

# Assignment 5: Data Visualization

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## OVERVIEW

This exercise accompanies the lessons in Environmental Data Analytics on Data Visualization

## Directions

1. Rename this file `<FirstLast>_A05_DataVisualization.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. Be sure your code is tidy; use line breaks to ensure your code fits in the knitted output.
  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.
- 

## Set up your session

1. Set up your session. Load the tidyverse, lubridate, here & cowplot packages, and verify your home directory. Read in the NTL-LTER processed data files for nutrients and chemistry/physics for Peter and Paul Lakes (use the tidy NTL-LTER\_Lake\_Chemistry\_Nutrients\_PeterPaul\_Processed.csv version in the Processed\_KEY folder) and the processed data file for the Niwot Ridge litter dataset (use the NEON\_NIWO\_Litter\_mass\_trap\_Processed.csv version, again from the Processed\_KEY folder).
2. Make sure R is reading dates as date format; if not change the format to date.

```
#1 set up session
```

```
library(tidyverse)
```

```
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr      1.1.3      v readr      2.1.4
## v forcats    1.0.0      v stringr   1.5.0
## v ggplot2    3.4.3      v tibble    3.2.1
## v lubridate  1.9.2      v tidyr     1.3.0
## v purrr      1.0.2
```

```
## -- Conflicts ----- tidyverse_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 errors
```

```
library(lubridate)
library(here)
```

```
## here() starts at C:/Users/Owner/OneDrive - Duke University/Documents/EDE_Fall2023
```

```
library(ggplot2)
library(cowplot)
```

```
##
## Attaching package: 'cowplot'
##
## The following object is masked from 'package:lubridate':
##
##     stamp
```

```
chem_nutrients <- read_csv("Data/Processed_KEY/NTL-LTER_Lake_Chemistry_Nutrients_PeterPaul_Processed.csv")
```

```
## Rows: 23008 Columns: 15
## -- Column specification -----
## Delimiter: ","
## chr   (1): lakename
## dbl  (13): year4, daynum, month, depth, temperature_C, dissolvedOxygen, irra...
## date  (1): sampleddate
##
## i Use 'spec()' to retrieve the full column specification for this data.
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
```

```
litter <- read_csv("Data/Processed_KEY/NEON_NIWO_Litter_mass_trap_Processed.csv")
```

```
## Rows: 1692 Columns: 13
## -- Column specification -----
## Delimiter: ","
## chr   (7): plotID, trapID, functionalGroup, qaDryMass, nlcdClass, plotType, g...
## dbl   (5): dryMass, subplotID, decimalLatitude, decimalLongitude, elevation
## date  (1): collectDate
##
## i Use 'spec()' to retrieve the full column specification for this data.
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
```

```
#2 reading dates as date format
```

```
chem_nutrients$sampleddate <- as.Date(chem_nutrients$sampleddate, format = "%Y/%m/%d")
```

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

## Define your theme

3. Build a theme and set it as your default theme. Customize the look of at least two of the following:

- Plot background

- Plot title
- Axis labels
- Axis ticks/gridlines
- Legend

```
#3 build theme
#change axis labels
#change legend position
my_theme <- theme(axis.text = element_text(color = "red"),
                  legend.position = "top")

#set as default theme
theme_set(my_theme)
```

## Create graphs

For numbers 4-7, create ggplot graphs and adjust aesthetics to follow best practices for data visualization. Ensure your theme, color palettes, axes, and additional aesthetics are edited accordingly.

