Intro to piping and Data Manipulation

JEPA

02/06/2017

Contents

Dplyr and Tidyr	า
Dplyr and ridyr Dplyr	2
Arrange	$\frac{2}{2}$
Basic structure:	$\frac{2}{2}$
Filter	3
Basic structure:	3
Group_by* (plus summarise)	4
Basic structure:	4
Mutate	5
Basic structure:	5
select	6
Basic structure:	6
slice	7
Basic Structure	7
Isining Data with dalam	8
Joining Data with dplyr The "bind" family	8
bind cols	8
bind rows	8
The "join" family	9
anti join	9
semi join	9
inner join	9
left join	10
right join	10
intersect	10
union	10
setdiff	10
Scium	10
The Piping opperator %>%	10

Libraries and Data

```
#install.packages('dplyr')
library(dplyr)

#install.packages('tidyr')
library(tidyr)

#install.packages('ggplot2')
library(ggplot2)
```

```
Alaska <- read.csv("./Data/Alaska.csv") #Sea around Us data for Alaska
USA <- read.csv("./Data/USAP.csv") #Sea around Us data for USA
```

Dplyr and Tidyr

Despite beign separate, these two packadges work together as one. Theyr main function is to manipulate data frames and keep things "tydi". In some cases you can also make basic data creation. Both packadges follow the same syntax and can use the pipe opperator, I normally don't even know which function is from what packadge so I oftenly just call both.

Plus: Most functions are self explanatory like select or filter!

Abalones Other demersal invertebrates

Dplyr

Arrange

1

The arrangefunction allows you to, literaly, arrenge your data by any value of a column

Basic structure:

```
New Table <- arrange(Data, column to arange by)
Note: If you eant to do from Top <- Bottom you can use desc() within the function
Note: when doing multiple variables the order is important since it will start with the first one
#You can arrange by characters (A -> Z)
Arrange_Example <- arrange(Alaska,common_name)</pre>
head(Arrange_Example[5:7], 3)
     year scientific_name common_name
## 1 1964
                  Haliotis
                              Abalones
## 2 1964
                  Haliotis
                              Abalones
## 3 1966
                 Haliotis
                              Abalones
#You can arrange by characters (A <- Z) using desc()
Arrange_Example2 <- arrange(Alaska,desc(common_name))</pre>
head(Arrange_Example2[5:7], 3)
     year
            scientific_name
                                      common_name
## 1 1984 Sebastes flavidus Yellowtail rockfish
## 2 1985 Sebastes flavidus Yellowtail rockfish
## 3 1987 Sebastes flavidus Yellowtail rockfish
# you can do multiple characters:
Arrange_Example3 <- arrange(Alaska,common_name,functional_group, desc(commercial_group))
head(Arrange Example3[7:9],3)
                              functional_group commercial_group
##
     common name
```

Molluscs

```
## 2
        Abalones Other demersal invertebrates
                                                        Molluscs
## 3
        Abalones Other demersal invertebrates
                                                        Molluscs
# And naturally, you can also arrange by numeric factors
Arrange_Example4 <- arrange(Alaska, uncertainty_score, desc(tonnes))
head(Arrange_Example4[4:6],3)
##
     uncertainty_score year
                                      scientific_name
## 1
                     1 2010 Oncorhynchus tshawytscha
## 2
                     1 1989 Oncorhynchus tshawytscha
## 3
                     1 1988 Oncorhynchus tshawytscha
Filter
The filterfunction allows you to, literaly, filter your data by any category or numer.
Basic structure:
New Table <- filter(Data, column_to_filter_by == "category")
#You can filter by character
Filter_Example <- filter(Alaska,common_name =="Clams")</pre>
head(Filter_Example[1:5], 5)
##
                   area_name area_type
                                                           data_layer
## 1 USA (Alaska, Subarctic)
                                    eez Reconstructed domestic catch
## 2 USA (Alaska, Subarctic)
                                    eez Reconstructed domestic catch
## 3 USA (Alaska, Subarctic)
                                    eez Reconstructed domestic catch
                                    eez Reconstructed domestic catch
## 4 USA (Alaska, Subarctic)
## 5 USA (Alaska, Subarctic)
                                    eez Reconstructed domestic catch
##
    uncertainty_score year
## 1
                     1 1950
## 2
                     1 1951
## 3
                     1 1952
## 4
                     1 1953
## 5
                     1 1954
#You can filter by numeric input too
Filter_Example2 <- filter(Alaska,</pre>
                          year == 2009)
head(Filter_Example2[1:5], 5)
##
                                                           data_layer
                   area_name area_type
## 1 USA (Alaska, Subarctic)
                                    eez Reconstructed domestic catch
## 2 USA (Alaska, Subarctic)
                                    eez Reconstructed domestic catch
## 3 USA (Alaska, Subarctic)
                                    eez Reconstructed domestic catch
## 4 USA (Alaska, Subarctic)
                                    eez Reconstructed domestic catch
## 5 USA (Alaska, Subarctic)
                                    eez Reconstructed domestic catch
    uncertainty_score year
##
## 1
                     4 2009
                     4 2009
## 2
```

