Filtering & Subsetting Data

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To filter or subset data is to take a dataframe or some other object, and select only part of that object based on some criteria. Most commonly we want to make a new dataframe from an existing dataframe, by filtering the rows of the dataframe based on one or more column.

As with all things data, we provide an option to perform the task using $\mathbf{core}\ \mathbf{R}$ functions, as well as in the $\mathbf{Tidyverse}$.

Filter Based on Single Value

We will use an example dataset here:

```
# remotes package is needed to get the most recent version of the package
install.packages("remotes")
# install package
remotes::install_github("lter/lterdatasampler")
library(lterdatasampler)
library(tidyverse)
```

If we want to filter a dataset based on a single value in a column, we can make use of Boolean operators.

```
# first call the dataset from the lterdatasampler package
crabs <- lterdatasampler::pie_crab
```

Let's look at the columns

```
# this will show us the first six rows
head(crabs)
```

```
## # A tibble: 6 x 9
##
                 latitude site
     date
                                  size air_temp air_temp_sd water_temp water_temp_sd
##
     <date>
                    <dbl> <chr> <dbl>
                                           <dbl>
                                                        <dbl>
                                                                    <dbl>
                                                                                   <dbl>
                       30 GTM
                                            21.8
## 1 2016-07-24
                                  12.4
                                                         6.39
                                                                    24.5
                                                                                   6.12
## 2 2016-07-24
                       30 GTM
                                  14.2
                                            21.8
                                                         6.39
                                                                    24.5
                                                                                   6.12
## 3 2016-07-24
                       30 GTM
                                  14.5
                                            21.8
                                                         6.39
                                                                    24.5
                                                                                   6.12
## 4 2016-07-24
                       30 GTM
                                  12.9
                                            21.8
                                                         6.39
                                                                    24.5
                                                                                   6.12
## 5 2016-07-24
                       30 GTM
                                  12.4
                                                                    24.5
                                                                                   6.12
                                            21.8
                                                         6.39
## 6 2016-07-24
                       30 GTM
                                  13.0
                                            21.8
                                                         6.39
                                                                    24.5
                                                                                   6.12
## # ... with 1 more variable: name <chr>
```

We see there is a column called site. We can get the unique values of that column:

```
unique(crabs$site)
```

```
## [1] "GTM" "SI" "NIB" "ZI" "RC" "VCR" "DB" "JC" "CT" "NB" "CC" "BC" ## [13] "PIE"
```

So we might want to filter our data such that we have a new dataframe, where site is only equal to "VCR".

Core R

Filtering in core R makes use of the square bracket indexing discussed elsewhere. Recall that the index goes [row,column]. In the row position, we use the which() function, which simply returns where some indices are TRUE. Since we want to keep all the columns in our dataset, we leave the column position empty, indicating that all columns should be retained. In our example that looks like this:

```
vcr_crabs <- crabs[which(crabs$site == "VCR"),]</pre>
```

We can check it worked by calling unique() on the new dataframe as above:

```
unique(vcr_crabs$site)
```

```
## [1] "VCR"
```

And we can see it worked as expected.

Tidyverse

Filtering in the Tidyverse uses mostly the dplyr package, which is EXTREMELY useful for most data management exercises. Here we use the aptly named filter() function to perform the operation.

```
vc_crabs <- crabs %>%
dplyr::filter(site == "VCR")
```

The %>% operator to pass the original data object crabs to our function, filter(), and the Boolean operator == is the only argument needed for this function.

There are many permutations of this single value option with Boolean operators. For example, using either approach, we could filter asking R to select rows where site == "VCR" | site == "BC" which would select rows where the site was either "VCR" or "BC".

In the case where we had a separate vector dictating the focal sites we were interested in, we could use that vector in combination with the %in% operator to deal with this:

Core R

```
focal_sites <- c("ZI", "RC", "VCR", "DB", "JC")
focal_crabs <- crabs[which(crabs$site %in% focal_sites),]</pre>
```

Tidyverse

```
focal_sites <- c("ZI", "RC", "VCR", "DB", "JC")

focal_crabs <- crabs %>%
   dplyr::filter(site %in% focal_sites)
```

Filter Based on Multiple Values

This filtering can be extended to use multiple columns relatively simply. There is a variable in this dataset called size.

```
hist(crabs$size)
```

Histogram of crabs\$size



want to filter crabs only in our focal sites, but also those above some size (e.g. 15).

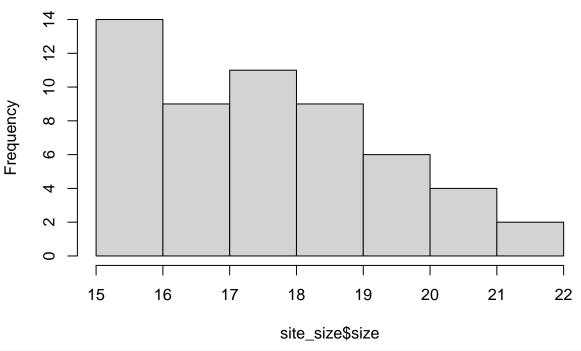
Core R

${\bf Tidy verse}$

```
site_size <- crabs %>%
dplyr::filter(site %in% focal_sites & size > 15)
```

Here we just use another logical operator, & to link our two conditions together. We can double check: hist(site_size\$size)

Histogram of site_size\$size



unique(site_size\$site)

```
## [1] "ZI" "RC" "VCR" "DB" "JC"
```

So we can see it worked well.

Filtering Out NA's

For this example, we will use a different dataset.

```
ice <- lterdatasampler::ntl_icecover
head(ice)</pre>
```

```
## # A tibble: 6 x 5
     lakeid
                                         ice_duration year
                  ice_on
                              ice_off
##
     <fct>
                  <date>
                              <date>
                                                 <dbl> <dbl>
## 1 Lake Mendota NA
                              1853-04-05
                                                   NA
                                                        1852
## 2 Lake Mendota 1853-12-27 NA
                                                   NA
                                                        1853
## 3 Lake Mendota 1855-12-18 1856-04-14
                                                   118
                                                        1855
## 4 Lake Mendota 1856-12-06 1857-05-06
                                                        1856
                                                   151
## 5 Lake Mendota 1857-11-25 1858-03-26
                                                   121
                                                        1857
## 6 Lake Mendota 1858-12-08 1859-03-14
                                                   96
                                                        1858
```

We can see already that there will be some values of $ice_duration$ that will have values of NA. To filter those out of the dataset, we can use the is.na() function:

Core R

```
trimmed_ice <- ice[which(!is.na(ice$ice_duration)),]</pre>
```

Again, we want to keep all the columns present.

Tidyverse

```
trimmed_ice <- ice %>%
  dplyr::filter(!is.na(ice_duration))
```

Here we use the combination of the ! and is.na() to first find the positions where the value is equal to NA, and then to remove them with the !.

Any conceivable set of criteria can be strung together (carefully!) to filter a dataset with as many conditions as necessary. It is always a good idea however when you are just beginning to use these functions, to work piece by piece, adding conditions one at a time and testing each one to make sure it's doing what you want. Then you can string multiple conditions together without worrying that an error is not being caught.