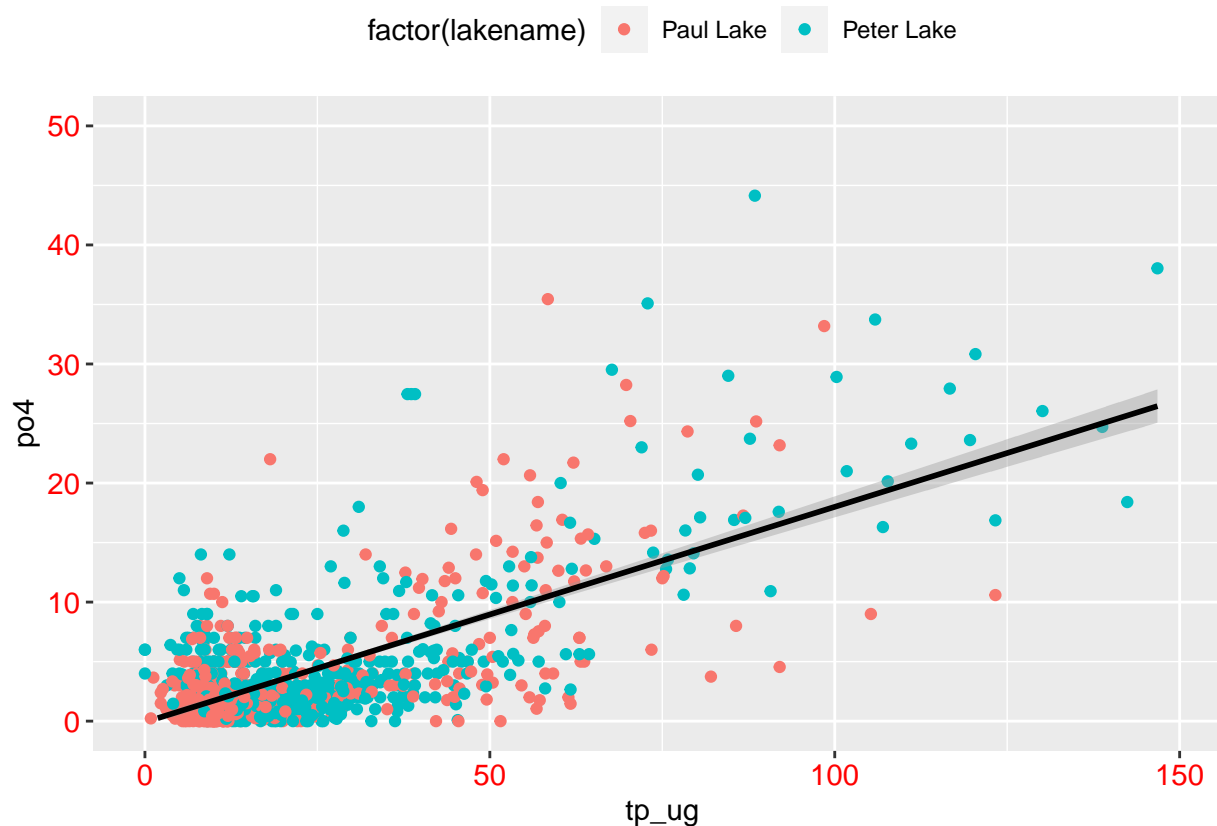
4. [NTL-LTER] Plot total phosphorus (tp<sub>ug</sub>) by phosphate (po<sub>4</sub>), with separate aesthetics for Peter and Paul lakes. Add a line of best fit and color it black. Adjust your axes to hide extreme values (hint: change the limits using `xlim()` and/or `ylim()`).

```
#4 total phosphorous v. phosphate
ggplot(chem_nutrients, aes(x = tp_ug, y = po4, color = factor(lakename)))+ #assign x and y, assign aest.
  geom_point()+ #scatterplot
  geom_smooth(method = "lm", formula = y ~ x, color = "black")+
#line of best fit
  xlim(0, 150) + #adjust x scale
  ylim(0, 50) #adjust y scale
```

```
## Warning: Removed 21948 rows containing non-finite values ('stat_smooth()').
```

```
## Warning: Removed 21948 rows containing missing values ('geom_point()').
```

```
## Warning: Removed 1 rows containing missing values ('geom_smooth()').
```



5. [NTL-LTER] Make three separate boxplots of (a) temperature, (b) TP, and (c) TN, with month as the x axis and lake as a color aesthetic. Then, create a cowplot that combines the three graphs. Make sure that only one legend is present and that graph axes are aligned.

