

# Data Exploration

```
# libraries used
library(ggplot2)
library(dplyr)
```

```
##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##   filter, lag

## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union
```

```
library(ggfortify)
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.3.2 --

## v tibble  3.1.8      v purrr  0.3.5
## v tidyr   1.2.1      v stringr 1.4.1
## v readr   2.1.3      v forcats 0.5.2
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()    masks stats::lag()
```

```
fish.data <- read.csv("TrawlCatch_SpringPreyfishBottomTrawl.csv")
fish.data.raw <- fish.data %>%
  mutate(year=as.factor(year)) #didn't want year to be a continuous variable, I wanted them as discrete
fish.data <- fish.data.raw %>%
  filter(!is.na(fishingTemperature_C), !is.na(latitude), !is.na(longitude), commonName!= "No fish caught")
#I am also removing unidentified or misc. fishes
```

```
fish.data %>%
  group_by(commonName) %>%
  tally() %>%
  arrange(desc(n)) #this helped us see how many observations we had per species
```

```
## # A tibble: 63 x 2
##   commonName      n
##   <chr>          <int>
```

```
## 1 Alewife 6659
## 2 Rainbow smelt 4646
## 3 Slimy sculpin 2328
## 4 Lake trout 1635
## 5 Round goby 1135
## 6 Dreissena spp. 1102
## 7 Johnny darter 1048
## 8 Trout-perch 714
## 9 Deepwater sculpin 618
## 10 Threespine stickleback 482
## # ... with 53 more rows
```

```
fish.list <- as.data.frame(unique(fish.data$commonName)) #this just made a data frame of the list so we
fish.data.exonats <- fish.data %>% #Here we made a new column that marks each fish species as exotic or
```

```
  mutate(inv.status = case_when(
    endsWith(commonName, "Alewife") ~ "exotic",
    endsWith(commonName, "Sea lamprey") ~ "exotic",
    endsWith(commonName, "Chinook salmon") ~ "exotic",
    endsWith(commonName, "Rainbow trout (Steelhead)") ~ "exotic",
    endsWith(commonName, "Carp") ~ "exotic",
    endsWith(commonName, "Brown trout") ~ "exotic",
    endsWith(commonName, "Rainbow smelt") ~ "exotic",
    endsWith(commonName, "Coho salmon") ~ "exotic",
    endsWith(commonName, "White perch") ~ "exotic",
    endsWith(commonName, "Blueback herring") ~ "exotic",
    endsWith(commonName, "Chain pickerel") ~ "exotic",
    endsWith(commonName, "Round goby") ~ "exotic",
    endsWith(commonName, "Tubenose goby") ~ "exotic",
    endsWith(commonName, "Threespine stickleback") ~ "native",
    endsWith(commonName, "Emerald shiner") ~ "native",
    endsWith(commonName, "Lake whitefish") ~ "native",
    endsWith(commonName, "Deepwater sculpin") ~ "native",
    endsWith(commonName, "Lake trout") ~ "native",
    endsWith(commonName, "Burbot") ~ "native",
    endsWith(commonName, "Slimy sculpin") ~ "native",
    endsWith(commonName, "Emerald shiner") ~ "native",
    endsWith(commonName, "Cisco (lake herring)") ~ "native",
    endsWith(commonName, "Whitefishes") ~ "native",
    endsWith(commonName, "Johnny darter") ~ "native",
    endsWith(commonName, "Trout-perch") ~ "native",
    endsWith(commonName, "Yellow perch") ~ "native",
    endsWith(commonName, "Spottail shiner") ~ "native"
  ))
fish.data.exonats %>%
  filter(is.na(inv.status)) %>%
  group_by(commonName) %>%
  tally() %>%
  arrange(desc(n))
```

```
## # A tibble: 37 x 2
##   commonName      n
##   <chr>          <int>
## 1 Dreissena spp. 1102
## 2 White bass    65
```

```
## 3 White sucker          65
## 4 Rockbass              64
## 5 American eel         60
## 6 Walleye              53
## 7 Freshwater drum      49
## 8 Vegetation/plant material 43
## 9 Brown bullhead       31
## 10 Pumpkinseed         29
## # ... with 27 more rows
```

```
#Checking to see which ones I hadn't researched yet to make sure I did not miss any important ones.
