Assignment 3: Data Exploration

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OVERVIEW

This exercise accompanies the lessons in Environmental Data Analytics on Data Exploration.

Directions

- 1. Rename this file <FirstLast>_A03_DataExploration.Rmd (replacing <FirstLast> with your first and last name).
- 2. Change "Student Name" on line 3 (above) with your name.
- 3. Work through the steps, **creating code and output** that fulfill each instruction.
- 4. Assign a useful name to each code chunk and include ample comments with your code.
- 5. Be sure to **answer the questions** in this assignment document.
- 6. When you have completed the assignment, **Knit** the text and code into a single PDF file.
- 7. After Knitting, submit the completed exercise (PDF file) to the dropbox in Sakai.

TIP: If your code extends past the page when knit, tidy your code by manually inserting line breaks.

TIP: If your code fails to knit, check that no install.packages() or View() commands exist in your code.

Set up your R session

1. Check your working directory, load necessary packages (tidyverse, lubridate), and upload two datasets: the ECOTOX neonicotinoid dataset (ECOTOX_Neonicotinoids_Insects_raw.csv) and the Niwot Ridge NEON dataset for litter and woody debris (NEON_NIWO_Litter_massdata_2018-08_raw.csv). Name these datasets "Neonics" and "Litter", respectively. Be sure to include the subcommand to read strings in as factors.

getwd() #qot my working directory

[1] "/home/guest/EDA_Spring2024"

#Loaded the corresponding packages library(tidyverse)

```
## v dplyr
              1.1.3
                        v readr
                                    2.1.4
## v forcats
              1.0.0
                        v stringr
                                    1.5.0
              3.4.3
                                    3.2.1
## v ggplot2
                        v tibble
## v lubridate 1.9.2
                        v tidyr
                                    1.3.0
## v purrr
              1.0.2
                               ----- tidyverse conflicts() --
## -- Conflicts -----
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                    masks stats::lag()
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become error
library(lubridate)
#Reading the datasets
Neonics <- read.csv("./Data/Raw/ECOTOX_Neonicotinoids_Insects_raw.csv",stringsAsFactors = T)</pre>
Litter <- read.csv("./Data/Raw/NEON NIWO Litter massdata 2018-08 raw.csv", stringsAsFactors = T)
```

---- tidyverse 2.0.0 --

Learn about your system

-- Attaching core tidyverse packages ----

2. The neonicotinoid dataset was collected from the Environmental Protection Agency's ECOTOX Knowledgebase, a database for ecotoxicology research. Neonicotinoids are a class of insecticides used widely in agriculture. The dataset that has been pulled includes all studies published on insects. Why might we be interested in the ecotoxicology of neonicotinoids on insects? Feel free to do a brief internet search if you feel you need more background information.

Answer: Because they migh have devastating ecological impacts and pose risk on agriculture production. First, because they affect both target (e.g., corn rootworm, flea beetle) and nontarget insects (e.g., bees), causing -when not death- chronic sublethal effects on aquatic insects, birds and pollinators, impacting agriculture. Second, due to they are highly soluble in water, easing their transport away from the area of initial application, making the problem easily widespread.

3. The Niwot Ridge litter and woody debris dataset was collected from the National Ecological Observatory Network, which collectively includes 81 aquatic and terrestrial sites across 20 ecoclimatic domains. 32 of these sites sample forest litter and woody debris, and we will focus on the Niwot Ridge long-term ecological research (LTER) station in Colorado. Why might we be interested in studying litter and woody debris that falls to the ground in forests? Feel free to do a brief internet search if you feel you need more background information.

Answer: In the case of the woody debris, because of their ecological functions, being habitat for organisms, serve as an nutrient source, are a site for nitrogen fixation, and influence soil and sediment transport and storage. Besides, they play an essential role to the freshwater and estuarine ecosystems (the HJ Andrews Experimental Forest, 1986). Regarding the litter, this is directly involved in plant-soil interaction, helping to incorporate carbon and nutrients from plants into the soil, so it's relevant for nutrient cycling and, hence, in productivity in forest ecosystems (Giweta, 2020).

