

# 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.

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## 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() # check working directory
```

```
## [1] "C:/Users/goode/OneDrive/Documents/Duke/ENV872_EDE/EDE_Fall2023"
```

```
library(tidyverse) # load in necessary packages
library(lubridate)
```

```
Neonics <- read.csv("../Data/Raw/ECOTOX_Neonicotinoids_Insects_raw.csv", stringsAsFactors = TRUE)
# upload the neonic dataset
```

```
Litter <- read.csv("../Data/Raw/NEON_NIWO_Litter_massdata_2018-08_raw.csv", stringsAsFactors = TRUE)
# upload the litter and woody debris dataset
```

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

Answer: Since neonicotinoids are insecticides, which means a chemical that has the purpose of killing insects, it is important to know how effective the neonicotinoids are at their job. Ecotox studies will provide information like what kind of dose and what duration of exposure/application methods are most effective at killing the desired species of insect, which is important knowledge for farmers or others trying to use the neonicotinoids in their agricultural work.

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: After a quick search of some publications from the USFS, I learned that woody debris in forests is important due to its role in the carbon budgets and nutrient cycling of forests. Woody debris also provides a source of energy for aquatic ecosystems and habitat for both terrestrial and aquatic organisms. It can also influence water flow and sediment transport.

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. Sampling takes place in terrestrial sites where there is woody vegetation that is more than 2m tall.  
2. Depending on vegetation coverage, the placement of traps can be random (sites with greater than 50% woody vegetation coverage) or targeted (sites with less than 50% woody vegetation coverage).  
3. In terms of temporal sampling, ground traps get samples once per year while elevated traps can get sampled once every 2 weeks or once every 1-2 months depending on the forest type.

## Obtain basic summaries of your data (Neonics)

5. What are the dimensions of the dataset?

```
dim(Neonics)
```

```
## [1] 4623 30
```

```
# finding the dimensions of the Neonics data set; 4623 rows and 30 columns
```

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
```

```
# finding how many of each type of effect are in the data set
```

Answer: Population is the most common effect studied (1803 counts) followed closely by mortality (1493 counts). Mortality refers to endpoints where the cause of death was directly related to the chemical of interest, and population refers to endpoints relating to a group of the same species in the same place at the same time. These are very important effects to study in insects because it will show if the chemical will perform its job and kill the insect and how the chemical effects a group of a particular species of insects.

