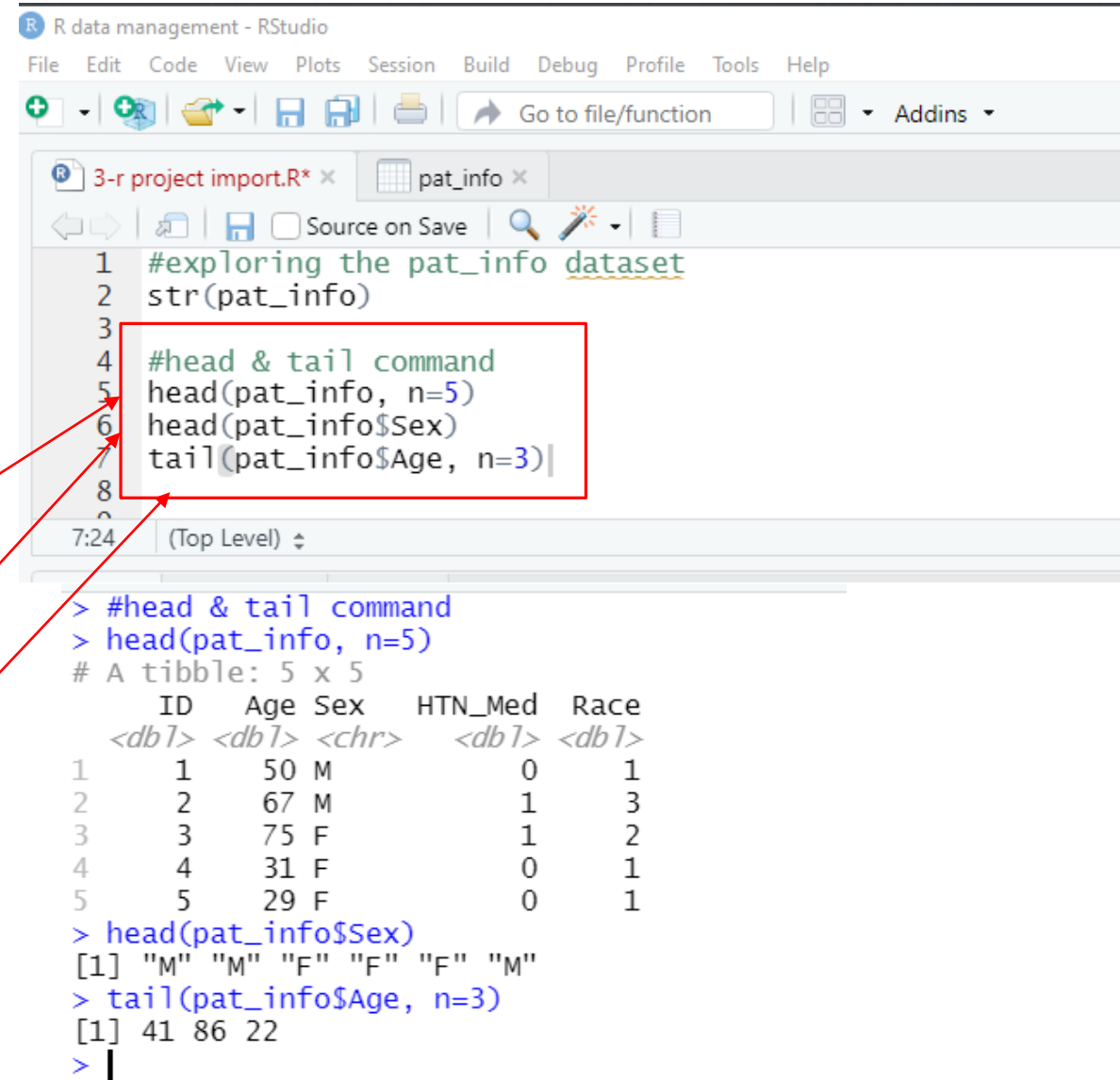
Note: If you don't specify the n option, the first six observations will be displayed by default.

The tail function in R, provides the last observations. Similarly, the last 6 are the default unless specified by the “n”.