4. How is litter and woody debris sampled as part of the NEON network? Read the NEON_Litterfall_UserGuide.pdf document to learn more. List three pieces of salient information about the sampling methods here:

Answer: 1.Spatial Sampling Design: elevated and ground traps for litter. 2. Randomized trap placement in sites with >50% aerial cover of woody vegetation >2m in height.Targeted trap

placement in sites with < 50% cover of woody vegetation. 3. Temporal sampling: ground traps are sampled once per year; while for elevated ones, varies according to vegetation present at the site (more frequent in deciduous forest sites during senescence, and infrequent year-round at evergreen sites).

Obtain basic summaries of your data (Neonics)

5. What are the dimensions of the dataset?

```
dim(Neonics) #Consulting the dimension of my DS, I got that it has 4623 rows and 30 columns of data.
## [1] 4623 30
```

6. Using the summary function on the "Effect" column, determine the most common effects that are studied. Why might these effects specifically be of interest?

```
# Getting the data only for Effect
sort(summary(Neonics$Effect), decreasing = TRUE)
```

##	Population	Mortality	Behavior	Feeding behavior
##	1803	1493	360	255
##	Reproduction	Development	Avoidance	Genetics
##	197	136	102	82
##	Enzyme(s)	Growth	Morphology	Immunological
##	62	38	22	16
##	Accumulation	Intoxication	Biochemistry	Cell(s)
##	12	12	11	9
##	Physiology	Histology	Hormone(s)	
##	7	5	1	

Answer: The 3 most common studied effects are Population, Mortality and Behavior. With such information, we can try to test the Neonicotinoids' impacts on different species, not only changes in the population by higher levels of mortality, but also in behaviors, and try to test if they are attributable to neonicotinoids or not.

7. Using the summary function, determine the six most commonly studied species in the dataset (common name). What do these species have in common, and why might they be of interest over other insects? Feel free to do a brief internet search for more information if needed.[TIP: The sort() command can sort the output of the summary command...]

sort(summary(Neonics\$Species.Common.Name), decreasing = T) #query for frequency by species (sorted from

```
##
                                 (Other)
                                                                     Honey Bee
##
                                     670
                                                                            667
                         Parasitic Wasp
##
                                                       Buff Tailed Bumblebee
##
                                     285
                                                                            183
##
                   Carniolan Honey Bee
                                                                    Bumble Bee
##
                                                                            140
##
                                                              Japanese Beetle
                       Italian Honeybee
##
                                     113
                                                                             94
```

##	Asian Lady Beetle	Euonymus Scale
##	76	75
##	Wireworm	European Dark Bee
##	69	66
## ##	Minute Pirate Bug 62	Asian Citrus Psyllid 60
##	Parastic Wasp	Colorado Potato Beetle
##	58	57
##	Parasitoid Wasp	Erythrina Gall Wasp
##	51	49
##	Beetle Order	Snout Beetle Family, Weevil
##	47	True Pur Orden
## ##	Sevenspotted Lady Beetle 46	True Bug Order 45
##	Buff-tailed Bumblebee	Aphid Family
##	39	38
##	Cabbage Looper	Sweetpotato Whitefly
##	38	37
##	Braconid Wasp	Cotton Aphid
## ##	33 Predatory Mite	33 Ladybird Beetle Family
##	33	30
##	Parasitoid	Scarab Beetle
##	30	29
##	Spring Tiphia	Thrip Order
##	29	29
## ##	Ground Beetle Family 27	Rove Beetle Family 27
##	Tobacco Aphid	Chalcid Wasp
##	27	25
##	Convergent Lady Beetle	Stingless Bee
##	. 25	25
##	Spider/Mite Class	Tobacco Flea Beetle
## ##	24 Citrus Leafminer	24 Ladybird Beetle
##	23	23
##	Mason Bee	Mosquito
##	22	22
##	Argentine Ant	Beetle
##	Elathonded Appletree Perer	Harmad Oak Call Harn
## ##	Flatheaded Appletree Borer 20	Horned Oak Gall Wasp 20
##	Leaf Beetle Family	Potato Leafhopper
##	20	20
##	Tooth-necked Fungus Beetle	Codling Moth
##	20	19
##	Black-spotted Lady Beetle	Calico Scale
## ##	18 Fairyfly Parasitoid	18 Lady Beetle
##	18	18
##	Minute Parasitic Wasps	Mirid Bug
##	18	18
##	Mulberry Pyralid	Silkworm
##	18	18

