Subsetting Data in R

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Overview

We showed one way to read data into R using read_csv() and read.csv(). In this module, we will show you how to:

- 1. Select specific elements of an object by an index or logical condition
- 2. Subset rows of a data.frame
- 3. Subset columns of a data.frame
- 4. Add/remove new columns to a data.frame
- 5. Order the columns of a data.frame
- 6. Order the rows of a data.frame

Setup

We will show you how to do each operation in base R then show you how to use the dplyr package to do the same operation (if applicable).

Many resources on how to use dplyr exist and are straightforward:

- https://cran.rstudio.com/web/packages/dplyr/vignettes/
- https://stat545-ubc.github.io/block009_dplyr-intro.html
- https://www.datacamp.com/courses/dplyr-data-manipulation-rtutorial

The dplyr package also interfaces well with tibbles.

Select specific elements using an index

Often you only want to look at subsets of a data set at any given time. As a review, elements of an R object are selected using the brackets ([and]).

For example, x is a vector of numbers and we can select the second element of x using the brackets and an index (2):

Select specific elements using an index

We can select the fifth or second AND fifth elements below:

```
x = c(1, 2, 4, 8, 10)
x[5]

[1] 10

x[c(2,5)]

[1] 2 10
```

Subsetting by deletion of entries

You can put a minus (-) before integers inside brackets to remove these indices from the data.

```
x[-2] # all but the second
[1] 1 4 8 10
```

Note that you have to be careful with this syntax when dropping more than 1 element:

```
x[-c(1,2,3)] # drop first 3
[1] 8 10
# x[-1:3] # shorthand. R sees as -1 to 3
x[-(1:3)] # needs parentheses
[1] 8 10
```

Select specific elements using logical operators

What about selecting rows based on the values of two variables? We use logical statements. Here we select only elements of \mathbf{x} greater than 2:

```
x
[1] 1 2 4 8 10
x > 2
[1] FALSE FALSE TRUE TRUE TRUE
x[ x > 2 ]
[1] 4 8 10
```

Select specific elements using logical operators

You can have multiple logical conditions using the following:

- &: AND
- · |: OR

[1] 4

$$x[x > 5 | x == 2]$$

[1] 2 8 10

which function

The which functions takes in logical vectors and returns the index for the elements where the logical value is TRUE.

```
which(x > 5 | x == 2) # returns index
[1] 2 4 5
x[ which(x > 5 | x == 2) ]
[1] 2 8 10
x[ x > 5 | x == 2 ]
[1] 2 8 10
```

Creating a data. frame to work with

Here we create a toy data.frame named df using random data:

Renaming Columns

Renaming Columns of a data.frame: base R

We can use the **colnames** function to directly reassign column names of **df**:

Renaming Columns of a data.frame: base R

We can assign the column names, change the ones we want, and then re-assign the column names:

library(dplyr)

Note, when loading dplyr, it says objects can be "masked". That means if you use a function defined in 2 places, it uses the one that is loaded in last.

For example, if we print filter, then we see at the bottom namespace:dplyr, which means when you type filter, it will use the one from the dplyr package.

```
filter

function (.data, ...)
{
    UseMethod("filter")
}
<environment: namespace:dplyr>
```

A filter function exists by default in the stats package, however. If you want to make sure you use that one, you use PackageName::Function with the colon-colon ("::") operator.

```
head(stats::filter,2)

1 function (x, filter, method = c("convolution", "recursive"),
2    sides = 2L, circular = FALSE, init = NULL)
```

This is important when loading many packages, and you may have some conflicts/masking:

To rename columns in dplyr, you use the rename command

Subsetting Columns

Subset columns of a data.frame:

We can grab the x column using the \$ operator.

df\$x

[1] 1 2 4 10 10

Subset columns of a data.frame:

We can also subset a data.frame using the bracket [,] subsetting.

For data.frames and matrices (2-dimensional objects), the brackets are [rows, columns] subsetting. We can grab the x column using the index of the column or the column name ("x")

```
df[, 1]
[1] 1 2 4 10 10
df[, "x"]
[1] 1 2 4 10 10
```

Subset columns of a data.frame:

We can select multiple columns using multiple column names:

```
df[, c("x", "y")]

x y
1 1 -0.2707606
2 2 -1.1179372
3 4 -1.3473558
4 10 0.4832675
5 10 0.1523950
```

Subset columns of a data.frame: dplyr

The select command from dplyr allows you to subset

```
x
1 1
2 2
3 4
4 10
5 10
```

Select columns of a data. frame: dplyr

The select command from dplyr allows you to subset columns of

```
select(df, x, x2)
  x x2
1 1 7
2 2 6
3 4 10
4 10 13
5 10 13
select(df, starts_with("x"))
  x x2
1 1 7
2 2 6
3 4 10
4 10 13
5 10 13
```

Subsetting Rows

Subset rows of a data.frame with indices:

Let's select **rows** 1 and 3 from **df** using brackets:

```
df[ c(1, 3), ]

x x2 y z
1 1 7 -0.2707606 6
3 4 10 -1.3473558 7
```

Subset rows of a data.frame:

Let's select the rows of df where the x column is greater than 5 or is equal to 2. Without any index for columns, all columns are returned:

Subset rows of a data.frame:

We can subset both rows and colums at the same time:

```
df[ df$x > 5 | df$x == 2, c("y", "z")]

y z
2 -1.1179372 4
4 0.4832675 10
5 0.1523950 5
```

Subset rows of a data.frame: dplyr

The command in dplyr for subsetting rows is filter. Try ?filter

filter(df,
$$x > 5 \mid x == 2$$
)

```
x x2 y z
1 2 6 -1.1179372 4
2 10 13 0.4832675 10
3 10 13 0.1523950 5
```

Note, no \$ or subsetting is necessary. R "knows" x refers to a column of df.