- 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))
```

```
##      Ant Family      Apple Maggot
##           9           9
##      Glasshouse Potato Wasp      Lacewing
##          10           10
##      Southern House Mosquito      Two Spotted Lady Beetle
##          10           10
##      Spotless Ladybird Beetle      Braconid Parasitoid
##          11           12
##      Common Thrip      Eastern Subterranean Termite
##          12           12
##      Jassid      Mite Order
##          12           12
##      Pea Aphid      Pond Wolf Spider
##          12           12
##      Armoured Scale Family      Diamondback Moth
##          13           13
##      Eulophid Wasp      Monarch Butterfly
##          13           13
##      Predatory Bug      Yellow Fever Mosquito
##          13           13
##      Corn Earworm      Green Peach Aphid
##          14           14
```

##	House Fly	Ox Beetle
##	14	14
##	Red Scale Parasite	Spined Soldier Bug
##	14	14
##	Western Flower Thrips	Hemlock Woolly Adelgid Lady Beetle
##	15	16
##	Hemlock Woolly Adelgid	Mite
##	16	16
##	Onion Thrip	Araneoid Spider Order
##	16	17
##	Bee Order	Egg Parasitoid
##	17	17
##	Insect Class	Moth And Butterfly Order
##	17	17
##	Oystershell Scale Parasitoid	Black-spotted Lady Beetle
##	17	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
##	Codling Moth	Flatheaded Appletree Borer
##	19	20
##	Horned Oak Gall Wasp	Leaf Beetle Family
##	20	20
##	Potato Leafhopper	Tooth-necked Fungus Beetle
##	20	20
##	Argentine Ant	Beetle
##	21	21
##	Mason Bee	Mosquito
##	22	22
##	Citrus Leafminer	Ladybird Beetle
##	23	23
##	Spider/Mite Class	Tobacco Flea Beetle
##	24	24
##	Chalcid Wasp	Convergent Lady Beetle
##	25	25
##	Stingless Bee	Ground Beetle Family
##	25	27
##	Rove Beetle Family	Tobacco Aphid
##	27	27
##	Scarab Beetle	Spring Tiphia
##	29	29
##	Thrip Order	Ladybird Beetle Family
##	29	30
##	Parasitoid	Braconid Wasp
##	30	33
##	Cotton Aphid	Predatory Mite
##	33	33
##	Sweetpotato Whitefly	Aphid Family
##	37	38

##	Cabbage Looper	Buff-tailed Bumblebee
##	38	39
##	True Bug Order	Sevenspotted Lady Beetle
##	45	46
##	Beetle Order	Snout Beetle Family, Weevil
##	47	47
##	Erythrina Gall Wasp	Parasitoid Wasp
##	49	51
##	Colorado Potato Beetle	Parastic Wasp
##	57	58
##	Asian Citrus Psyllid	Minute Pirate Bug
##	60	62
##	European Dark Bee	Wireworm
##	66	69
##	Euonymus Scale	Asian Lady Beetle
##	75	76
##	Japanese Beetle	Italian Honeybee
##	94	113
##	Bumble Bee	Carniolan Honey Bee
##	140	152
##	Buff Tailed Bumblebee	Parasitic Wasp
##	183	285
##	Honey Bee	(Other)
##	667	670

```
# finding how many of each type of species are in the data set and
# ordering them from lowest to highest count
```

Answer: Not including “other” the six most commonly studeid species are honey bees, parasitic wasps, buff tailed bumblebees, Carniolan honey bees, bumble bees, and Italian Honeybees. These are all pollinators and are crucial for healthy ecosystems and successful agricultural operations (wasps are pollinators but are also beneficial becuase they can kill pests that would be harmful to crops). None of these species are harmful to crops and are beneficial to crops and ecosystems. There are likely lots of studies of these species to ensure that the insecticides aren’t harmful to or target these species.

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"
```