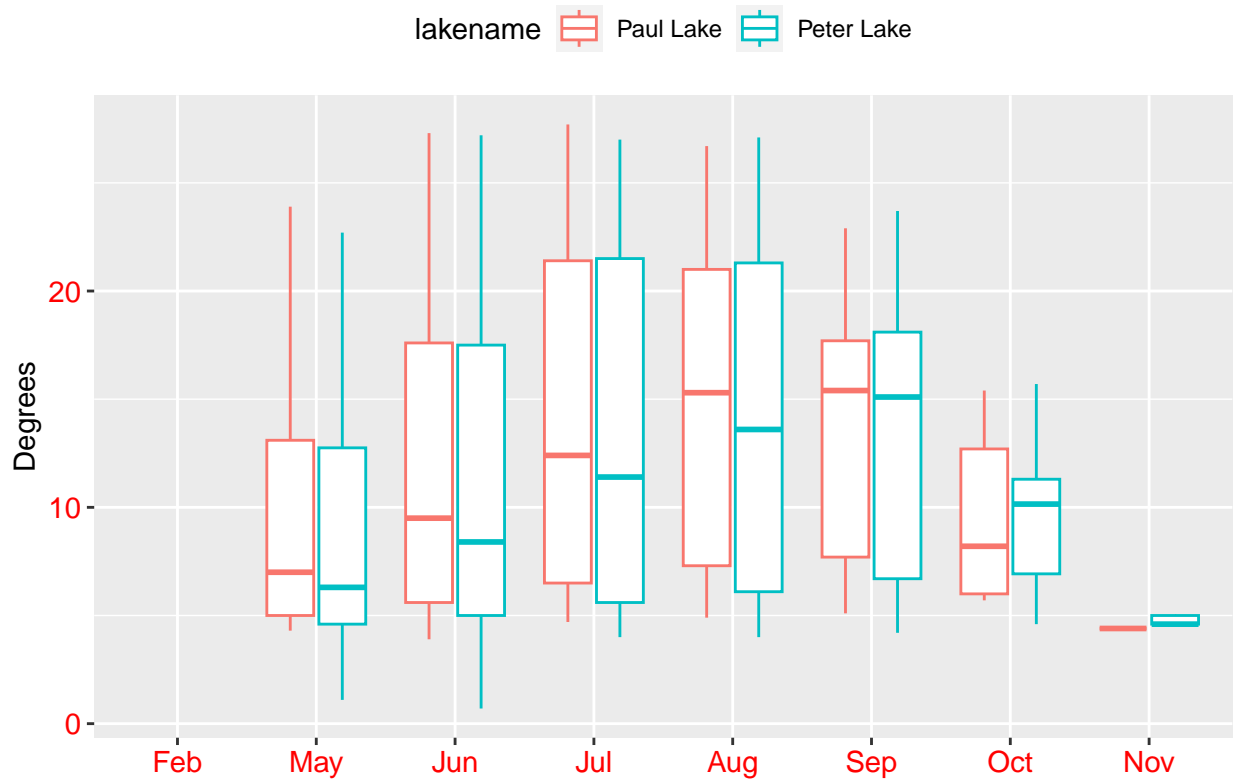
Tip: \* Recall the discussion on factors in the previous section as it may be helpful here. \* R has a built-in variable called `month.abb` that returns a list of months; see <https://r-lang.com/month-abb-in-r-with-example>

```
#5 3 boxplots
chem_nutrients$month <- factor(chem_nutrients$month,
  levels = 1:12,
  labels = month.abb)

#temperature boxplot
plot_temp <- ggplot(chem_nutrients, aes(x = factor(month), y = temperature_C, color = lakename))+
  geom_boxplot()+
  labs(
    x = NULL,
    y = "Degrees",
    title = "Temperature")
print(plot_temp)
```

```
## Warning: Removed 3566 rows containing non-finite values ('stat_boxplot()').
```