##	Vedalia Beetle	Araneoid Spider Order
##	18	17
##	Bee Order	Egg Parasitoid
##	17	17
##	Insect Class	Moth And Butterfly Order
##	17	17
##	Oystershell Scale Parasitoid	Hemlock Woolly Adelgid Lady Beetle
##	17	16
##	Hemlock Wooly Adelgid	Mite
##	16	16
##	Onion Thrip	Western Flower Thrips
##	16	15
##	Corn Earworm	Green Peach Aphid
##	14	14
##	House Fly	Ox Beetle
##	14	14
##	Red Scale Parasite	Spined Soldier Bug
##	14	14
##	Armoured Scale Family	Diamondback Moth
##	13	13
##	Eulophid Wasp	Monarch Butterfly
##	13	13
##	Predatory Bug	Yellow Fever Mosquito
##	13	13
##	Braconid Parasitoid	Common Thrip
##	12	12
##	Eastern Subterranean Termite	Jassid
##	12	12
##	Mite Order 12	Pea Aphid 12
## ##		
##	Pond Wolf Spider 12	Spotless Ladybird Beetle 11
##		Lacewing
##	Glasshouse Potato Wasp 10	Lacewing 10
##	Southern House Mosquito	Two Spotted Lady Beetle
##	10	10 spotted Lady Beetle
##	Ant Family	Apple Maggot
##	Ant ramily	4ppie Haggot 9
ππ	3	3

Answer: The most common estudied species are the Honey Bee, the Parasitic Wasp, the Buff Tailed Bumblebee, the Carniolan Honey Bee, the Bumble Bee, and the Italian Honeybee.

8. Concentrations are always a numeric value. What is the class of Conc.1..Author. column in the dataset, and why is it not numeric?

```
class(Neonics$Conc.1..Author.)
```

[1] "factor"

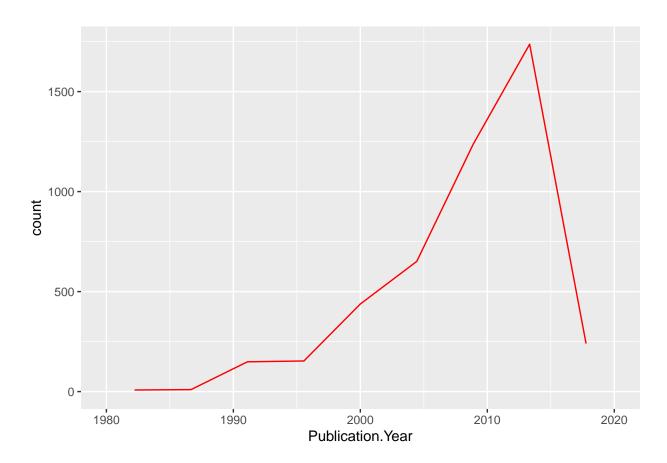
Answer: It is a Factor type, as contains symbols such as " \sim " and "/".

Explore your data graphically (Neonics)

9. Using geom_freqpoly, generate a plot of the number of studies conducted by publication year.

```
#Create the plot
ggplot(Neonics) +
  geom_freqpoly(aes(x = Publication.Year), bins = 10, color = "red", lty = 1) + scale_x_continuous(line)
```

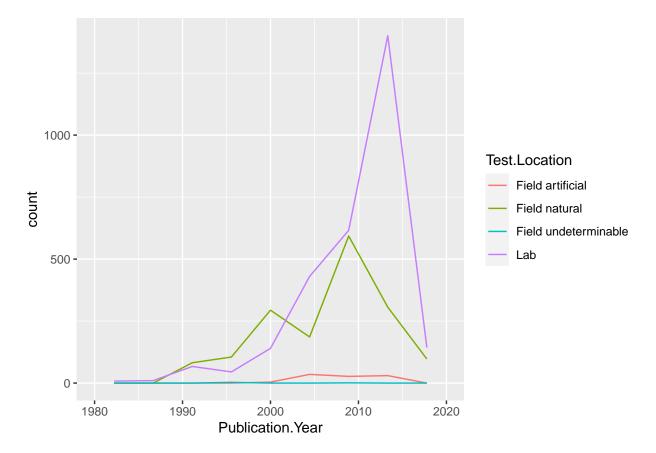
Warning: Removed 2 rows containing missing values ('geom_path()').