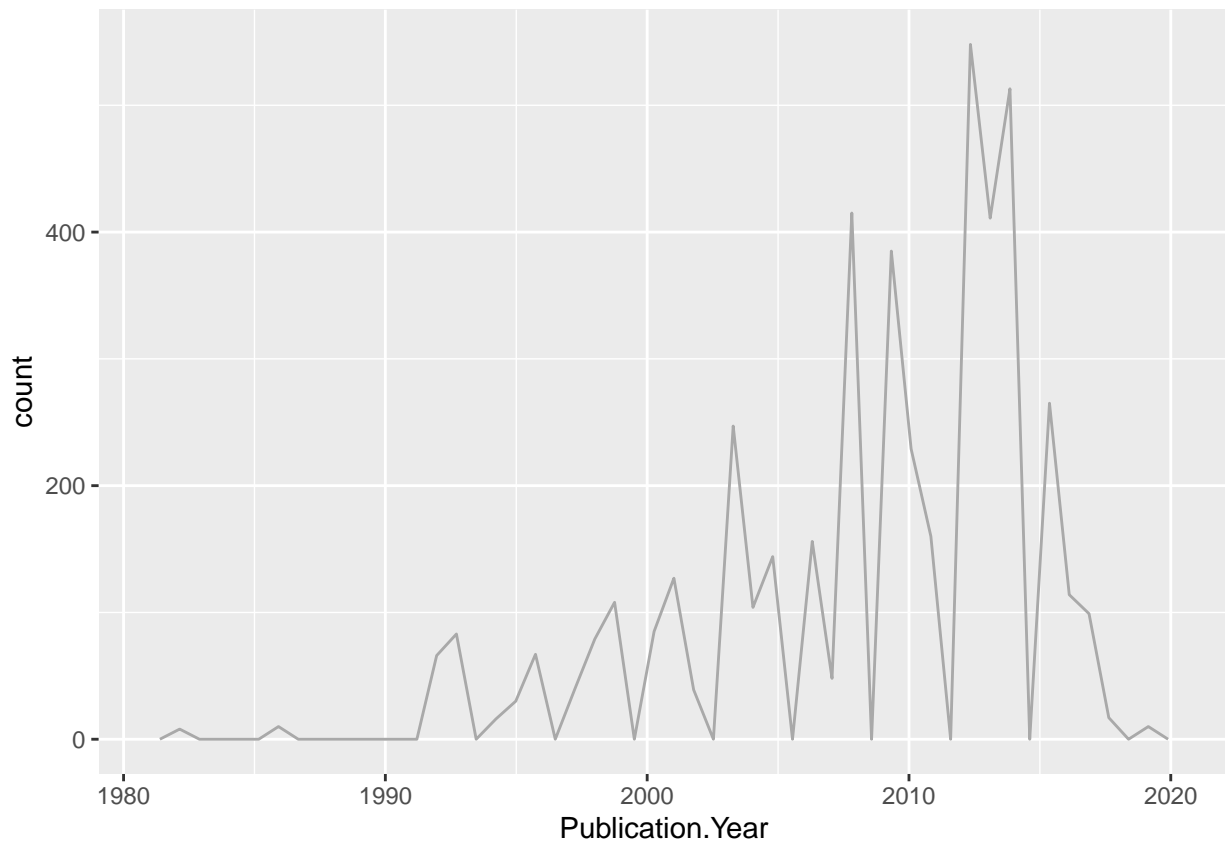
```
# class factor
```

Answer: This concentration variable is in the class factor. I’m guessing it’s factor and not numeric because the concentrations are discreet values in this case and are not continuous. There are also unusual notations in the variable, with some numbers written with a forward slash after them and a few cases where the variable is NR for not reported.