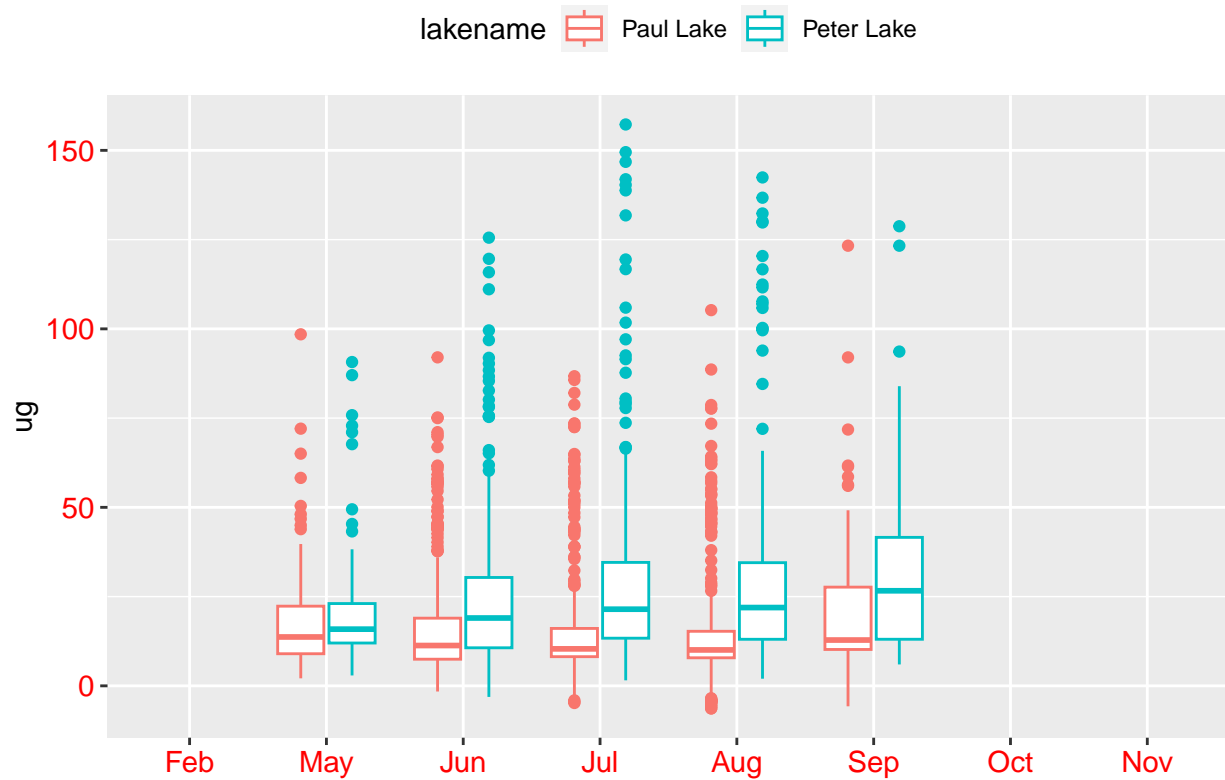
## Temperature



```
#TP boxplot
plot_tp <- ggplot(chem_nutrients, aes(x = factor(month), y = tp_ug, color = lakename))+
  geom_boxplot()+
  labs(
    x = NULL,
    y = "ug",
    title = "Total Phosphorus")
print(plot_tp)
```

## Warning: Removed 20729 rows containing non-finite values ('stat\_boxplot()').

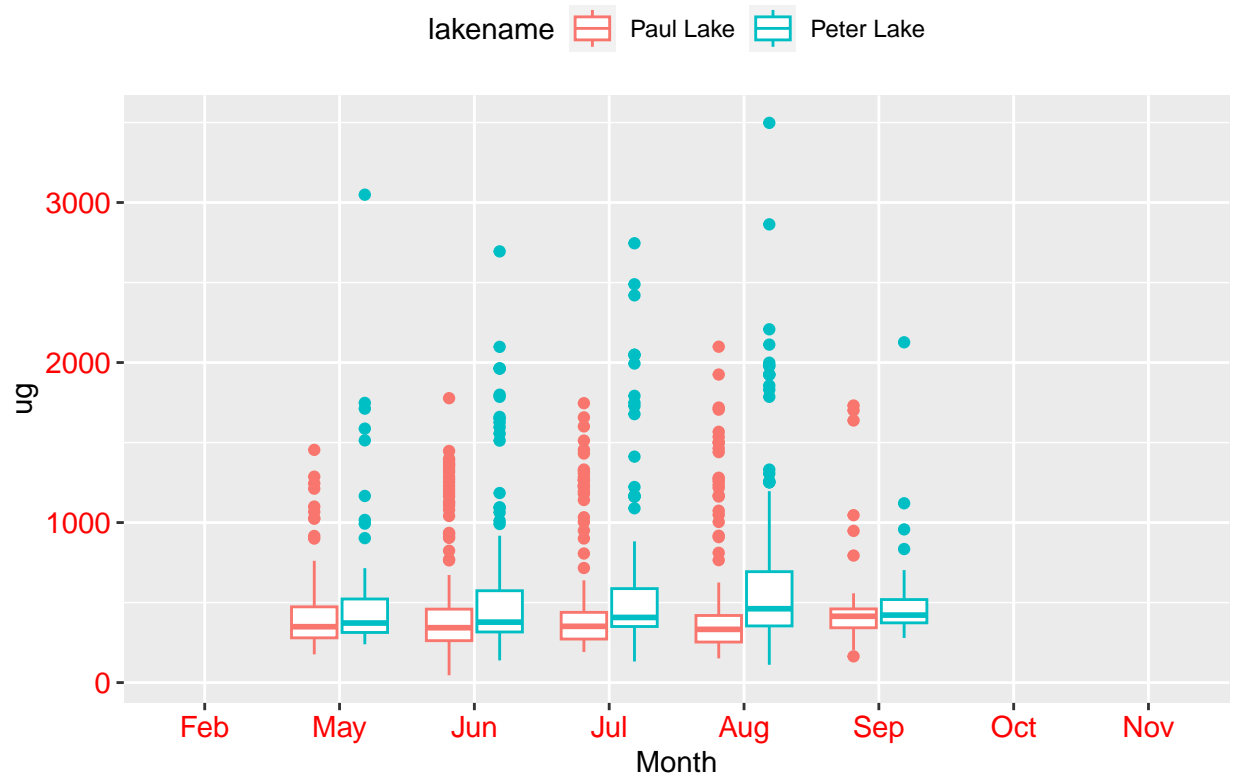
## Total Phosphorus



```
#TN boxplot
plot_tn <- ggplot(chem_nutrients, aes(x = factor(month), y = tn_ug, color = lakename))+
  geom_boxplot()+
  labs(
    x = "Month",
    y = "ug",
    title = "Total Nitrogen")
print(plot_tn)
```