Here, I'm requesting the first five observations from the entire dataset pat\_info

Next, I specify that I want to display the first default observations for the variable Sex in the pat\_info dataset

Finally, I am requesting the last three observations of the variable Age.



```
R data management - RStudio
File Edit Code View Plots Session Build Debug Profile Tools Help
Go to file/function
3-r project import.R* x pat_info x
Source on Save
1 #exploring the pat_info dataset
2 str(pat_info)
3
4 #head & tail command
5 head(pat_info, n=5)
6 head(pat_info$Sex)
7 tail(pat_info$Age, n=3)
8
7:24 (Top Level)
> #head & tail command
> head(pat_info, n=5)
# A tibble: 5 x 5
  ID Age Sex HTN_Med Race
<dbl> <dbl> <chr> <dbl> <dbl>
1 1 50 M 0 1
2 2 67 M 1 3
3 3 75 F 1 2
4 4 31 F 0 1
5 5 29 F 0 1
> head(pat_info$Sex)
[1] "M" "M" "F" "F" "F" "M"
> tail(pat_info$Age, n=3)
[1] 41 86 22
> |
```

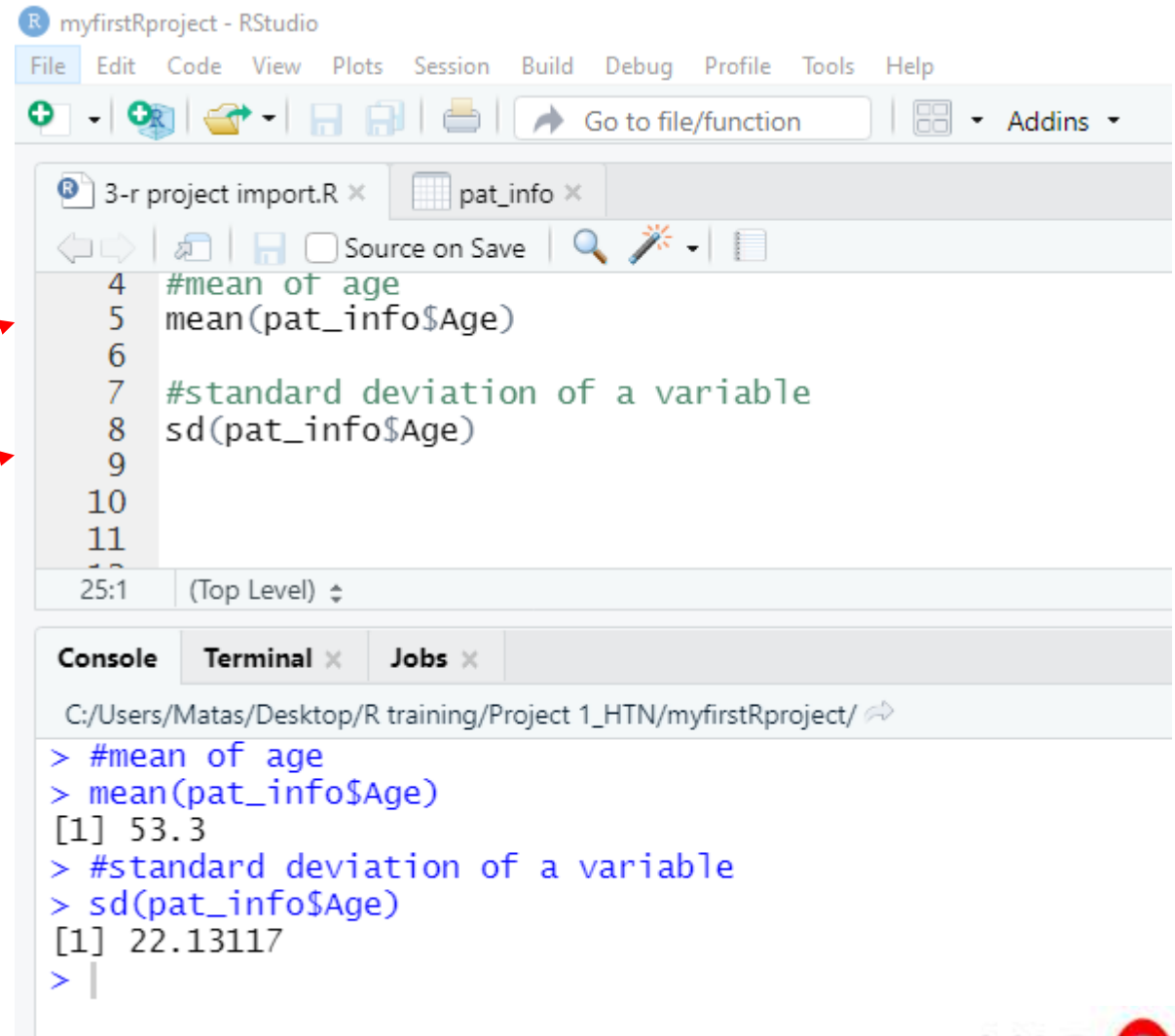
# Identifying the mean & standard deviation of a variable

Type:

Mean(pat\_info\$Age)

sd(pat\_info\$Age)

The first part represents the name of the data frame and the second part after the dollar sign represents the specific variable



The screenshot shows the RStudio interface with a script editor and a console. The script editor contains the following R code:

```
4 #mean of age
5 mean(pat_info$Age)
6
7 #standard deviation of a variable
8 sd(pat_info$Age)
9
10
11
```

The console shows the output of the code:

```
> #mean of age
> mean(pat_info$Age)
[1] 53.3
> #standard deviation of a variable
> sd(pat_info$Age)
[1] 22.13117
> |
```

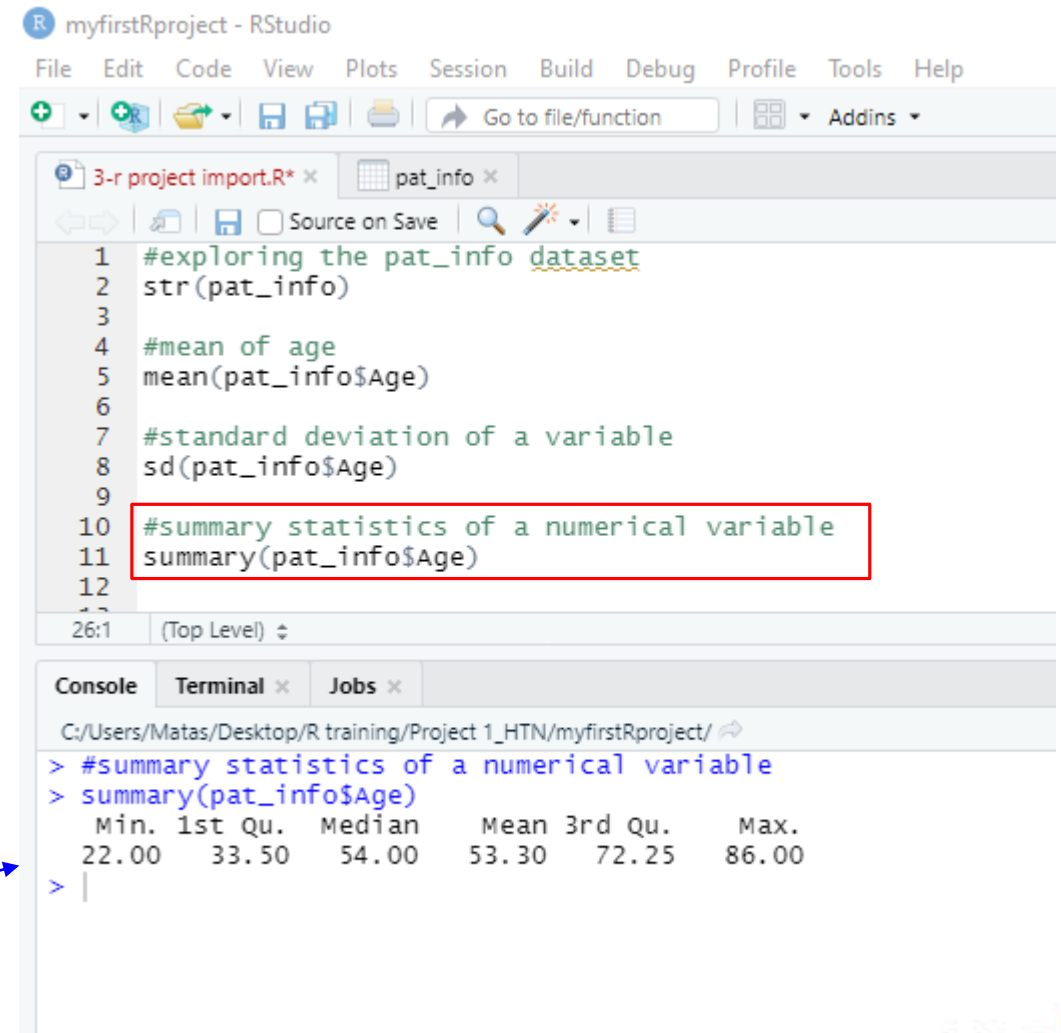
Red arrows point from the text on the left to the corresponding lines in the R code.

# summary of a variable

Type: `summary(pat_info$Age)` →

The first part represents the name of the data frame and the second part after the dollar sign represents the specific variable

Summary statistics are here →



The screenshot shows the RStudio interface with a script editor and a console. The script editor contains the following R code:

```
1 #exploring the pat_info dataset
2 str(pat_info)
3
4 #mean of age
5 mean(pat_info$Age)
6
7 #standard deviation of a variable
8 sd(pat_info$Age)
9
10 #summary statistics of a numerical variable
11 summary(pat_info$Age)
12
```

The line `summary(pat_info$Age)` is highlighted with a red box. The console shows the output of this command:

```
> #summary statistics of a numerical variable
> summary(pat_info$Age)
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
22.00  33.50   54.00   53.30  72.25   86.00
> |
```

# Management of categorical variables

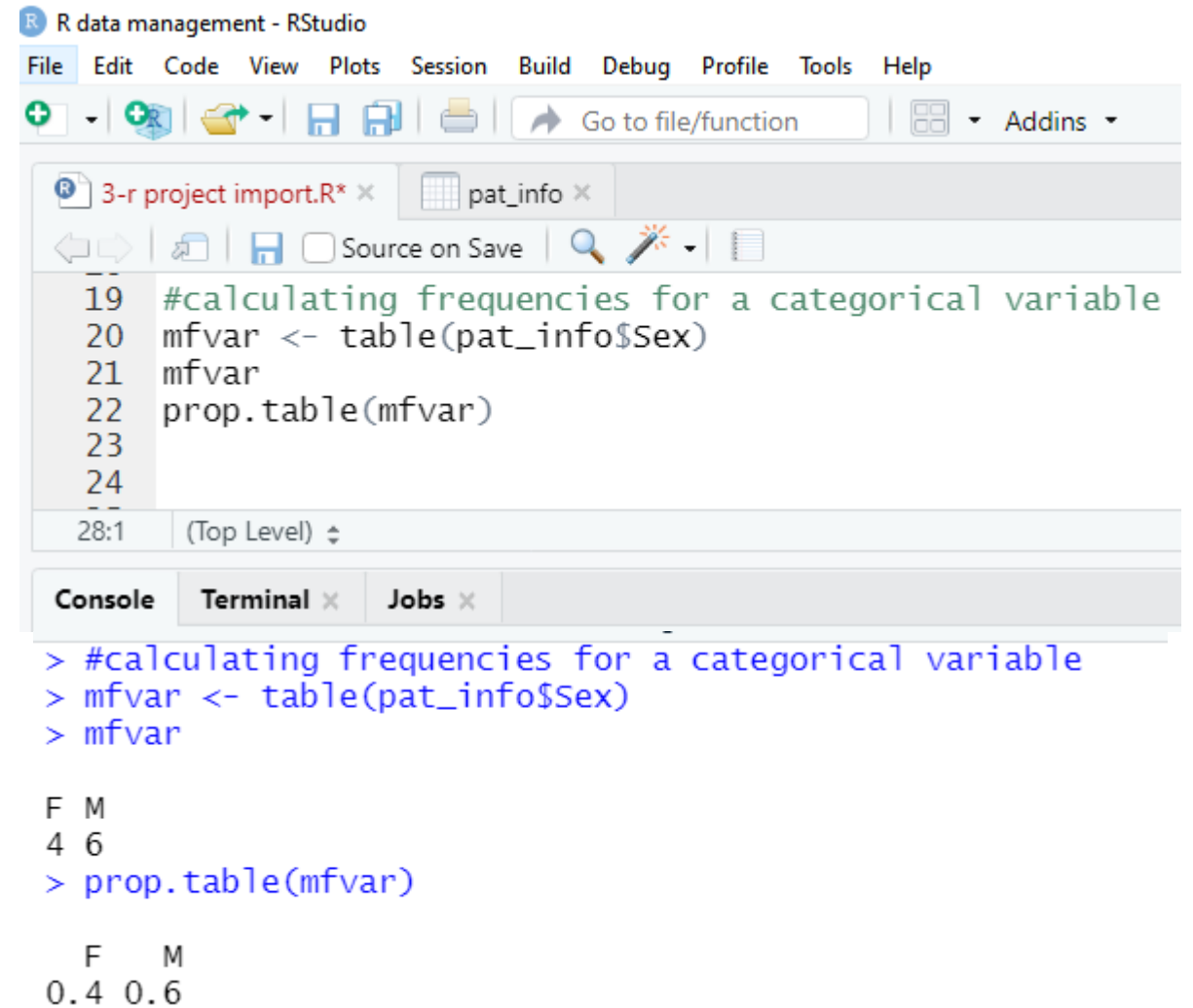
---

# Calculating a frequency of a categorical variable

We create a vector called "mfvar" to include the table counts of Sex.

The table() command will provide you the count of the variable. In the dataset, we have 4 females and 6 males.

The prop.table(table) command will provide you with the proportion. 40% females and 60% males.



```
R data management - RStudio
File Edit Code View Plots Session Build Debug Profile Tools Help
3-r project import.R* x pat_info x
Source on Save
19 #calculating frequencies for a categorical variable
20 mfvar <- table(pat_info$Sex)
21 mfvar
22 prop.table(mfvar)
23
24
28:1 (Top Level)
Console Terminal x Jobs x
> #calculating frequencies for a categorical variable
> mfvar <- table(pat_info$Sex)
> mfvar
F M
4 6
> prop.table(mfvar)
      F      M
0.4 0.6
```



# Adding labels

Notice we have a variable called race which identifies patients' self-reported race as:

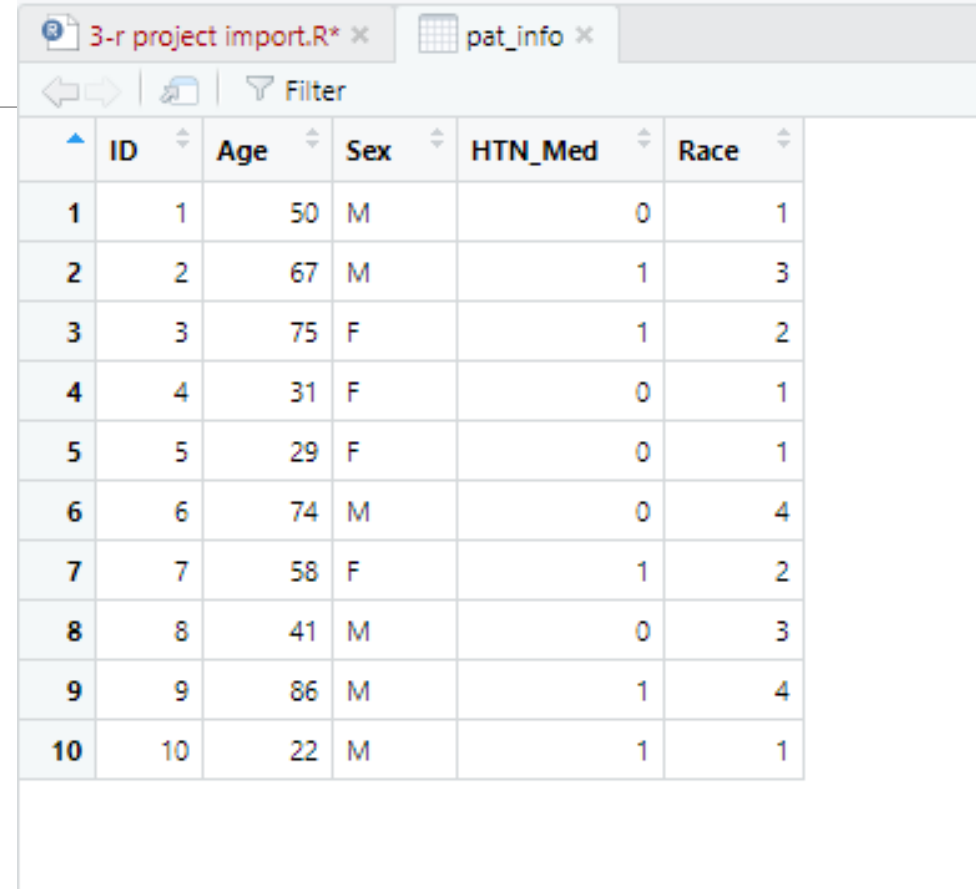
1= Non-Hispanic White (NHW)

2= Non-Hispanic Black (NHB)

3= Hispanic (HIS)

4= Other (OTH)

We will demonstrate how to add labels to the numerical values (1-4) of the qualitative variable "race".



	ID	Age	Sex	HTN_Med	Race
1	1	50	M	0	1
2	2	67	M	1	3
3	3	75	F	1	2
4	4	31	F	0	1
5	5	29	F	0	1
6	6	74	M	0	4
7	7	58	F	1	2
8	8	41	M	0	3
9	9	86	M	1	4
10	10	22	M	1	1

# Adding labels

We call out the variable Race by locating it in the pat\_info dataset, convert it to a factor with 4 levels and label them in order as NHW, NHB, HIS, OTH.

Recall for categorical variables, we use factors. This is especially useful for statistical modeling.



The screenshot shows the RStudio interface. The top menu bar includes File, Edit, Code, View, Plots, Session, Build, Debug, Profile, Tools, and Help. Below the menu is a toolbar with icons for file operations and a search bar. The main editor window shows a script with the following code:

```
25  
26  
27 #labels  
28 pat_info$Race <- factor(pat_info$Race, levels=c(1,2,3,4), labels=c("NHW", "NHB", "HIS", "OTH"))  
29  
30
```

At the bottom, the Console window shows the execution of the code:

```
> #labels  
> pat_info$Race <- factor(pat_info$Race, levels=c(1,2,3,4), labels=c("NHW", "NHB", "HIS", "OTH"))  
> |
```



# Management of both categorical and numerical variables

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**SUBGROUP ANALYSES**

# Subgroup analysis

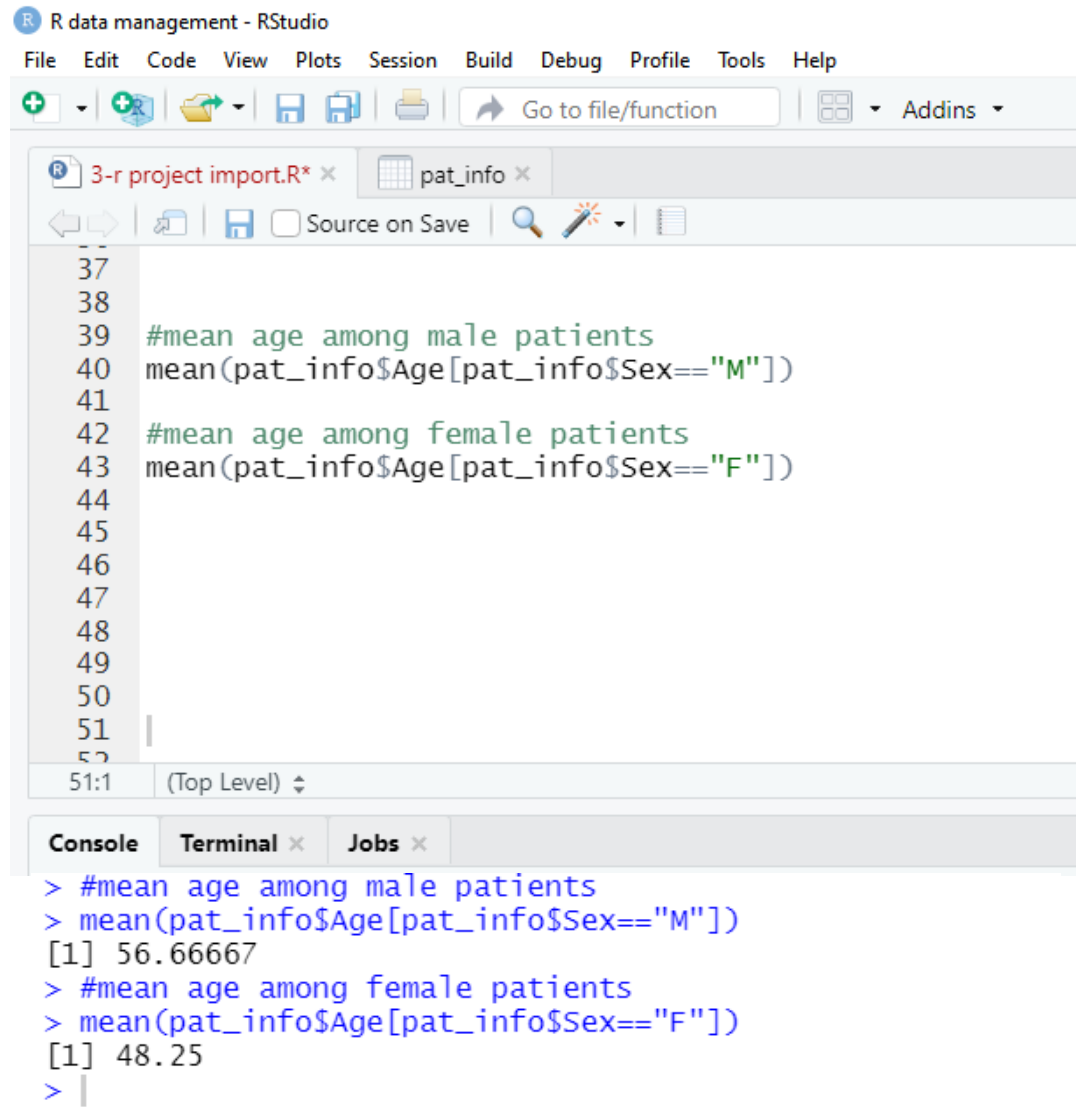
---

- Suppose you are interested in identifying the mean age of male and female patients.
- Code is provided on the next slide

This first part will give you the mean age of the patients. Notice how you have to include the dataframe name both times to identify the numerical variable, age and the categorical variable "M"

