Overview

We showed one way to read data into R using `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. Renaming columns of a `data.frame`
3. Subset rows of a `data.frame`
4. Subset columns of a `data.frame`
5. Add/remove new columns to a `data.frame`
6. Order the columns of a `data.frame`
7. 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://cran.rstudio.com/web/packages/dplyr/vignettes/)
- [https://stat545-ubc.github.io/block009_dplyr-intro.html](https://stat545-ubc.github.io/block009_dplyr-intro.html)
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):

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

\[
[1] 4
\]
Select specific elements using an index

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

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

```
[1] 10
```

```r
x[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

\[
\begin{array}{cccc}
1 & 4 & 8 & 10 \\
\end{array}
\]

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

\[ x[-c(1,2,3)] \]  # drop first 3

\[
\begin{array}{cccc}
8 & 10 \\
\end{array}
\]

\# x[-1:3]  # shorthand. R sees as -1 to 3
\[ x[-(1:3)] \]  # needs parentheses

\[
\begin{array}{cccc}
8 & 10 \\
\end{array}
\]
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 $x$ 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

\[ x[ x > 2 \& x < 5 ] \]

[1] 4

\[ x[ x > 5 \mid x == 2 ] \]

[1] 2 8 10
which function

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

```r
which(x > 5 | x == 2)  # returns index
```

```
[1] 2 4 5
```

```r
x[ which(x > 5 | x == 2) ]
```

```
[1] 2 8 10
```

```r
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:

```r
set.seed(2016) # reproducibility
df = data.frame(x = c(1, 2, 4, 10, 10),
                 x2 = rpois(5, 10),
                 y = rnorm(5),
                 z = rpois(5, 6)
)
```
Renaming Columns of a `data.frame`: base R

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

```r
colnames(df) = c("x", "X", "y", "z")
head(df)
```

```
x   X       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
```

```r
colnames(df) = c("x", "x2", "y", "z") #reset
```
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:

```r
cn = colnames(df)
cn[ cn == "x2"] = "X"
colnames(df) = cn
head(df)

   x  X     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
```

```r
colnames(df) = c("x", "x2", "y", "z") #reset
```
Renaming Columns of a `data.frame`: `dplyr`

```r
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`. 
Renaming Columns of a data.frame: dplyr

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.

```r
filter

function (.data, ...) {
    filter_(.data, .dots = lazyeval::lazy_dots(...))
}
<environment: namespace:dplyr>
```
Renaming Columns of a data.frame: 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.

```r
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:
Renaming Columns of a `data.frame`: dplyr

To rename columns in `dplyr`, you use the `rename` command

```r
df = dplyr::rename(df, X = x2)
head(df)
```

```
x  X        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
```

```r
df = dplyr::rename(df, x2 = X) # reset
```
Subset columns of a data.frame:

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

```r
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”)

```r
df[, 1]
```

```
[1] 1 2 4 10 10
```

```r
df[, "x"]
```

```
[1] 1 2 4 10 10
```
Subset columns of a `data.frame`:

We can select multiple columns using multiple column names:

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

<table>
<thead>
<tr>
<th></th>
<th>x</th>
<th>y</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>-0.2707606</td>
</tr>
<tr>
<td>2</td>
<td>2</td>
<td>-1.1179372</td>
</tr>
<tr>
<td>3</td>
<td>4</td>
<td>-1.3473558</td>
</tr>
<tr>
<td>4</td>
<td>10</td>
<td>0.4832675</td>
</tr>
<tr>
<td>5</td>
<td>10</td>
<td>0.1523950</td>
</tr>
</tbody>
</table>
Subset columns of a data.frame: dplyr

The `select` command from `dplyr` allows you to subset

```r
select(df, x)
```

<table>
<thead>
<tr>
<th>x</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
</tr>
<tr>
<td>2</td>
</tr>
<tr>
<td>3</td>
</tr>
<tr>
<td>4</td>
</tr>
<tr>
<td>4</td>
</tr>
<tr>
<td>5</td>
</tr>
</tbody>
</table>
Select columns of a data.frame: dplyr

The `select` command from `dplyr` allows you to subset columns of a data.frame.