## Warning: Removed 21583 rows containing non-finite values ('stat\_boxplot()').

## Total Nitrogen



```
#combine boxplots
combined_plots <- plot_grid(plot_temp + theme(legend.position = "top"),
                             plot_tp + theme(legend.position = "none"),
                             plot_tn + theme(legend.position = "none"), nrow = 3, ncol = 1)
```

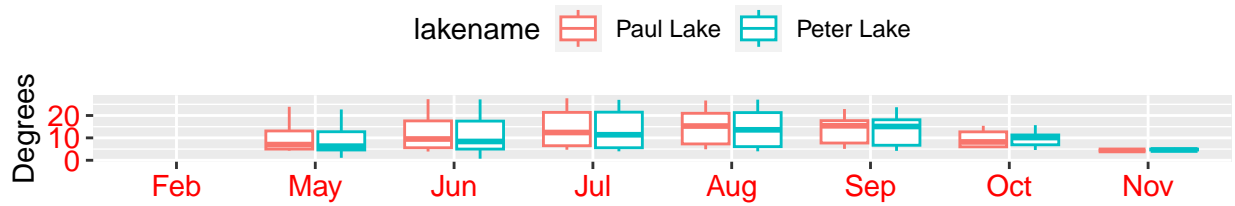
```
## Warning: Removed 3566 rows containing non-finite values ('stat_boxplot()').
```

```
## Warning: Removed 20729 rows containing non-finite values ('stat_boxplot()').
```

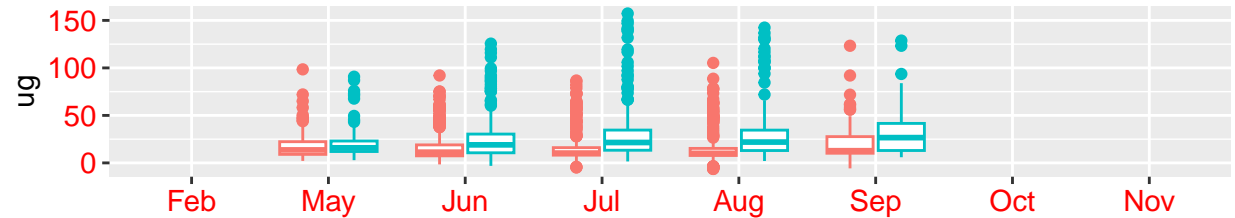
```
## Warning: Removed 21583 rows containing non-finite values ('stat_boxplot()').
```

```
print(combined_plots)
```

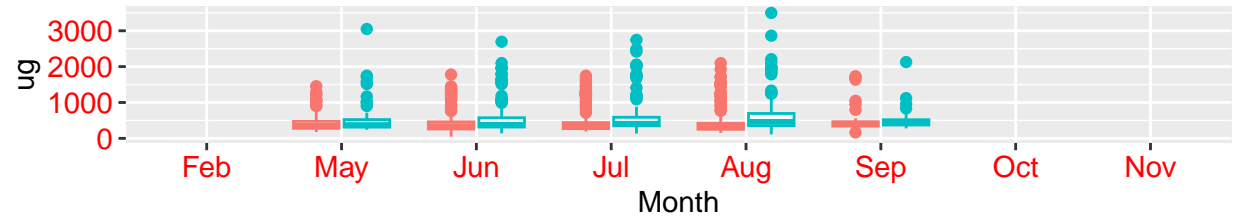
## Temperature



## Total Phosphorus



## Total Nitrogen



Question: What do you observe about the variables of interest over seasons and between lakes?