#Dreissena are mussels and we are only focused on fishes so we will be cutting those out anyway
#We ignore everything below 200 observations on this list because they do not have enough observations
fish.data.exonat %>% #now that we have labeled each species, we can display our native species of interest
  filter(inv.status=="native") %>%
  group_by(commonName) %>%
  tally() %>%
  arrange(desc(n))
```

```
## # A tibble: 13 x 2
##   commonName      n
##   <chr>          <int>
## 1 Slimy sculpin   2328
## 2 Lake trout     1635
## 3 Johnny darter  1048
## 4 Trout-perch    714
## 5 Deepwater sculpin 618
## 6 Threespine stickleback 482
## 7 Yellow perch    396
## 8 Spottail shiner 271
## 9 Lake whitefish  131
## 10 Emerald shiner 110
## 11 Cisco (lake herring) 65
## 12 Burbot        19
## 13 Whitefishes    1
```

```
#based on this, we can choose only species with more than 300 observations. In this case that means Yellow perch
```

```
fish.data.clean <- fish.data.exonat %>% #this is now the data we are interested in, including only the species we want
  filter(commonName=="Yellow perch" | commonName=="Threespine stickleback" | commonName=="Deepwater sculpin")
head(fish.data.clean)
```

```
##   opId year      vesselName serial  opDate
## 1 30247 1992 Kaho              2 19920421
## 2 30248 1992 Kaho              3 19920421
## 3 48124 1984 Kaho             35 19840420
## 4 48124 1984 Kaho             35 19840420
## 5 48124 1984 Kaho             35 19840420
## 6 48125 1984 Kaho             36 19840420
##   latitude longitude fishingTemperature_C fishingDepth_m towTime_min
## 1 43.38000 -77.51833          2.9          150          10
```

```
## 2 43.38667 -77.55833          2.6          130          10
## 3 43.37167 -78.75000          2.7           55          10
## 4 43.37167 -78.75000          2.7           55          10
## 5 43.37167 -78.75000          2.7           55          10
## 6 43.37833 -78.75000          2.9           65          10
##   speed_mpsec wingSpreadModeled_m extraBottomContactTime_sec
## 1    1.251712          9.081622          9.191982
## 2    1.251712          9.081138          9.185680
## 3    1.251712          8.979826          8.960727
## 4    1.251712          8.979826          8.960727
## 5    1.251712          8.979826          8.960727
## 6    1.251712          9.029780          9.039374
##   areaSampledDoors_m2      lifeStageName  commonName  n weight_g
## 1          11803.029 Life Stage Not Recorded Slimy sculpin 33      309
## 2          10730.085 Life Stage Not Recorded Slimy sculpin 193     1919
## 3           7756.949 Life Stage Not Recorded  Lake trout  20     7768
## 4           7756.949 Life Stage Not Recorded Johnny darter 256     380
## 5           7756.949 Life Stage Not Recorded Slimy sculpin 280    1337
## 6           8093.266 Life Stage Not Recorded  Lake trout   2     390
##   inv.status
## 1      native
## 2      native
## 3      native
## 4      native
## 5      native
## 6      native
```

```
unique(fish.data$year) #we have data from 1997 to 2022
```

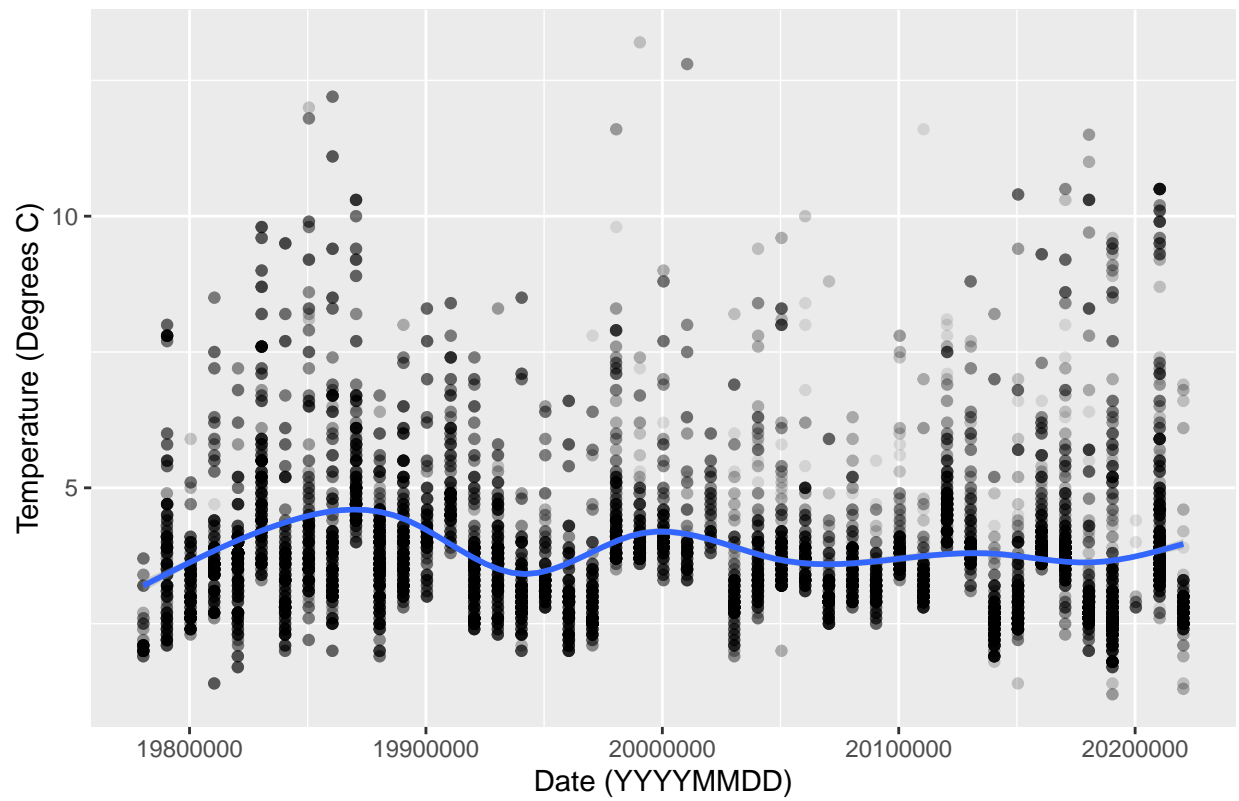
```
## [1] 1992 1984 1985 2004 1983 1979 1978 1996 1995 1998 1999 1994 1986 1993 2000
## [16] 1988 1987 1997 1982 1981 1980 1990 1989 2005 2006 2003 2002 2001 1991 2009
## [31] 2011 2007 2010 2013 2008 2012 2014 2017 2018 2015 2016 2022 2019 2021 2020
## 45 Levels: 1978 1979 1980 1981 1982 1983 1984 1985 1986 1987 1988 1989 ... 2022
```

```
#the dates should be converted into a more readable format. I just don't know how to do that so I need
```

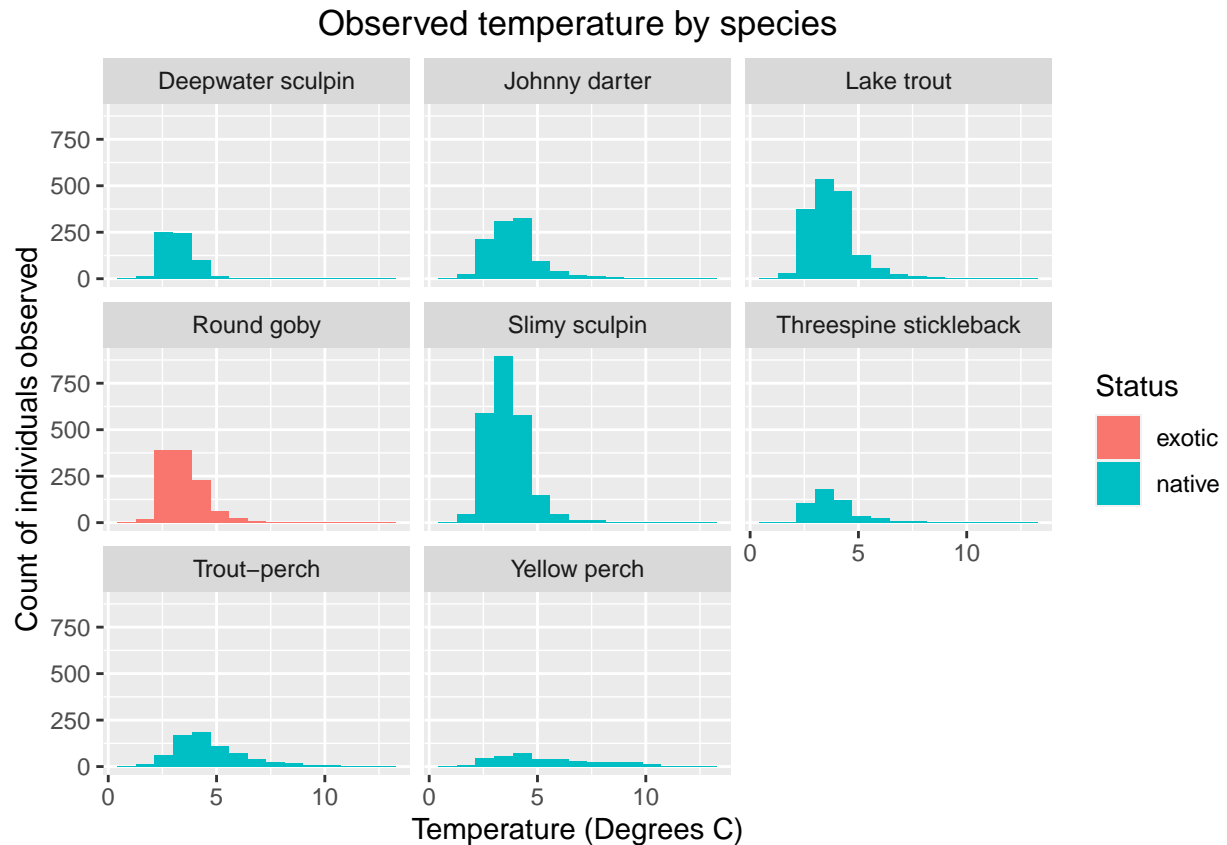
```
ggplot(fish.data, aes(x=opDate, y=fishingTemperature_C)) + geom_point(alpha=0.1) + geom_smooth() + labs
```

```
## 'geom_smooth()' using method = 'gam' and formula 'y ~ s(x, bs = "cs")'
```

Temperature values by date



```
ggplot(fish.data.clean, aes(x=fishingTemperature_C, fill=inv.status)) + geom_histogram(bins=15) + facet.
```



*#plotting the count of observations of each species depending on the temperature.*

Lets visualize the percentage of each species

```
# Proportion of the total catch from the first siting in 1997 -->
# Based on abundance
fish.data.clean %>%
  group_by(commonName, year) %>%
  # filtering out year based on first time a goby was sighted --> in 1997
  filter(year %in% seq(1997, 2022)) %>%
  tally(n) %>% # tallying up occurrences of each species
  ggplot(aes(x=year, y=n, fill=commonName)) + geom_bar(position="fill", stat="identity") + labs(title="")
```