```r
select(df, x, x2)
```

<p>| | | |</p>
<table>
<thead>
<tr>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>7</td>
</tr>
<tr>
<td>2</td>
<td>2</td>
<td>6</td>
</tr>
<tr>
<td>3</td>
<td>4</td>
<td>10</td>
</tr>
<tr>
<td>4</td>
<td>10</td>
<td>13</td>
</tr>
<tr>
<td>5</td>
<td>10</td>
<td>13</td>
</tr>
</tbody>
</table>

```r
select(df, starts_with("x"))
```

<p>| | | |</p>
<table>
<thead>
<tr>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>7</td>
</tr>
<tr>
<td>2</td>
<td>2</td>
<td>6</td>
</tr>
<tr>
<td>3</td>
<td>4</td>
<td>10</td>
</tr>
<tr>
<td>4</td>
<td>10</td>
<td>13</td>
</tr>
<tr>
<td>5</td>
<td>10</td>
<td>13</td>
</tr>
</tbody>
</table>
Subset rows of a data.frame with indices:

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

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

<table>
<thead>
<tr>
<th></th>
<th>x</th>
<th>x2</th>
<th>y</th>
<th>z</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>7</td>
<td>-0.2707606</td>
<td>6</td>
</tr>
<tr>
<td>3</td>
<td>4</td>
<td>10</td>
<td>-1.3473558</td>
<td>7</td>
</tr>
</tbody>
</table>
Subset rows of a data.frame:

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

\[
df[ \ df$x > 5 \ | \ df$x == 2, ]
\]

<table>
<thead>
<tr>
<th>x</th>
<th>x2</th>
<th>y</th>
<th>z</th>
</tr>
</thead>
<tbody>
<tr>
<td>2</td>
<td>2</td>
<td>6</td>
<td>-1.1179372</td>
</tr>
<tr>
<td>4</td>
<td>10</td>
<td>13</td>
<td>0.4832675</td>
</tr>
<tr>
<td>5</td>
<td>10</td>
<td>13</td>
<td>0.1523950</td>
</tr>
</tbody>
</table>
Subset rows of a data.frame:

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

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

<table>
<thead>
<tr>
<th></th>
<th>y</th>
<th>z</th>
</tr>
</thead>
<tbody>
<tr>
<td>2</td>
<td>-1.1179372</td>
<td>4</td>
</tr>
<tr>
<td>4</td>
<td>0.4832675</td>
<td>10</td>
</tr>
<tr>
<td>5</td>
<td>0.1523950</td>
<td>5</td>
</tr>
</tbody>
</table>
Subset rows of a data.frame: dplyr

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

```r
filter(df, x > 5 | x == 2)
```

<table>
<thead>
<tr>
<th>x</th>
<th>x2</th>
<th>y</th>
<th>z</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>2</td>
<td>6</td>
<td>-1.1179372</td>
</tr>
<tr>
<td>2</td>
<td>10</td>
<td>13</td>
<td>0.4832675</td>
</tr>
<tr>
<td>3</td>
<td>10</td>
<td>13</td>
<td>0.1523950</td>
</tr>
</tbody>
</table>

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 

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

```
x  x2       y   z
1  4  10 -1.347356 7
```

```r
filter(df, x > 2, y < 0)
```

```
x  x2       y   z
1  4  10 -1.347356 7
```
Combining filter and select

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

```r
select(filter(df, x > 2 & y < 0), y, z)
```

```
y z
1  -1.347356 7
```

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:

```python
df2 = filter(df, x > 2 & y < 0)
df2 = select(df2, y, z)
```
There is another (newer) way of performing these operations, called “piping”. It is becoming more popular as it’s easier to read:

```r
df %>% filter(x > 2 & y < 0) %>% select(y, z)
```

```
y z
1  -1.347356 7
```

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

```r
df$newcol = NULL
```

or selecting only the columns that were not `newcol`:

```r
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:

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

<table>
<thead>
<tr>
<th></th>
<th>x</th>
<th>x2</th>
<th>y</th>
<th>z</th>
<th>newcol</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>7</td>
<td>-0.2707606</td>
<td>6</td>
<td>5</td>
</tr>
<tr>
<td>2</td>
<td>2</td>
<td>6</td>
<td>-1.1179372</td>
<td>4</td>
<td>4</td>
</tr>
<tr>
<td>3</td>
<td>4</td>
<td>10</td>
<td>-1.3473558</td>
<td>7</td>
<td>3</td>
</tr>
<tr>
<td>4</td>
<td>10</td>
<td>13</td>
<td>0.4832675</td>
<td>10</td>
<td>2</td>
</tr>
<tr>
<td>5</td>
<td>10</td>
<td>13</td>
<td>0.1523950</td>
<td>5</td>
<td>1</td>
</tr>
</tbody>
</table>
Adding columns to a `data.frame`: `dplyr`

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

```r
mutate(df, newcol = 5:1)
```

<table>
<thead>
<tr>
<th>x</th>
<th>x2</th>
<th>y</th>
<th>z</th>
<th>newcol</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>7</td>
<td>-0.2707606</td>
<td>5</td>
</tr>
<tr>
<td>2</td>
<td>2</td>
<td>6</td>
<td>-1.1179372</td>
<td>4</td>
</tr>
<tr>
<td>3</td>
<td>4</td>
<td>10</td>
<td>-1.3473558</td>
<td>3</td>
</tr>
<tr>
<td>4</td>
<td>10</td>
<td>13</td>
<td>0.4832675</td>
<td>2</td>
</tr>
<tr>
<td>5</td>
<td>10</td>
<td>13</td>
<td>0.1523950</td>
<td>1</td>
</tr>
</tbody>
</table>

```r
print({df = mutate(df, newcol = x + 2)})
```