## Explore your data graphically (Neonics)

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

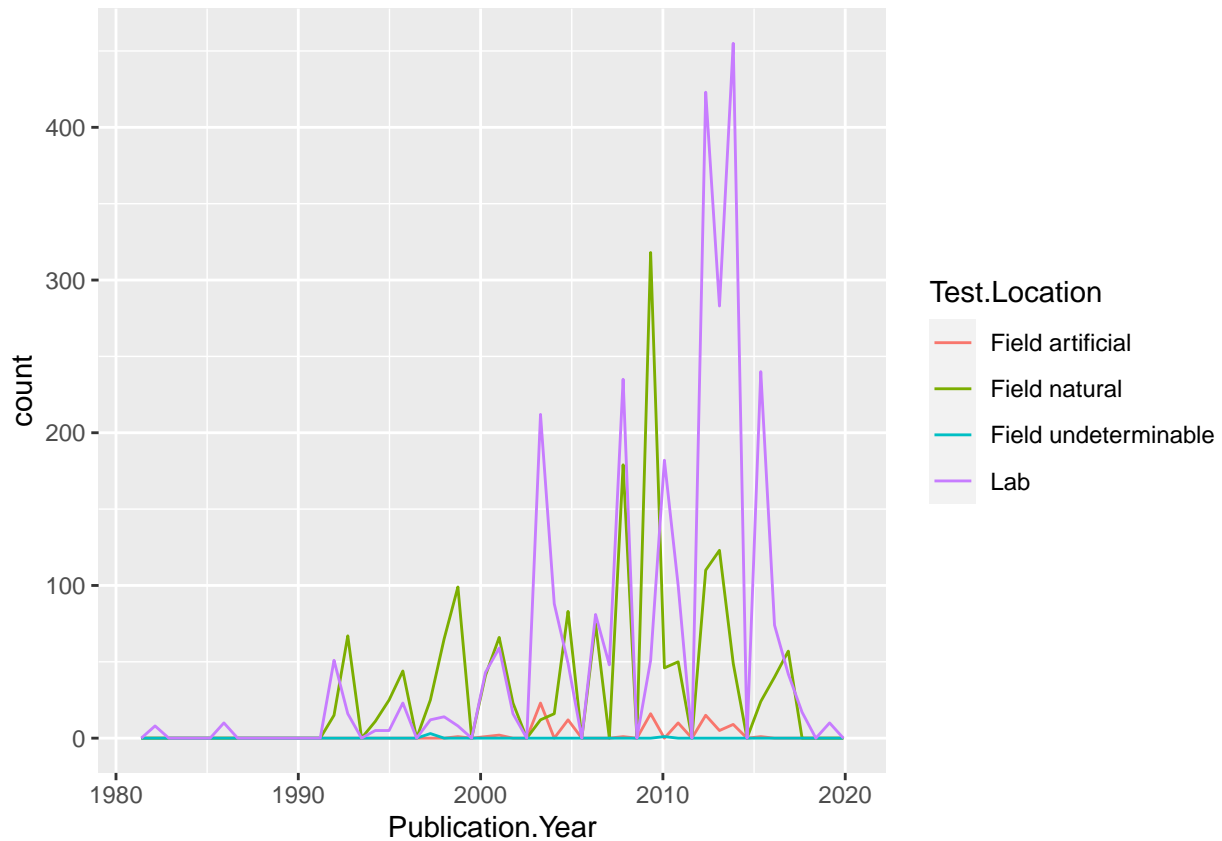
```
ggplot(Neonics) +  
  geom_freqpoly(aes(x = Publication.Year), bins = 50, color = "darkgray")
```



```
# creates a plot showing the publication year on the x axis and a count of  
# the number of studies published each year on the y axis
```

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, color = Test.Location), bins = 50)
```