10. Reproduce the same graph but now add a color aesthetic so that different Test.Location are displayed as different colors.

```
ggplot(Neonics) +
  geom_freqpoly(aes(x = Publication.Year, group = Test.Location, color = Test.Location), bins = 10, lty
  scale_x_continuous(limits = c(1980, 2020))
```

Warning: Removed 8 rows containing missing values ('geom_path()').



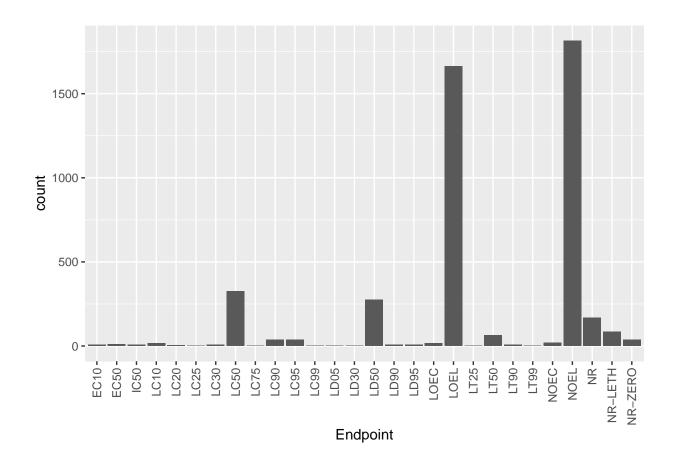
Interpret this graph. What are the most common test locations, and do they differ over time?

Answer: All along the plot, we observe that Lab and Field natural are the most common test locations. Although "Field natural" were the main type of test location for decades, in early 2000's years the use of lab gain relevance an became the most used type. Even for the last years, when the number of publications dropped dramatically, the lab is still more used than field natural.

11. Create a bar graph of Endpoint counts. What are the two most common end points, and how are they defined? Consult the ECOTOX_CodeAppendix for more information.

[TIP: Add theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1)) to the end of your plot command to rotate and align the X-axis labels...]

```
ggplot(Neonics, aes(x = Endpoint)) +
geom_bar() + theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))
```



Answer: The two most common are: 1) No-observable-effect-level (NOEL), which means that the highest dose (concentration) does not produce effects significantly different from responses of controls; and 2) Lowest-observable-effect-level (LOEL), that means that lowest dose (concentration) produce effects significantly different from responses of controls.

Explore your data (Litter)

12. Determine the class of collectDate. Is it a date? If not, change to a date and confirm the new class of the variable. Using the unique function, determine which dates litter was sampled in August 2018.

```
class(Litter$collectDate) #It's a factor type, so below I will put it as a date type

## [1] "factor"

Litter$collectDate <- as.Date(Litter$collectDate, format = "%Y/%m/%d")

class(Litter$collectDate) #Now it is Date type

## [1] "Date"

Augustsampled_litter <- unique(Litter$collectDate[format(Litter$collectDate, "%Y/%m") == 2018/08])

print(Augustsampled_litter) #None of the values were unique, so all of them were sampled in August 2018</pre>
```

[1] NA

13. Using the unique function, determine how many plots were sampled at Niwot Ridge. How is the information obtained from unique different from that obtained from summary?