<table>
<thead>
<tr>
<th>x</th>
<th>x2</th>
<th>y</th>
<th>z</th>
<th>newcol</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>7</td>
<td>-0.2707606</td>
<td>3</td>
</tr>
<tr>
<td>2</td>
<td>2</td>
<td>6</td>
<td>-1.1179372</td>
<td>4</td>
</tr>
<tr>
<td>3</td>
<td>4</td>
<td>10</td>
<td>-1.3473558</td>
<td>6</td>
</tr>
<tr>
<td>4</td>
<td>10</td>
<td>13</td>
<td>0.4832675</td>
<td>12</td>
</tr>
<tr>
<td>5</td>
<td>10</td>
<td>13</td>
<td>0.1523950</td>
<td>12</td>
</tr>
</tbody>
</table>
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:

```r
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

```
select(df, -one_of("newcol", "y"))
```

<table>
<thead>
<tr>
<th>x</th>
<th>x2</th>
<th>z</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>7</td>
<td>6</td>
</tr>
<tr>
<td>2</td>
<td>6</td>
<td>4</td>
</tr>
<tr>
<td>3</td>
<td>10</td>
<td>7</td>
</tr>
<tr>
<td>4</td>
<td>13</td>
<td>10</td>
</tr>
<tr>
<td>5</td>
<td>13</td>
<td>5</td>
</tr>
</tbody>
</table>
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:

```r
cn = colnames(df)
df[, c("newcol", cn[cn != "newcol"])]
```

```
newcol  x  x2  y       z
 1   3  1  7 -0.2707606 6
 2   4  2  6 -1.1179372 4
 3   6  4 10 -1.3473558 7
 4  12 10 13  0.4832675 10
 5  12 10 13  0.1523950 5
```
Ordering the columns of a data.frame: dplyr

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

```
select(df, newcol, everything())
```

<table>
<thead>
<tr>
<th>newcol</th>
<th>x</th>
<th>x2</th>
<th>y</th>
<th>z</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>3</td>
<td>1</td>
<td>7</td>
<td>-0.2707606</td>
</tr>
<tr>
<td>2</td>
<td>4</td>
<td>2</td>
<td>6</td>
<td>-1.1179372</td>
</tr>
<tr>
<td>3</td>
<td>6</td>
<td>4</td>
<td>10</td>
<td>-1.3473558</td>
</tr>
<tr>
<td>4</td>
<td>12</td>
<td>10</td>
<td>13</td>
<td>0.4832675</td>
</tr>
<tr>
<td>5</td>
<td>12</td>
<td>10</td>
<td>13</td>
<td>0.1523950</td>
</tr>
</tbody>
</table>
Ordering the rows of a data.frame: base R

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

```
setosa <- iris[1:5, ]
setosa$Species

iris[order(setosa$Sepal.Length), ]
```

<table>
<thead>
<tr>
<th>x</th>
<th>x2</th>
<th>y</th>
<th>z</th>
<th>newcol</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>7</td>
<td>-0.2707606</td>
<td>6</td>
</tr>
<tr>
<td>2</td>
<td>2</td>
<td>6</td>
<td>-1.1179372</td>
<td>4</td>
</tr>
<tr>
<td>3</td>
<td>4</td>
<td>10</td>
<td>-1.3473558</td>
<td>7</td>
</tr>
<tr>
<td>4</td>
<td>10</td>
<td>13</td>
<td>0.4832675</td>
<td>10</td>
</tr>
<tr>
<td>5</td>
<td>10</td>
<td>13</td>
<td>0.1523950</td>
<td>5</td>
</tr>
</tbody>
</table>
Ordering the rows of a data.frame: base R

The decreasing argument will order it in decreasing order:

df[ order(df$x, decreasing = TRUE), ]

<p>| | | | | | | |</p>
<table>
<thead>
<tr>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>4</td>
<td>10</td>
<td>13</td>
<td>0.4832675</td>
<td>10</td>
<td>12</td>
<td></td>
</tr>
<tr>
<td>5</td>
<td>10</td>
<td>13</td>
<td>0.1523950</td>
<td>5</td>
<td>12</td>
<td></td>
</tr>
<tr>
<td>3</td>
<td>4</td>
<td>5</td>
<td>-1.3473558</td>
<td>7</td>
<td>6</td>
<td></td>
</tr>
<tr>
<td>2</td>
<td>2</td>
<td>6</td>
<td>-1.1179372</td>
<td>4</td>
<td>4</td>
<td></td>
</tr>
<tr>
<td>1</td>
<td>1</td>
<td>7</td>
<td>-0.2707606</td>
<td>6</td>
<td>3</td>
<td></td>
</tr>
</tbody>
</table>
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 `x` and decreasing on `y`):

```r
df[ order(df$x, -df$y), ]
```

<table>
<thead>
<tr>
<th>x</th>
<th>x2</th>
<th>y</th>
<th>z</th>
<th>newcol</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>7</td>
<td>-0.2707606</td>
<td>6</td>
</tr>
<tr>
<td>2</td>
<td>2</td>
<td>6</td>
<td>-1.1179372</td>
<td>4</td>
</tr>
<tr>
<td>3</td>
<td>4</td>
<td>10</td>
<td>-1.3473558</td>
<td>7</td>
</tr>
<tr>
<td>4</td>
<td>10</td>
<td>13</td>
<td>0.4832675</td>
<td>10</td>
</tr>
<tr>
<td>5</td>
<td>10</td>
<td>13</td>
<td>0.1523950</td>
<td>5</td>
</tr>
</tbody>
</table>
Ordering the rows of a `data.frame`: `dplyr`

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

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

<p>| | | | | | | |</p>
<table>
<thead>
<tr>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>7</td>
<td>-0.2707606</td>
<td>6</td>
<td>3</td>
<td></td>
</tr>
<tr>
<td>2</td>
<td>2</td>
<td>6</td>
<td>-1.1179372</td>
<td>4</td>
<td>4</td>
<td></td>
</tr>
<tr>
<td>3</td>
<td>4</td>
<td>10</td>
<td>-1.3473558</td>
<td>7</td>
<td>6</td>
<td></td>
</tr>
<tr>
<td>4</td>
<td>10</td>
<td>13</td>
<td>0.4832675</td>
<td>10</td>
<td>12</td>
<td></td>
</tr>
<tr>
<td>5</td>
<td>10</td>
<td>13</td>
<td>0.1523950</td>
<td>5</td>
<td>12</td>
<td></td>
</tr>
</tbody>
</table>
Ordering the rows of a `data.frame`: **dplyr**

Use the `desc` to arrange the rows in descending order:

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

<table>
<thead>
<tr>
<th>x</th>
<th>x2</th>
<th>y</th>
<th>z</th>
<th>newcol</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>10</td>
<td>13</td>
<td>0.4832675</td>
<td>10</td>
</tr>
<tr>
<td>2</td>
<td>10</td>
<td>13</td>
<td>0.1523950</td>
<td>5</td>
</tr>
<tr>
<td>3</td>
<td>4</td>
<td>10</td>
<td>-1.3473558</td>
<td>7</td>
</tr>
<tr>
<td>4</td>
<td>2</td>
<td>6</td>
<td>-1.1179372</td>
<td>4</td>
</tr>
<tr>
<td>5</td>
<td>1</td>
<td>7</td>
<td>-0.2707606</td>
<td>6</td>
</tr>
</tbody>
</table>
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))
```

<table>
<thead>
<tr>
<th></th>
<th>x</th>
<th>x2</th>
<th>y</th>
<th>z</th>
<th>newcol</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>7</td>
<td>-0.2707606</td>
<td>6</td>
<td>3</td>
</tr>
<tr>
<td>2</td>
<td>2</td>
<td>6</td>
<td>-1.1179372</td>
<td>4</td>
<td>4</td>
</tr>
<tr>
<td>3</td>
<td>4</td>
<td>10</td>
<td>-1.3473558</td>
<td>7</td>
<td>6</td>
</tr>
<tr>
<td>4</td>
<td>10</td>
<td>13</td>
<td>0.4832675</td>
<td>10</td>
<td>12</td>
</tr>
<tr>
<td>5</td>
<td>10</td>
<td>13</td>
<td>0.1523950</td>
<td>5</td>
<td>12</td>
</tr>
</tbody>
</table>
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:

```r
transmute(df, newcol2 = x * 3, x, y)
```

<table>
<thead>
<tr>
<th></th>
<th>newcol2</th>
<th>x</th>
<th>y</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>3</td>
<td>1</td>
<td>-0.2707606</td>
</tr>
<tr>
<td>2</td>
<td>6</td>
<td>2</td>
<td>-1.1179372</td>
</tr>
<tr>
<td>3</td>
<td>12</td>
<td>4</td>
<td>-1.3473558</td>
</tr>
<tr>
<td>4</td>
<td>30</td>
<td>10</td>
<td>0.4832675</td>
</tr>
<tr>
<td>5</td>
<td>30</td>
<td>10</td>
<td>0.1523950</td>
</tr>
</tbody>
</table>