# MODULE 22

## FILTERING DATA IN R.

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In the Module 4, you learned how to retrieve data from the class webpage, enter your own data into a CSV file, load that data into R, and how to view that data in R. In this module, we will learn how to create subsets (i.e., filter) a data.frame into smaller data.frames. For example, you may want to create a data.frame that contains just male bears from a data.frame with both male and female bears, or a data.frame that only contains sales during summer months from a data.frame that contains all sales. Less often you may wish to eliminate a particular individual from the data.frame, perhaps if it is considered to be erroneous.

### 22.1 Filtering a data.frame

It is common to create a new data.frame that contains only some of the individuals from an existing data.frame. The process of creating the newer, smaller data.frame is called filtering (or subsetting) and is accomplished with filterD(). The filterD() function requires the original data.frame as the first argument and a condition statement as the second argument. The condition statement is used to either include or exclude individuals from the original data.frame. Condition statements consist of the name of a variable in the original data.frame, a comparison operator, and a comparison value (Table 22.1). The results from filterD() should be assigned to an object, which is then the name of the new data.frame.

The following are examples of new data.frames created from bears (which was created in the previous module). The name of the new data.frame (i.e., object left of the assignment operator) can be any valid object name. As demonstrated below, the new data.frame (or its structure) should be examined after each filtering to ensure that the data.frame actually contains the items that you desire.

Table 22.1. Condition operators used in filterD() and their results. Note that *var* generically represents a variable in the original data frame and *value* is a generic value or level. Both *variable* and *value* would be replaced with specific items (see examples in main text).

Condition Operator	Individuals Returned from Original Data Frame
var == value	all individual that are equal to the given value
var ! = value	all individuals that are ${f NOT}$ equal to the given value
var > value	all individuals that are greater than the given value
var >= value	all individuals that are greater than or equal to the given value
var < value	all individuals that are less than the given value
var <= value	all individuals that are less than or equal to the given value
$condition\ ,\ condition$	all individuals that meet both conditions
$condition \mid condition$	all individuals that $\mathbf{meet}$ one or both $\mathbf{conditions}^1$

• Only individuals from *Bayfield* county.

```
> bf <- filterD(bears,loc=="Bayfield")</pre>
> bf
 length.cm weight.kg
                 110 Bayfield
      139.0
2
      138.0
                   60 Bayfield
3
      139.0
                   90 Bayfield
4
      120.5
                    60 Bayfield
5
      149.0
                    85 Bayfield
```

• Individuals from both Bayfield and Ashland counties.

```
> bfash <- filterD(bears,loc %in% c("Bayfield","Ashland"))</pre>
> bfash
 length.cm weight.kg
                           loc
     139.0
                110 Bayfield
2
     138.0
                  60 Bayfield
3
      139.0
                  90 Bayfield
4
     120.5
                   60 Bayfield
5
     149.0
                   85 Bayfield
6
      141.0
                  100 Ashland
7
      141.0
                   95
                      Ashland
```

• Individuals with a weight greater than 100 kg.

```
> gt100 <- filterD(bears, weight.kg>100)
> gt100
 length.cm weight.kg
                           loc
1
      139.0
                 110 Bayfield
2
      166.0
                  155 Douglas
3
     151.5
                  140 Douglas
4
      129.5
                  105 Douglas
5
      150.0
                  110 Douglas
```

• Individuals from *Douglas* County that weighed at least 150 kg.

```
> do150 <- filterD(bears,loc=="Douglas",weight.kg>=150)
> do150
  length.cm weight.kg   loc
1   166  155 Douglas
```

### 22.2 Selecting Individuals

In some instances, you may need to select or exclude an individual from a data frame. Positions within an object are identified within square brackets. As data frames are two-dimensional objects they are indexed by a row and a column, in that order. For example, the item in the third row and second column of bears is selected below.

```
> bears[3,2]
[1] 90
```

An entire row or column may be selected by omitting the other dimension. For example, one could select the entire second column with bears[,2], but this is also the weight.kg variable and is better selected with bears\$weight.kg. As a better example, the entire third row is selected below (note that the column designation was omitted).

```
> bears[3,]
  length.cm weight.kg loc
3 139 90 Bayfield
```

Multiple rows are selected by combining row indices together with c(). For example, the third, fifth, and eighth rows are selected below (again, the column index is omitted).

```
> bears[c(3,5,8),]
  length.cm weight.kg    loc
3    139    90 Bayfield
5    149    85 Bayfield
8    150    85 Douglas
```

Finally, rows can be excluded by preceding the row indices with a negative sign.

```
> bears[-c(3,5,8,10,12),]
   length.cm weight.kg
                             loc
1
       139.0
                    110 Bayfield
2
       138.0
                     60 Bayfield
4
       120.5
                     60 Bayfield
6
       141.0
                    100 Ashland
7
       141.0
                    95
                        Ashland
9
       166.0
                    155
                         Douglas
11
       129.5
                    105
                         Douglas
```