Subset rows of a data.frame: dplyr

By default, you can separate conditions by commas, and filter assumes these statements are joined by &

Combining filter and select

You can combine filter and select to subset the rows and columns, respectively, of a data.frame:

In R, the common way to perform multiple operations is to wrap functions around each other in a nested way such as above

Assigning Temporary Objects

One can also create temporary objects and reassign them:

```
df2 = filter(df, x > 2 & y < 0)

df2 = select(df2, y, z)
```

Piping - a new concept

There is another (newer) way of performing these operations, called "piping". It is becoming more popular as it's easier to read:

It is read: "take df, then filter the rows and then select y, z".

Adding/Removing Columns

Adding new columns to a data.frame: base R

You can add a new column, called newcol to df, using the \$ operator:

```
df$newcol = 5:1
df$newcol = df$x + 2
```

Removing columns to a data.frame: base R

You can remove a column by assigning to NULL:

```
df$newcol = NULL
or selecing only the columns that were not newcol:
df = df[, colnames(df) != "newcol"]
```

Adding new columns to a data.frame: base R

You can also "column bind" a data.frame with a vector (or series of vectors), using the cbind command:

```
cbind(df, newcol = 5:1)
```

```
x x2 y z newcol

1 1 7 -0.2707606 6 5

2 2 6 -1.1179372 4 4

3 4 10 -1.3473558 7 3

4 10 13 0.4832675 10 2

5 10 13 0.1523950 5 1
```

Adding columns to a data.frame: dplyr

The mutate function in dplyr allows you to add or replace columns of a data.frame:

```
mutate(df, newcol = 5:1)
  x x2 y z newcol
1 1 7 -0.2707606 6
2 2 6 -1.1179372 4
3 4 10 -1.3473558 7
4 10 13 0.4832675 10
5 10 13 0.1523950 5
print({df = mutate(df, newcol = x + 2)})
  x x2 y z newcol
1 1 7 -0.2707606 6
2 2 6 -1.1179372 4
3 4 10 -1.3473558 7
4 10 13 0.4832675 10 12
5 10 13 0.1523950 5
                     12
```

Removing columns to a data.frame: dplyr

The **NULL** method is still very common.

The select function can remove a column with a minus (-), much like removing rows:

```
select(df, -newcol)
```

```
x x2 y z

1 1 7 -0.2707606 6

2 2 6 -1.1179372 4

3 4 10 -1.3473558 7

4 10 13 0.4832675 10

5 10 13 0.1523950 5
```

Removing columns to a data.frame: dplyr

Remove newcol and y

Ordering columns

Ordering the columns of a data.frame: base R

We can use the colnames function to get the column names of df and then put newcol first by subsetting df using brackets:

Ordering the columns of a data.frame: dplyr

The select function can reorder columns. Put newcol first, then select the rest of columns:

Ordering rows

Ordering the rows of a data.frame: base R

We use the **order** function on a vector or set of vectors, in increasing order:

```
df[ order(df$x), ]

x x2 y z newcol
1 1 7 -0.2707606 6 3
2 2 6 -1.1179372 4 4
3 4 10 -1.3473558 7 6
4 10 13 0.4832675 10 12
5 10 13 0.1523950 5 12
```

Ordering the rows of a data.frame: base R

The decreasing argument will order it in decreasing order:

Ordering the rows of a data.frame: base R

You can pass multiple vectors, and must use the negative (using -) to mix decreasing and increasing orderings (sort increasing on \mathbf{x} and decreasing on \mathbf{y}):

Ordering the rows of a data.frame: dplyr

The arrange function can reorder rows By default, arrange orders in ascending order:

```
arrange(df, x)
```

```
x x2 y z newcol

1 1 7 -0.2707606 6 3

2 2 6 -1.1179372 4 4

3 4 10 -1.3473558 7 6

4 10 13 0.4832675 10 12

5 10 13 0.1523950 5 12
```

Ordering the rows of a data. frame: dplyr

Use the desc to arrange the rows in descending order:

```
arrange(df, desc(x))
```

```
x x2 y z newcol

1 10 13 0.4832675 10 12

2 10 13 0.1523950 5 12

3 4 10 -1.3473558 7 6

4 2 6 -1.1179372 4 4

5 1 7 -0.2707606 6 3
```

Ordering the rows of a data.frame: dplyr

It is a bit more straightforward to mix increasing and decreasing orderings:

```
arrange(df, x, desc(y))
```

```
x x2 y z newcol

1 1 7 -0.2707606 6 3

2 2 6 -1.1179372 4 4

3 4 10 -1.3473558 7 6

4 10 13 0.4832675 10 12

5 10 13 0.1523950 5 12
```

Transmutation

The transmute function in dplyr combines both the mutate and select functions. One can create new columns and keep the only the columns wanted: