Assignment 3: Data Exploration

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OVERVIEW

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

Directions

- 1. Change "Student Name, Section #" on line 3 (above) with your name and section number.
- 2. Work through the steps, creating code and output that fulfill each instruction.
- 3. Be sure to **answer the questions** in this assignment document.
- 4. When you have completed the assignment, **Knit** the text and code into a single PDF file.
- 5. After Knitting, submit the completed exercise (PDF file) to the dropbox in Sakai. Add your last name into the file name (e.g., "FirstLast_A03_DataExploration.Rmd") prior to submission.

The completed exercise is due on <>.

Set up your R session

1. Check your working directory, load necessary packages (tidyverse), 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 add the stringsAsFactors = TRUE parameter to the function when reading in the CSV files.

```
#Check working directory
getwd()
```

[1] "/Users/clara/Desktop/DATAN/Environmental_Data_Analytics_2022"

```
#Install necessary packages
require("tidyverse")

#Upload necessary datasets
Neonics<-read.csv("./Data/Raw/ECOTOX_Neonicotinoids_Insects_raw.csv", stringsAsFactors = TRUE)
Litter <-read.csv("./Data/Raw/NEON_NIWO_Litter_massdata_2018-08_raw.csv", stringsAsFactors = TRUE)</pre>
```

Learn about your system

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 ecotoxicologoy of neonicotinoids on insects? Feel free to do a brief internet search if you feel you need more background information.

Answer: We are interested in the ecotoxicology of neonicotinoids on insects to evaluate the ramifications of using this to the environment. The insectiside may do a lot more harm than good to surrounding biodiversity. For example, the insectiside may be absorbed by plants and become present in pollen and nectar which consequently poisons feeding bees and other pollen vectors. It is also necessary to evaluate whether these insectisides are directly or indirectly harmful to humans.

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: There are multiple reasons why we may be interested in studying litter and woody debris on forest grounds. Firstly, it is one way to get more information on how and at what rate nutrients are cycled in an ecosystem. Secondly, as woody debris and forest litter increases fuel load in many cases, it is important to keep track of their abundance and other relevant patterns. Thirdly, they may serve as indicators of the health of the forest. Additionally this data may simply reveal what species are present and abundant in the forest canopy.

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: * Litter and fine woody debris sampled using tower plots, which are selected randomly within a 90% flux footprint of primary and secondary airsheds. * In sites with forested tower airsheds, litter sampling takes place in 20 40mx40m tower plots. * In sites with low-statured vegetation over the tower airsheds, litter sampling takes place in 4 40mx40m and 26 20mx20m tower plots.

Obtain basic summaries of your data (Neonics)

5. What are the dimensions of the dataset?

print(dim(Neonics))

[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?

summary(Neonics\$Effect)

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

Answer: The most common effects studied are population (1803), mortality (1493), and behavior (360). These are of interest to determine whether the insectiside has altered population sizes of insects, in other words, to evaluate whether the insectiside harms insects, and/or alters their feeding patterns.

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.

summary(Neonics\$Species.Common.Name)

##	Honey Bee	Parasitic Wasp
##	667	285
##	Buff Tailed Bumblebee	Carniolan Honey Bee
##	183	152
##	Bumble Bee	Italian Honeybee
##	140	113
##	Japanese Beetle	Asian Lady Beetle
##	94	76
##	Euonymus Scale	Wireworm
##	75	Minute Pinete Pun
##	European Dark Bee	Minute Pirate Bug
##	66	62 Paranti a Mara
##	Asian Citrus Psyllid	Parastic Wasp
##	60	Demographed Hoge
## ##	Colorado Potato Beetle 57	Parasitoid Wasp 51
##	Erythrina Gall Wasp	Beetle Order
##	Erythiina Gair wasp 49	beetle bider 47
##	Snout Beetle Family, Weevil	Sevenspotted Lady Beetle
##	47	46
##	True Bug Order	Buff-tailed Bumblebee
##	45	39
##	Aphid Family	Cabbage Looper
##	38	38
##	Sweetpotato Whitefly	Braconid Wasp
##	37	33
##	Cotton Aphid	Predatory Mite
##	33	33
##	Ladybird Beetle Family	Parasitoid
##	30	30
##	Scarab Beetle	Spring Tiphia
##	29	29
##	Thrip Order	Ground Beetle Family
##	29	27
##	Rove Beetle Family	Tobacco Aphid
##	27	27
##	Chalcid Wasp	Convergent Lady Beetle
##	25	25
##	Stingless Bee	Spider/Mite Class
##	25	24
##	Tobacco Flea Beetle	Citrus Leafminer
##	24	23

##	Ladybird Beetle	Mason Bee
##	23	22
##	Mosquito	Argentine Ant
##	22	21
##	Beetle	Flatheaded Appletree Borer
##	21	20
##	Horned Oak Gall Wasp	Leaf Beetle Family
##	20	Tooth marked Fungua Bootle
##	Potato Leafhopper 20	Tooth-necked Fungus Beetle 20
##	Codling Moth	Black-spotted Lady Beetle
##	19	18
##	Calico Scale	Fairyfly Parasitoid
##	18	18
##	Lady Beetle	Minute Parasitic Wasps
##	18	18
##	Mirid Bug	Mulberry Pyralid
##	18	18
##	Silkworm	Vedalia Beetle
##	18	18
##	Araneoid Spider Order	Bee Order
##	17	17
##	Egg Parasitoid	Insect Class
##	17	17
##	Moth And Butterfly Order	Oystershell Scale Parasitoid
##	17 Hemlock Woolly Adelgid Lady Beetle	17 Hemlock Wooly Adelgid
##	16	nemiock wooly kdeigid
##	Mite	Onion Thrip
##	16	16
##	Western Flower Thrips	Corn Earworm
##	15	14
##	Green Peach Aphid	House Fly
##	14	14
##	Ox Beetle	Red Scale Parasite
##	14	14
##	Spined Soldier Bug	Armoured Scale Family
##	14	13
##	Diamondback Moth	Eulophid Wasp
##	13	13
##	Monarch Butterfly	Predatory Bug
## ##	Yallar Farran Magazita	13 Braconid Parasitoid
##	Yellow Fever Mosquito	braconid Parasitoid 12
##	Common Thrip	Eastern Subterranean Termite
##	12	12
##	Jassid	Mite Order
##	12	12
##	Pea Aphid	Pond Wolf Spider
##	12	12
##	Spotless Ladybird Beetle	Glasshouse Potato Wasp
##	11	10
##	Lacewing	Southern House Mosquito
##	10	10

##	Two Spotted Lady Beetle	Ant Family
##	10	9
##	Apple Maggot	(Other)
##	9	670

Answer: The six most commonly studied species are: Honey bee; Parasitic wasp; Buff tailed bumblebee; Carniolan honey bee; Bumblee bee; and the Italian honeybee. These are all pollinator species/vectors of pollen. They may be of interest because they determine plant yield and consequently directly impact human food abundance.

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

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

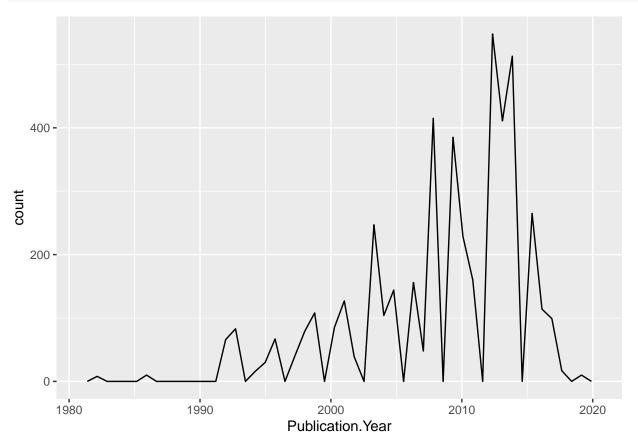
[1] "factor"

Answer: The class of Conc.1..Author is factor. It is not numeric because there are characters present in the data.

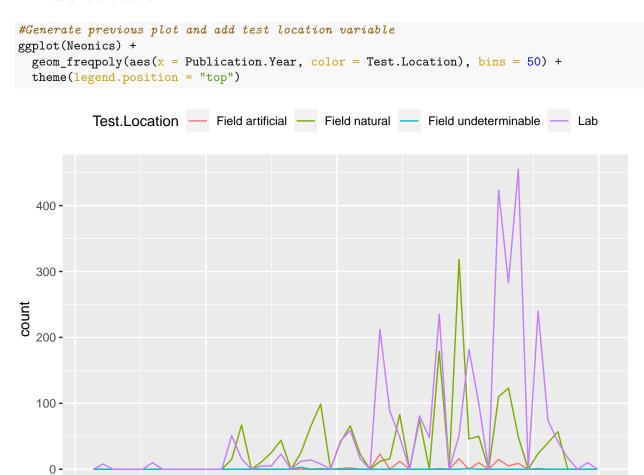
Explore your data graphically (Neonics)

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

```
#Generate plot of number of studies conducted by publication year
ggplot(Neonics) +
geom_freqpoly(aes(x = Publication.Year), bins = 50)
```



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



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

1990

1980

Answer: The most common test locations are the lab and the natural field. Previously the natural field and the lab test locations alternated in popularity, but more recently, the lab appears to be the most common location to conduct the tests.

2000

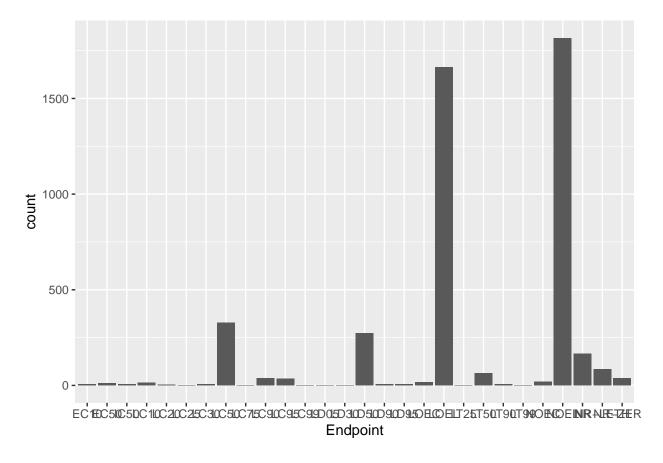
Publication.Year

2010

2020

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.

```
#Generate bar graph of Endpoint counts
ggplot(Neonics, aes(x = Endpoint)) +
  geom_bar()
```



Answer: The two most common end points are LOEL and NOEL. LOEL: Lowest-observable-effect-level; It is the lowest dose (concentration) producing effects that were significantly different from responses of controls. NOEL: No-observable-effect-level; It is the highest dose (concentration) producing effects not significantly different from responses of controls according to the author's reported statistical test.

Explore your data (Litter)

[1] "Date"

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.

```
#Determine class of collectDate
print(class(Litter$collectDate))

## [1] "factor"

#Change class to date
Litter$collectDate<-as.Date(Litter$collectDate)
print(class(Litter$collectDate))</pre>
```

```
#Determine which dates litter was sampled in August 2018
print(unique(Litter$collectDate, incomparables = FALSE))
```

```
## [1] "2018-08-02" "2018-08-30"
```

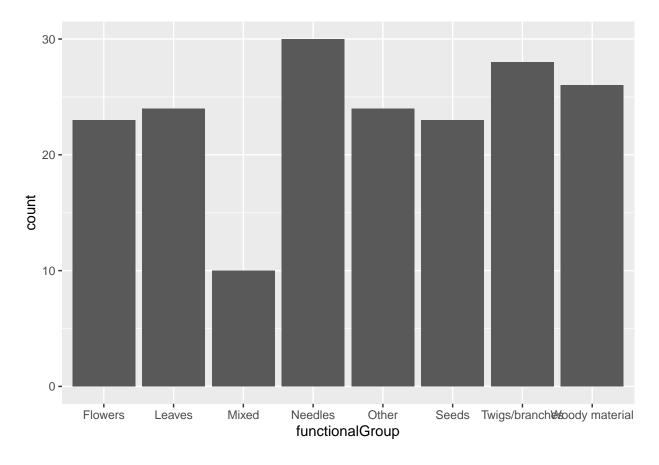
13. Using the unique function, determine how many plots were sampled at Niwot Ridge. How is the information obtained from unique different from that obtained from summary?

```
#Determine how many plots sampled at Niwot Ridge
print(unique(Litter$plotID, incomparables = FALSE))
   [1] NIWO_061 NIWO_064 NIWO_067 NIWO_040 NIWO_041 NIWO_063 NIWO_047 NIWO_051
   [9] NIWO_058 NIWO_046 NIWO_062 NIWO_057
## 12 Levels: NIWO_040 NIWO_041 NIWO_046 NIWO_047 NIWO_051 NIWO_057 ... NIWO_067
summary(Litter$plotID)
## NIWO_040 NIWO_041 NIWO_046 NIWO_047 NIWO_051 NIWO_057 NIWO_058 NIWO_061
##
         20
                                              14
                                                        8
                                                                         17
                  19
                           18
                                    15
## NIWO_062 NIWO_063 NIWO_064 NIWO_067
##
         14
                  14
                           16
                                    17
```