```
# creates a plot of publications counts per year with a line for each type of
# test location; a legend of test locations is generated to the right of the
# plot
```

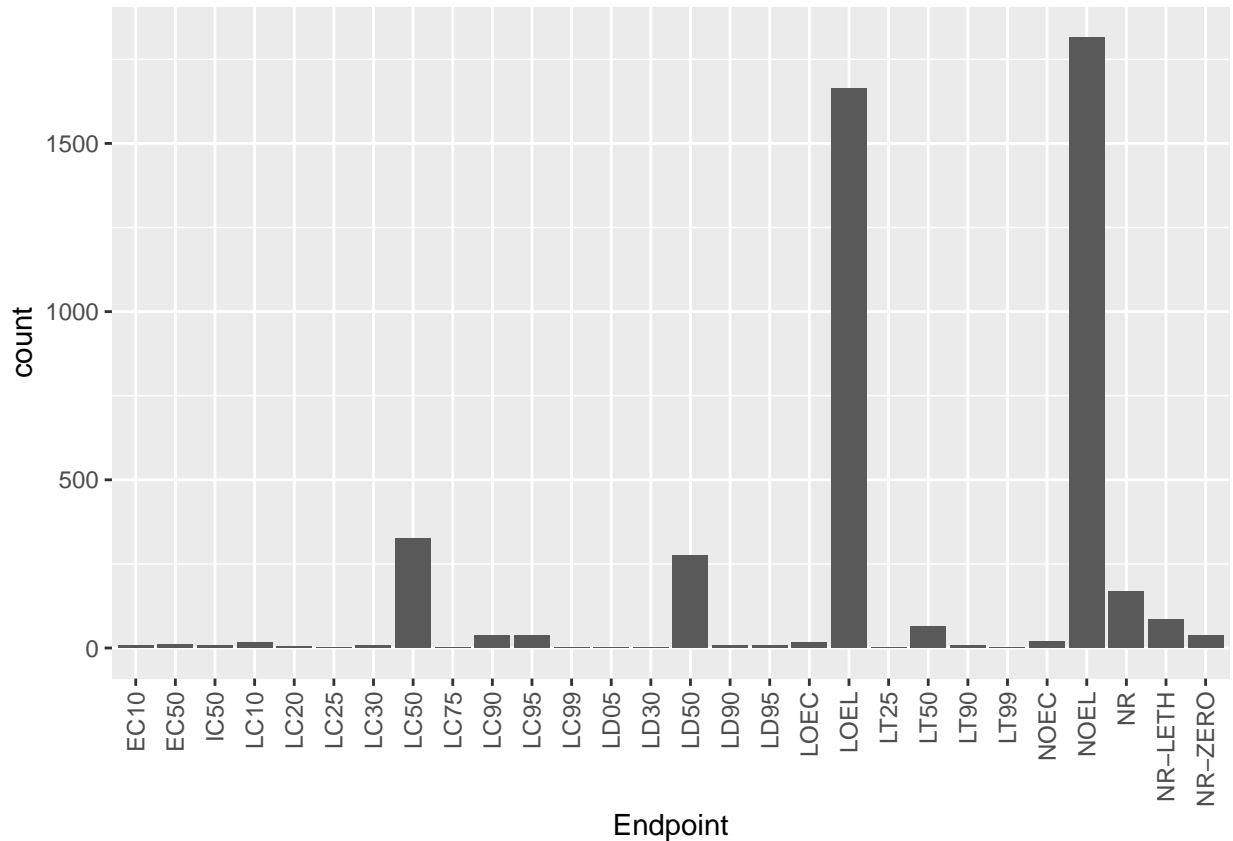
Interpret this graph. What are the most common test locations, and do they differ over time?

Answer: The most common test locations are lab and field natural. It looks like they started out being used relatively equally, with maybe the field natural being slightly more popular. After 2010, the lab studies became by far the most popular test location. It looks like field artificial and field undeterminable were never that popular.

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))
```



```
# creates a barplot with endpoint on the x axis and the count of each on
# the y axis
```

Answer: The two most common endpoints are NOEL and LOEL. NOEL stands for no observed effect level and is the highest dose producing effects not significantly different from responses of controls (the highest dose that is “safe”). LOEL stands for the lowest observed effect level, which means it was the lowest dose producing effects that were significantly different from responses of controls (the lowest dose that is harmful).

## Explore your data (Litter)

- 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 is not a date it is a factor
```

```
## [1] "factor"
```

```
library(lubridate)
Litter$collectDate <- lubridate::ymd(Litter$collectDate) # convert to class date
# using lubridate
```

```
class(Litter$collectDate) # the class is now date
```



```
## [1] "Date"
```

```
unique(Litter$collectDate) # sampling was done on 2018-08-02 and 2018-08-30
```

```
## [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`?

```
unique(Litter$plotID) # there were 12 plots sampled at Niwot
```

```
## [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) # so I can see the comparison of unique and summary
```