3

4

2 2009

4 2009

```
## 5
                     4 2009
# Note: you can do =>, <= or !=
# you can do multiple characters:
Selection <- c("Clams", "Octopuses")</pre>
Filter_Example3 <- filter(Alaska,common_name %in% Selection)</pre>
head(Filter_Example3[4:8], 5)
##
     uncertainty_score year scientific_name common_name
## 1
                     1 1950
                                    Bivalvia
                                                   Clams
## 2
                     1 1950
                                Octopodidae
                                               Octopuses
## 3
                     1 1951
                                    Bivalvia
                                                   Clams
## 4
                                Octopodidae
                                               Octopuses
                     1 1951
## 5
                     1 1952
                                    Bivalvia
                                                   Clams
##
                 functional_group
## 1 Other demersal invertebrates
## 2
                      Cephalopods
## 3 Other demersal invertebrates
## 4
                      Cephalopods
## 5 Other demersal invertebrates
# NOTE: remember that in R there are multiple ways to get to the same result!
#Wait! What if I want to filter by multiple columns!?
Filter_Example4 <- filter(Alaska,common_name == "Clams" &
                            reporting_status =="Unreported")
#You can also filter by NA
Filter_NA_Example1 <- filter(Alaska,is.na(uncertainty_score)) #Extract only NA's
head(Filter_NA_Example1[1:4],3)
                   area_name area_type
                                                    data layer
## 1 USA (Alaska, Subarctic)
                              eez Inferred foreign catch
## 2 USA (Alaska, Subarctic)
                                   eez Inferred foreign catch
## 3 USA (Alaska, Subarctic)
                                   eez Inferred foreign catch
##
    uncertainty_score
## 1
                    NA
## 2
                    NA
## 3
                    NA
Filter NA Example2 <- filter(Alaska,!is.na(uncertainty score)) #Clear NA's
```

Group_by* (plus summarise)

The <code>group_byfunction</code> allows you to group your data by common variables for future (inmidiate) calculations. This function needs the "pipe opperator"

Basic structure:

```
New Table <- Data %>% group by(column 1,column 2...) %>% second function()
#Simple group_by
Group_by_Example <- Alaska %>%
  group_by(common_name) %>%
  summarise(n()) #tells you how many rows of each common_name you have
head(Group_by_Example, 3)
## # A tibble: 3 x 2
##
        common_name
                      n()
##
             <fctr> <int>
## 1
           Abalones
                       52
## 2 Alaska plaice
## 3 Alaska pollock
                      290
#Multiple
Group_by_Example2 <- Alaska %>%
  group_by(common_name,uncertainty_score) %>%
  summarise(n()) %>% #tells you how many rows of each common name you have
  arrange(uncertainty_score)
head(Group_by_Example, 3)
## # A tibble: 3 x 2
##
        common_name
                      n()
##
             <fctr> <int>
## 1
           Abalones
                       52
## 2 Alaska plaice
                        9
## 3 Alaska pollock
                      290
```

Mutate

The mutatefunction allows you to create a new column in the dataset. The new columb can have characters or numbers.

Basic structure:

```
New Table <- mutate(Data, Name New Column = action)
#Functions
Mutate_Example1 <- mutate(Alaska, Log = log(tonnes))</pre>
head(Mutate_Example1[13:16], 3)
##
     reporting_status
                          tonnes landed_value
                                                    Log
## 1
           Unreported
                         13.8030
                                      20235.2 2.624886
## 2
             Reported 1483.9740
                                    2175505.9 7.302479
                                     571724.0 5.966119
## 3
           Unreported 389.9891
#In data calculations (per row)
Mutate_Example2 <- mutate(Alaska, Price_plus_Ton = (landed_value+tonnes))</pre>
head(Mutate_Example2[13:16], 3)
```

```
## 1
           Unreported
                        13.8030
                                       20235.2
                                                         20249
## 2
                                    2175505.9
                                                      2176990
             Reported 1483.9740
## 3
           Unreported 389.9891
                                     571724.0
                                                        572114
#Or characters...
Mutate_Example3 <- mutate(Alaska, Country = "USA")</pre>
head(Mutate_Example3[13:16], 3)
##
     reporting_status
                          tonnes landed_value Country
                                       20235.2
## 1
           Unreported
                         13.8030
                                                   USA
## 2
             Reported 1483.9740
                                     2175505.9
                                                   USA
## 3
           Unreported 389.9891
                                     571724.0
                                                   USA
Mutate_Example4 <- mutate(Mutate_Example3, Country = paste("In", year, Country, "harvested", round(tonnes, 2</pre>
paste(Mutate_Example4[1,16])
## [1] "In 1950 USA harvested 13.8 tonnes of Marine fishes nei"
paste(Mutate_Example4[5387,16])
## [1] "In 1979 USA harvested 18.7 tonnes of Squids"
select
The selectfunction is one of those "of-course it does that" function cus it allows you to, wait for it...
SELECT any column you want.
Basic structure:
New_Table <- select(Data,number or name of colum)
#Select by column number
Select_Example1 <- select(Alaska, 6)</pre>
head(Select_Example1,3)
##
                  scientific_name
## 1 Marine fishes not identified
## 2 Marine fishes not identified
## 3 Marine fishes not identified
#Select by multiple column numbers
Select_Example2 <- select(Alaska, 4,5,6,7)</pre>
head(Select_Example2, 3)
     uncertainty_score year
##
                                           scientific_name
                                                                  common_name
## 1
                      1 1950 Marine fishes not identified Marine fishes nei
## 2
                      3 1950 Marine fishes not identified Marine fishes nei
                      3 1950 Marine fishes not identified Marine fishes nei
# You can also do (4:7) and even (4:6,15)
```