The same applies for "F"

The results show down here.



```
R data management - RStudio
File Edit Code View Plots Session Build Debug Profile Tools Help
Go to file/function
3-r project import.R* pat_info
Source on Save
37
38
39 #mean age among male patients
40 mean(pat_info$Age[pat_info$Sex=="M"])
41
42 #mean age among female patients
43 mean(pat_info$Age[pat_info$Sex=="F"])
44
45
46
47
48
49
50
51
52
51:1 (Top Level)
Console Terminal x Jobs x
> #mean age among male patients
> mean(pat_info$Age[pat_info$Sex=="M"])
[1] 56.66667
> #mean age among female patients
> mean(pat_info$Age[pat_info$Sex=="F"])
[1] 48.25
> |
```

# Criterion-based selection

---

Suppose you are interested in obtaining the ID numbers of patients above the age of 60.

Type `pat_info$ID[pat_info$Age > 60]`

The first part depicts the main variable of interest and the part inside the brackets depicts the specific criteria.

IDs are displayed down here:

The screenshot shows the RStudio interface. At the top right, a data table is displayed with columns: ID, Age, Sex, HTN\_Med, and Race. The table contains 10 rows of patient data. Rows 2, 3, 6, and 9 are highlighted with red boxes, indicating patients whose ages are greater than 60. Below the table, the R console shows the execution of the command `pat_info$ID[pat_info$Age > 60]`, resulting in the output `[1] 2 3 6 9`.

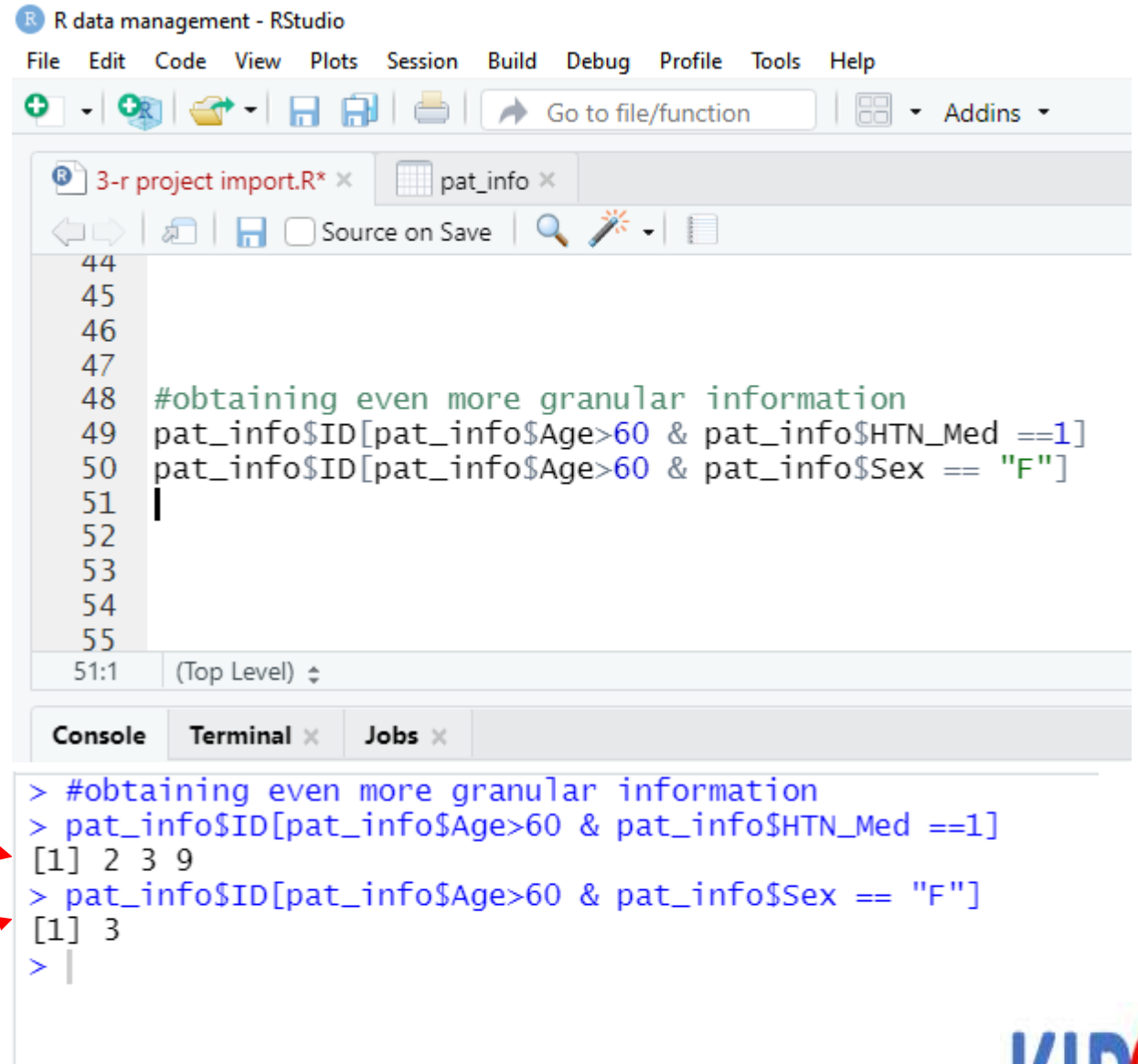
ID	Age	Sex	HTN_Med	Race	
1	1	50	M	0	1
2	2	67	M	1	3
3	3	75	F	1	2
4	4	31	F	0	1
5	5	29	F	0	1
6	6	74	M	0	4
7	7	58	F	1	2
8	8	41	M	0	3
9	9	86	M	1	4
10	10	22	M	1	1

```
R data management - RStudio
File Edit Code View Plots Session Build Debug Profile Tools Help
3-r project import.R* x pat_info x
38
39
40 #criterion based selection
41 pat_info$ID[pat_info$Age > 60]
42
43
44
45
46
47
48
49
53:1 (Top Level)
Console Terminal x Jobs x
> #criterion based selection
> pat_info$ID[pat_info$Age > 60]
[1] 2 3 6 9
> |
```