Answer: 12 plots were sampled at Niwot Ridge. Summary gives an overview of the plot ID data whereas the unique function allows you to identify and return a specific vector, data frame, or array without including duplicates. The unique function explicitly identifies that number of plots, but

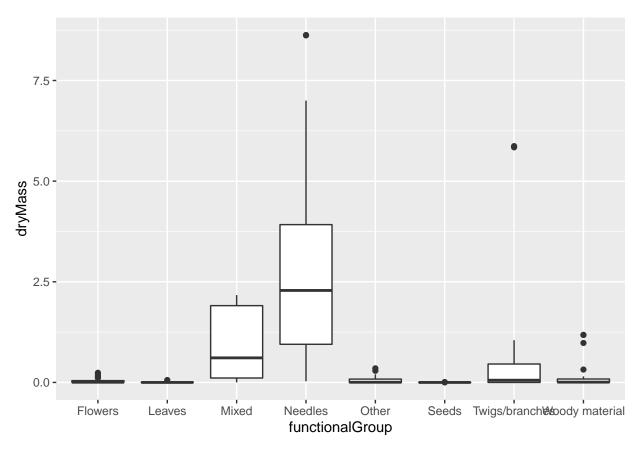
14. Create a bar graph of functional Group counts. This shows you what type of litter is collected at the Niwot Ridge sites. Notice that litter types are fairly equally distributed across the Niwot Ridge sites.

```
#Generate bar graph of functionalGroup counts
ggplot(Litter, aes(x = functionalGroup)) +
  geom_bar()
```



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

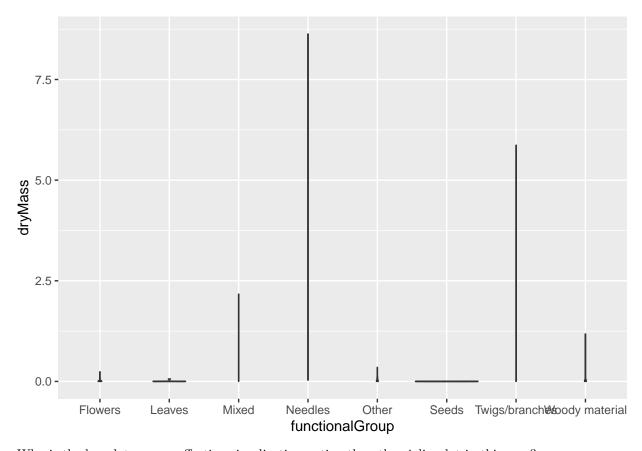
```
#Generate boxplot of dryMass by functionalGroup
ggplot(Litter) +
  geom_boxplot(aes(x = functionalGroup, y = dryMass))
```



```
## Warning in regularize.values(x, y, ties, missing(ties), na.rm = na.rm):
## collapsing to unique 'x' values

## Warning in regularize.values(x, y, ties, missing(ties), na.rm = na.rm):
## collapsing to unique 'x' values

## Warning in regularize.values(x, y, ties, missing(ties), na.rm = na.rm):
## collapsing to unique 'x' values
```



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

Answer: The violin plot is not an effective visualization option in this case because it does not depict the spread of the data.

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

Answer: Needles tends to have the highest biomass at these sites, as evident from the boxplot.