Data Cleaning and Manipulation

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```
library(dplyr)
library(tidyr)
# To call stats filter -- stats::filter()
```

Reformat the catch data

- Remove "all" column
- Remove "notes" column
- Create a "species" column
 - Move from wide to long
- General QA

Mike Byerly. Alaska commercial salmon catches by management region (1886- 1997). Gulf of Alaska Data Portal. df35b.304.2.

```
catch_original = read.csv("https://knb.ecoinformatics.org/knb/d1/mn/v2/object/df35b.302.1", stringsAsFa
head(catch_original)
```

```
Region Year Chinook Sockeye Coho Pink Chum All notesRegCode
##
## 1
        SSE 1886
                        0
                                 5
                                      0
## 2
        SSE 1887
                        0
                               155
                                      0
                                           0
                                                 0 155
        SSE 1888
## 3
                        0
                               224
                                     16
                                           0
                                                 0 240
## 4
        SSE 1889
                        0
                               182
                                     11
                                           92
                                                 0 285
        SSE 1890
                        0
                               251
                                     42
                                                 0 292
## 5
                                           0
## 6
        SSE 1891
                        0
                               274
                                     24
                                                 0 298
```

Remove the "all" and "notesRegCode" columns using "select"

```
##
     Region Year species catch
## 1
        SSE 1886 Chinook
                              0
## 2
        SSE 1887 Chinook
        SSE 1888 Chinook
                              0
## 3
## 4
        SSE 1889 Chinook
                             0
## 5
        SSE 1890 Chinook
                             0
## 6
        SSE 1891 Chinook
```

```
catch_wide = catch_long %>% spread(key = species, value = catch)
head(catch_wide)
     Region Year Chinook Chum Coho Pink Sockeye
        ALU 1911
## 1
                         0
                              0
## 2
        ALU 1912
                         0
                              0
                                   0
                                         0
                                                 0
## 3
        ALU 1913
                        0
                              0
                                   0
                                         0
                                                 0
                        0
                              0
                                                 0
        ALU 1914
                                   0
                                         0
## 5
                        0
                              0
                                                 0
        ALU 1915
                                   0
                                         0
## 6
        ALU 1916
                                      180
                                                76
```

Clean up our data

- Rename catch to catch_thousands
- Change "catch" column to numeric
- Create a new "catch" column in units num. of fish

```
##
    Region Year species catch
## 1
        SSE 1886 Chinook
## 2
        SSE 1887 Chinook
                              0
## 3
        SSE 1888 Chinook
                              0
## 4
        SSE 1889 Chinook
                              0
## 5
        SSE 1890 Chinook
                              0
## 6
        SSE 1891 Chinook
                              0
```

Split - Apply - Combine

• Calculate mean catch by species

```
species_mean = catch_clean %>% group_by(species, Region) %>% summarise(catch_mean = mean(catch), num_ob
head(species_mean)
```

```
## # A tibble: 6 x 4
## # Groups:
               species [1]
##
     species Region catch_mean num_obs
##
     <chr>
             <chr>>
                          <dbl>
                                   <int>
## 1 Chinook ALU
                           23.0
                                      87
## 2 Chinook BER
                           19.6
                                     102
## 3 Chinook BRB
                        76211.
                                     114
## 4 Chinook CHG
                         1536.
                                     110
## 5 Chinook CKI
                        43876.
                                     105
## 6 Chinook COP
                        19798.
                                      94
```

Join the region definitions

```
regions_original = read.csv("https://knb.ecoinformatics.org/knb/d1/mn/v2/object/df35b.303.1")
```

Miscellaneous functions