Select_Example3 <- select(Alaska, area_name, year, scientific_name, tonnes)</pre>

#Select by name

```
head(Select_Example3, 3)
                   area_name year
                                                scientific_name
                                                                   tonnes
## 1 USA (Alaska, Subarctic) 1950 Marine fishes not identified
                                                                   13.8030
## 2 USA (Alaska, Subarctic) 1950 Marine fishes not identified 1483.9740
## 3 USA (Alaska, Subarctic) 1950 Marine fishes not identified
# You can substract columns from a dataframe
Select_Example4 <- select(Select_Example3, -area_name, year)</pre>
head(Select_Example4, 3)
##
                       scientific_name
    year
                                           tonnes
## 1 1950 Marine fishes not identified
                                          13.8030
## 2 1950 Marine fishes not identified 1483.9740
## 3 1950 Marine fishes not identified 389.9891
#Note, you can also substract using -1
#And you can also re-order your columns!
Select_Example5 <- select(Select_Example3, scientific_name, year, tonnes, area_name)</pre>
head(Select_Example5, 3)
                  scientific_name year
                                           tonnes
                                                                 area_name
## 1 Marine fishes not identified 1950
                                          13.8030 USA (Alaska, Subarctic)
## 2 Marine fishes not identified 1950 1483.9740 USA (Alaska, Subarctic)
## 3 Marine fishes not identified 1950 389.9891 USA (Alaska, Subarctic)
```

slice

The slicefunction works like the selectfunction but for rows. So, if you want to extract an specific row, a set of rows, or a range between values, use slice!