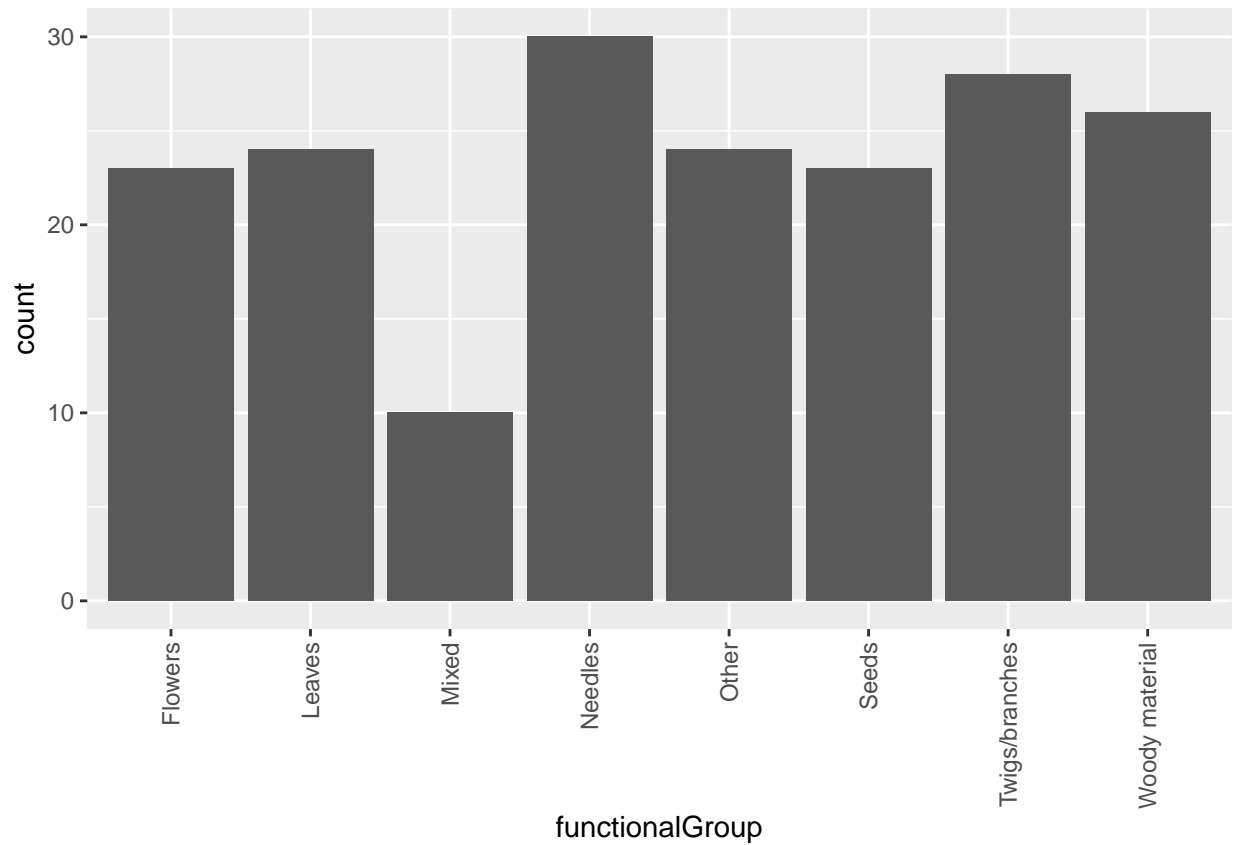
```
## NIWO_040 NIWO_041 NIWO_046 NIWO_047 NIWO_051 NIWO_057 NIWO_058 NIWO_061  
##      20      19      18      15      14       8      16      17  
## NIWO_062 NIWO_063 NIWO_064 NIWO_067  
##      14      14      16      17
```

```
# outputs
```

Answer: The `unique` function provides a list of each of the plots that was sampled at the site. It's picking out each unique `plotID` regardless of how many of each plot are in the variable to show that there were 12 plots sampled at the site. The `summary` function then provides a count of how many of each of the plots are in the data set. So it provides the list of the 12 plots and then a count of how many times they're in the data set (for example, NIWO\_040 has 20 entries).