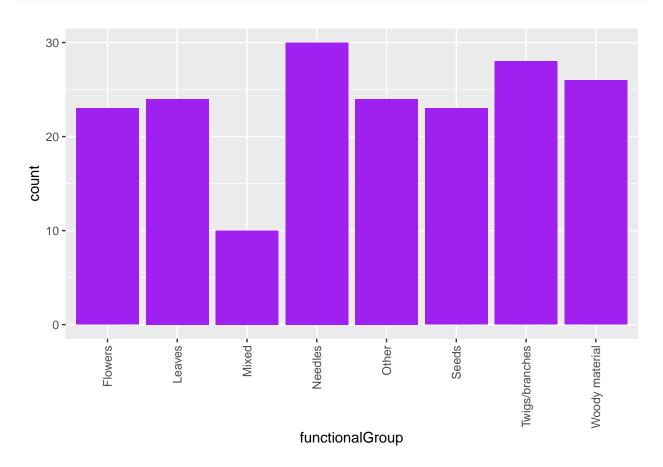
```
Niwot_Ridged_sampled <- unique(Litter$plotID)
print(Niwot_Ridged_sampled) #12 Plots were sampled</pre>
```

```
## [1] NIWO_061 NIWO_064 NIWO_067 NIWO_040 NIWO_041 NIWO_063 NIWO_047 NIWO_051 ## [9] NIWO_058 NIWO_046 NIWO_062 NIWO_057 ## 12 Levels: NIWO_040 NIWO_041 NIWO_046 NIWO_047 NIWO_051 NIWO_057 ... NIWO_067
```

Answer: unique() function eliminates the duplicate values or the rows, and returns only those that are unique; while summary returns the frequency of each value.

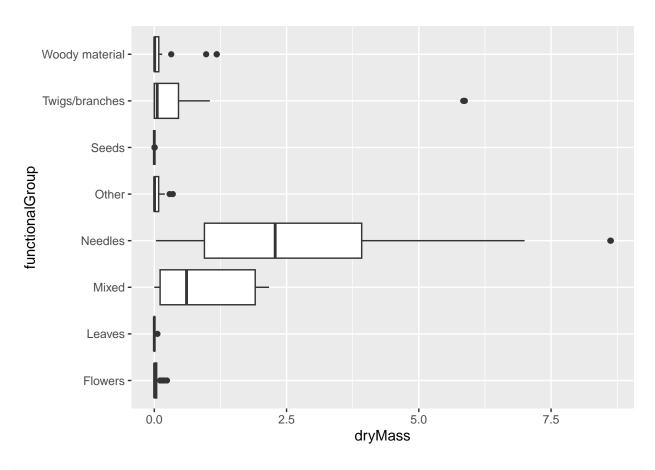
14. Create a bar graph of functional Group counts. This shows you what type of litter is collected at the Niwot Ridge sites. Notice that litter types are fairly equally distributed across the Niwot Ridge sites.

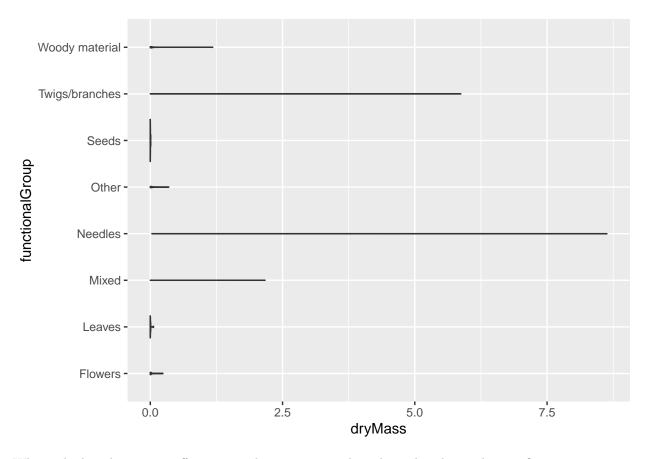
```
ggplot(Litter, aes(x = functionalGroup)) +
geom_bar(fill="purple") + theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))
```



15. Using geom_boxplot and geom_violin, create a boxplot and a violin plot of dryMass by functional-Group.

```
#For geom_boxplot
ggplot(Litter) +
geom_boxplot(aes(x = dryMass, y = functionalGroup))
```





Why is the boxplot a more effective visualization option than the violin plot in this case?

Answer: Because the violin plot shows distribution and as the size of the sample is small with several functional groups, it's not the most suitable option.

What type(s) of litter tend to have the highest biomass at these sites?

Answer: The needles.