Basic Structure

```
New_Data <- slice(Old_Data, number)
```

```
#Select by row number
Slice_Example1 <- slice(Alaska, 3948)
Slice_Example1</pre>
```

```
area_name area_type
                                                         data_layer
## 1 USA (Alaska, Subarctic)
                                   eez Reconstructed domestic catch
##
    uncertainty_score year
                                    scientific_name
                                                         common_name
## 1
                     3 1973 Clupea pallasii pallasii Pacific herring
##
                 functional_group commercial_group fishing_entity
## 1 Medium pelagics (30 - 89 cm)
                                     Herring-likes
    fishing sector catch type reporting status tonnes landed value
                                      Reported 15792.9
## 1
        Industrial Landings
                                                            23152391
```

```
#Select by multiple rows
Slice_Example2 <- slice(Alaska, 1000:3948)
head(Slice_Example2, 3)
##
                   area_name area_type
                                                          data_layer
## 1 USA (Alaska, Subarctic)
                                   eez Reconstructed domestic catch
## 2 USA (Alaska, Subarctic)
                                   eez Reconstructed domestic catch
## 3 USA (Alaska, Subarctic)
                                    eez Reconstructed domestic catch
     uncertainty_score year
                                      scientific_name
                                                          common_name
## 1
                     3 1957 Hippoglossus stenolepis Pacific halibut
## 2
                     1 1957 Hippoglossus stenolepis Pacific halibut
## 3
                     3 1957 Clupea pallasii pallasii Pacific herring
##
                 functional_group commercial_group fishing_entity
                                        Flatfishes
## 1
       Large flatfishes (>=90 cm)
                                                               USA
                                                               USA
       Large flatfishes (>=90 cm)
                                        Flatfishes
## 3 Medium pelagics (30 - 89 cm)
                                     Herring-likes
                                                               USA
     fishing_sector catch_type reporting_status
                                                      tonnes landed_value
## 1
          Artisanal
                      Landings
                                       Reported 12564.60000 18419703.60
## 2
       Recreational
                      Landings
                                      Unreported
                                                    11.14694
                                                                 16341.42
## 3
         Industrial
                      Landings
                                       Reported 53656.10001 78659842.61
Joining Data with dplyr
The "bind" family
bind_cols
#Lets just asume that we have two different data sets
Data1 <- select(Alaska, 1)
Data2 <- select(Alaska, 2)
#Now we bind the columns together
Bind_Cols_1 <- bind_cols(Data1,Data2)</pre>
head(Bind_Cols_1, 3)
##
                   area_name area_type
## 1 USA (Alaska, Subarctic)
                                    eez
## 2 USA (Alaska, Subarctic)
                                    eez
## 3 USA (Alaska, Subarctic)
                                    eez
bind_rows
#Lets just asume that we have two different data sets
Data1 <- slice(Alaska, 1:3)
Data2 <- slice(Alaska, 10800:10802)</pre>
#Now we bind the columns together
Bind_Row_1 <- bind_cols(Data1,Data2)</pre>
head(Bind_Row_1, 6)
```

```
area_name area_type
##
                                                          data_layer
                                   eez Reconstructed domestic catch
## 1 USA (Alaska, Subarctic)
## 2 USA (Alaska, Subarctic)
                                   eez Reconstructed domestic catch
## 3 USA (Alaska, Subarctic)
                                    eez Reconstructed domestic catch
##
     uncertainty score year
                                          scientific name
                                                                 common name
## 1
                     1 1950 Marine fishes not identified Marine fishes nei
## 2
                     3 1950 Marine fishes not identified Marine fishes nei
## 3
                     3 1950 Marine fishes not identified Marine fishes nei
##
                  functional_group
                                          commercial_group fishing_entity
## 1 Medium demersals (30 - 89 cm) Other fishes & inverts
                                                                       USA
## 2 Medium demersals (30 - 89 cm) Other fishes & inverts
                                                                       USA
## 3 Medium demersals (30 - 89 cm) Other fishes & inverts
                                                                       USA
     fishing_sector catch_type reporting_status
                                                    tonnes landed value
## 1
        Subsistence
                      Landings
                                      Unreported
                                                   13.8030
                                                                 20235.2
## 2
          Artisanal
                                        Reported 1483.9740
                                                               2175505.9
                      Landings
## 3
          Artisanal
                      Landings
                                      Unreported 389.9891
                                                                571724.0
##
                   area_name area_type
                                                          data_layer
## 1 USA (Alaska, Subarctic)
                                   eez Reconstructed domestic catch
## 2 USA (Alaska, Subarctic)
                                   eez Reconstructed domestic catch
## 3 USA (Alaska, Subarctic)
                                    eez Reconstructed domestic catch
##
     uncertainty_score year
                                scientific_name common_name
## 1
                     4 2009 Anoplopoma fimbria
                                                  Sablefish
## 2
                     2 2009 Anoplopoma fimbria
                                                  Sablefish
## 3
                     4 2009 Anoplopoma fimbria
                                                  Sablefish
##
                   functional_group commercial_group fishing_entity
## 1 Large bathydemersals (>=90 cm)
                                       Scorpionfishes
                                                                  USA
## 2 Large bathydemersals (>=90 cm)
                                       Scorpionfishes
                                                                  USA
## 3 Large bathydemersals (>=90 cm)
                                       Scorpionfishes
                                                                  USA
     fishing_sector catch_type reporting_status
                                                       tonnes landed_value
## 1
         Industrial
                      Landings
                                        Reported
                                                  1074.516856
                                                               1575241.711
## 2
        Subsistence
                      Landings
                                      Unreported
                                                     5.002588
                                                                   7333.794
## 3
          Artisanal
                      Landings
                                        Reported 11175.083144 16382671.889
The "join" family
anti_join
#Lets asume we want to know how many species are fished in Alaska and not in the continental US
Similar_Species <- anti_join(Alaska, USA, by="scientific_name")</pre>
#You can also do it by more than one variable
Similar Species2 <- anti join(Alaska, USA, by=c("scientific name", "reporting status"))
semi_join
#Now we want to know how many species are fished in BOTH Alaska and the continental US
Diff_Species <- semi_join(Alaska, USA, by="scientific_name")</pre>
```

 $inner_join$

#Not just like anti_join, you can do it for more than one variable

left_join
right_join
intersect
union
setdiff

The Piping opperator %>%

Many R packadges like dplyr, tidyr and leaflet, allows you to use the pipe (%>%) operator to chain functions together. Chaining code allows you to streamline your workflow and make it easier to read.

When using the %>% operator, first specify the data frame that all following functions will use. For the rest of the chain the data frame argument can be omitted from the remaining functions.

NOTE: for Mac users the pipe simbol "%>%" shortcut is command + shit + m