14. Create a bar graph of `functionalGroup` 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.

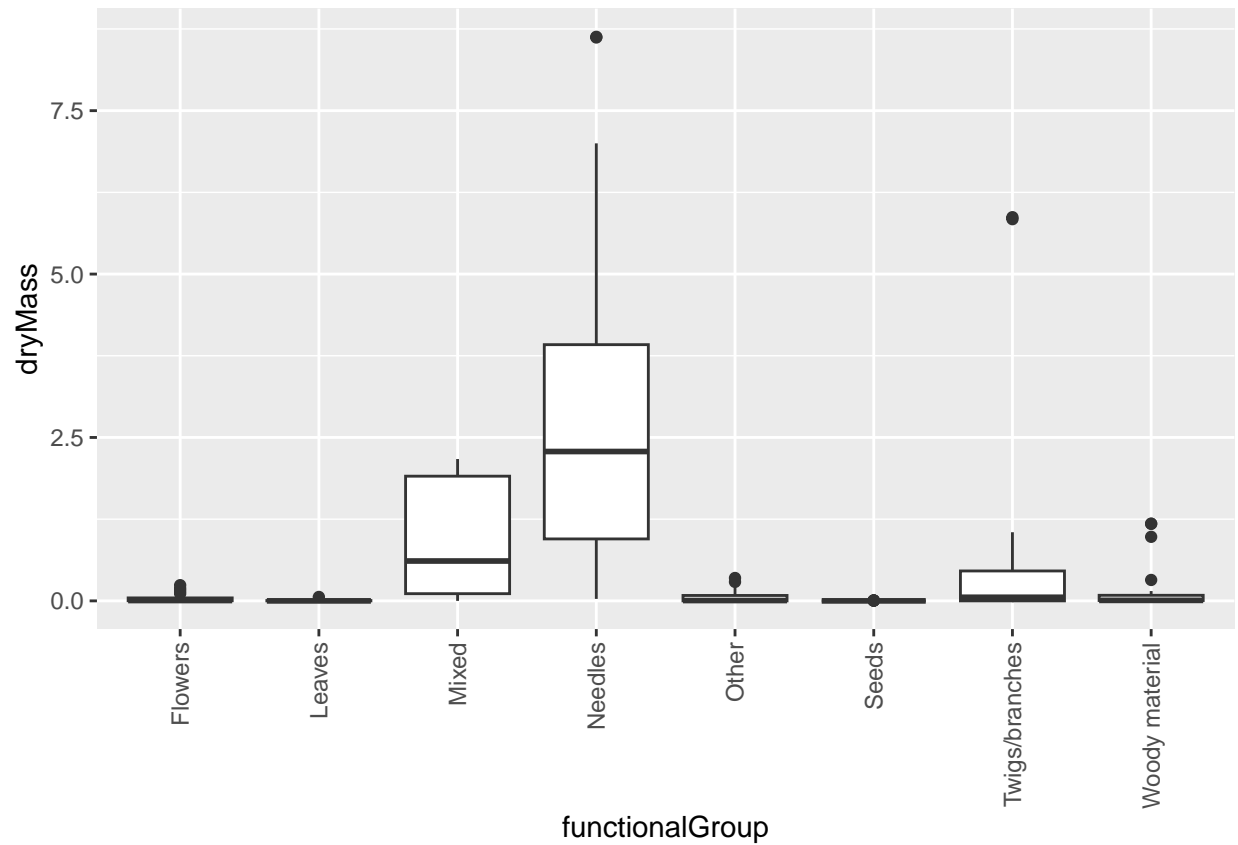
```
ggplot(Litter, aes(x = functionalGroup)) +  
  geom_bar() +  
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))
```



*# creates a bar plot with the type of litter on the x axis and the count of  
# each type on the y axis*