Obtaining IDs for patients above the age of 60 AND an indicator for being on hypertensive medication. Output: IDs 2,3,9

Obtaining IDs for patients about the age of 60 and female sex. Output: ID #3



The screenshot shows the RStudio interface with the following content:

```
44
45
46
47
48 #obtaining even more granular information
49 pat_info$ID[pat_info$Age>60 & pat_info$HTN_Med ==1]
50 pat_info$ID[pat_info$Age>60 & pat_info$Sex == "F"]
51 |
52
53
54
55
```

The console output shows the execution of the two R commands:

```
> #obtaining even more granular information
> pat_info$ID[pat_info$Age>60 & pat_info$HTN_Med ==1]
[1] 2 3 9
> pat_info$ID[pat_info$Age>60 & pat_info$Sex == "F"]
[1] 3
> |
```

Two red arrows point from the text on the left to the corresponding lines in the console output.

# Class Exercise #1: Module 03

---

The data in the Table below pertains to five patients with certain characteristics.

ID	Age	Sex	Diabetes	Race
1	55	M	1	2
2	70	F	0	1
3	40	M	0	1
4	20	M	1	3
5	63	F	1	2

# Class Exercise #1: Module 03

---

Do the following:

- a. Create the relevant variables in R
- b. Frame the variables into a dataset
- c. Calculate the frequency and proportion of individuals with diabetes by sex

# Class Exercise #1: Module 03

---

d. Compare the mean age of individuals with versus those without diabetes

e. Label the race variable as follows: 1 = “White”, 2 = “Black”, and 3 = “Hispanic”.

# Class Exercise #2: Module 03

The following Table contains information on age (in years), marital status (1= married, 2=divorced, 3=single), obesity status (1=obese,2=overweight, 3=normal weight) and race (1=White,2=Black, 3=Other) of ten patients as follows:

ID	Age	Marital status	Obesity status	Race
1	45	2	3	2
2	28	3	1	2
3	59	1	2	1
4	33	3	2	3
5	39	1	1	1
6	52	2	3	1
7	61	1	3	2
8	39	3	2	1
9	46	1	1	2
10	29	3	1	2

# Class Exercise #2: Module 03

---

1. Create a vector for each of the variables in the Table
  - ii. Compile/frame the vectors into a dataset and give that dataset a name
  - iii. Change the numeric labels of variables marital status, obesity status and race to character labels
  - iv. Create a 3 by 3 table between race and obesity
  - v. Create a 3 by 3 table between race and marital status

# Class Exercise #2: Module 03

---

- vi. Create a 3 by 3 table between marital status and obesity
- vii. Find the mean age for patients who are obese, for Blacks, Whites, and for those who are single
- viii. How many patients are less than 40 years and obese?
- ix. How many patients are black and less than 40?
- x. How many patients are white and older than 40?

# Summary

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➤ In this lecture you learned:

- How to import excel files into R environment
- Numerical data manipulation
- Categorical data manipulation
- Basic sub-analyses incorporating both categorical and numerical values

➤ Next, we will discuss packages in R and how to use them.