

Intro to piping and Data Manipulation

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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 being separate, these two packages work together as one. Their main function is to manipulate data frames and keep things “tidy”. In some cases you can also make basic data creation. Both packages follow the same syntax and can use the pipe operator, I normally don’t even know which function is from what package so I often just call both.

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

Dplyr

Arrange

The `arrange` function allows you to, literally, arrange your data by any value of a column

Basic structure:

```
New_Table <- arrange(Data, column_to_arrange_by)
```

Note: If you want 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)

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))

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))

knitr::kable(head(Arrange_Example3[7:9],3))
```

common_name	functional_group	commercial_group
Abalones	Other demersal invertebrates	Molluscs
Abalones	Other demersal invertebrates	Molluscs
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 `filter` function allows you to, literally, filter your data by any category or number.

Basic structure:

```
New_Table <- filter(Data, column_to_filter_by == "category")
#You can filter by character
Filter_Example <- filter(Alaska,common_name == "Clams")

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
## 4 USA (Alaska, Subarctic)    eez Reconstructed domestic catch
## 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,
                          year == 2009)
head(Filter_Example2[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
## 4 USA (Alaska, Subarctic)    eez Reconstructed domestic catch
## 5 USA (Alaska, Subarctic)    eez Reconstructed domestic catch
## uncertainty_score year
## 1          4 2009
## 2          4 2009
## 3          2 2009
## 4          4 2009
## 5          4 2009
```

```
# Note: you can do ==, <= or !=
```

```
# you can do multiple characters:
```

```
Selection <- c("Clams","Octopuses")
Filter_Example3 <- filter(Alaska,common_name %in% Selection)
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          1 1951    Octopodidae    Octopuses
## 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 `group_by` function allows you to group your data by common variables for future (immediate) calculations. This function needs the “pipe operator”

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    9
## 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 `mutate` function allows you to create a new column in the dataset. The new column can have characters or numbers.

Basic structure:

```
New_Table <- mutate(Data, Name_New_Column = action)
```

#Functions

```
Mutate_Example1 <- mutate(Alaska, Log = log(tonnes))
```

```
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
## 3      Unreported  389.9891      571724.0 5.966119
```

#In data calculations (per row)

```
Mutate_Example2 <- mutate(Alaska, Price_plus_Ton = (landed_value+tonnes))
```

```
head(Mutate_Example2[13:16], 3)
```

```
##   reporting_status   tonnes landed_value Price_plus_Ton
## 1      Unreported   13.8030      20235.2      20249
## 2        Reported 1483.9740     2175505.9     2176990
## 3      Unreported  389.9891      571724.0      572114
```

#Or characters...

```
Mutate_Example3 <- mutate(Alaska, Country = "USA")
```

```
head(Mutate_Example3[13:16], 3)
```

```
##   reporting_status   tonnes landed_value Country
## 1      Unreported   13.8030      20235.2     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),
```

```
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 `select` function 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)

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)

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                   3 1950 Marine fishes not identified Marine fishes nei

# You can also do (4:7) and even (4:6,15)

#Select by name
Select_Example3 <- select(Alaska, area_name,year,scientific_name,tonnes)

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 389.9891

# You can subtract columns from a dataframe

Select_Example4 <- select(Select_Example3, -area_name,year)

head(Select_Example4, 3)

##   year           scientific_name      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 subtract using -1

#And you can also re-order your columns!

Select_Example5 <- select(Select_Example3, scientific_name,year,tonnes,area_name)

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 `slice` function works like the `select` function 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
```

```
##           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           USA
##   fishing_sector catch_type reporting_status tonnes landed_value
## 1   Industrial   Landings      Reported 15792.9    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
## 1   Large flatfishes (>=90 cm)   Flatfishes           USA
## 2   Large flatfishes (>=90 cm)   Flatfishes           USA
## 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)

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)
```

```
#Now we bind the columns together
Bind_Row_1 <- bind_rows(Data1,Data2)

head(Bind_Row_1, 6)
```

```
##           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              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   Landings      Reported    1483.9740    2175505.9
## 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")

#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")

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

inner_join

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`