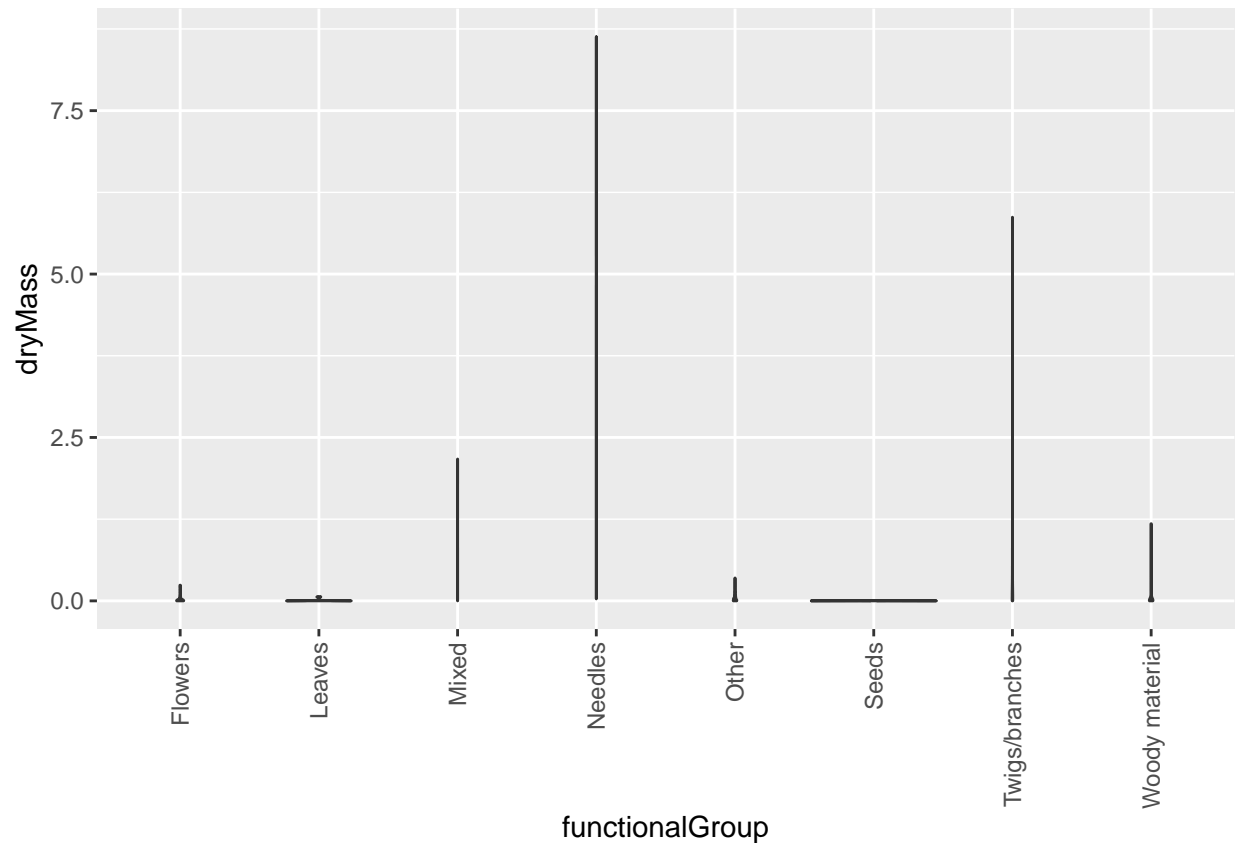
15. Using `geom_boxplot` and `geom_violin`, create a boxplot and a violin plot of `dryMass` by `functionalGroup`.

```
ggplot(Litter) +
  geom_boxplot(aes(x = functionalGroup, y = dryMass)) +
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))
```



```
# generate the boxplot
```

```
ggplot(Litter) +  
  geom_violin(aes(x = functionalGroup, y = dryMass)) +  
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))
```



```
# generate the violin plot
```

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

Answer: Violin plots display density distributions, but since there doesn't seem to be a large variation in density distribution in the types of litter with the largest dry masses, all you see are straight lines where you can't discern the summary statistics. Because the boxplot doesn't take into account the density distributions, you're able to see where the summary statistics are on the plot, making it more useful for this visualization.

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

Answer: Needles, mixed, and twigs/branches are the types of litter which tend to have the highest biomass at these sites.