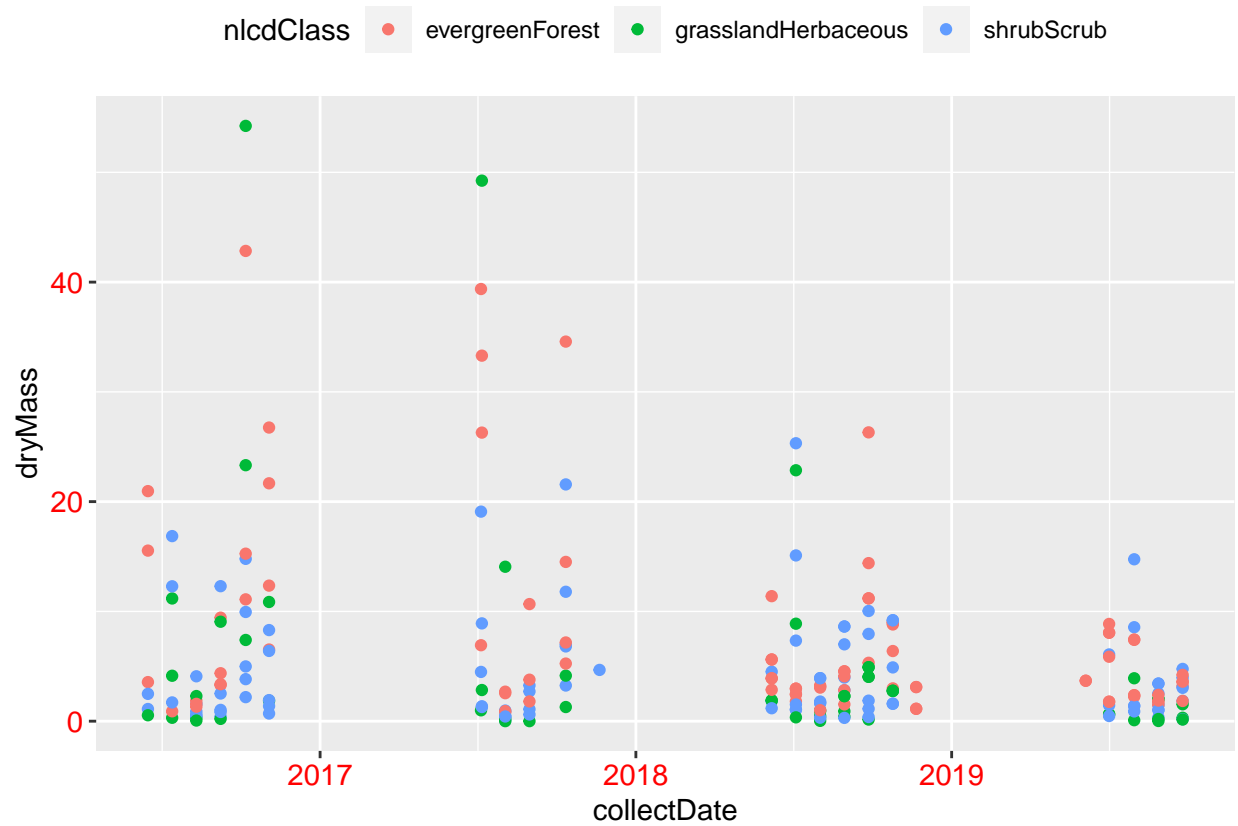
Answer: Peter Lake has higher temperature, tp, and tn than Paul Lake. Tn and Tp increase and decrease with temperature.

- [Niwot Ridge] Plot a subset of the litter dataset by displaying only the “Needles” functional group. Plot the dry mass of needle litter by date and separate by NLCD class with a color aesthetic. (no need to adjust the name of each land use)
- [Niwot Ridge] Now, plot the same plot but with NLCD classes separated into three facets rather than separated by color.

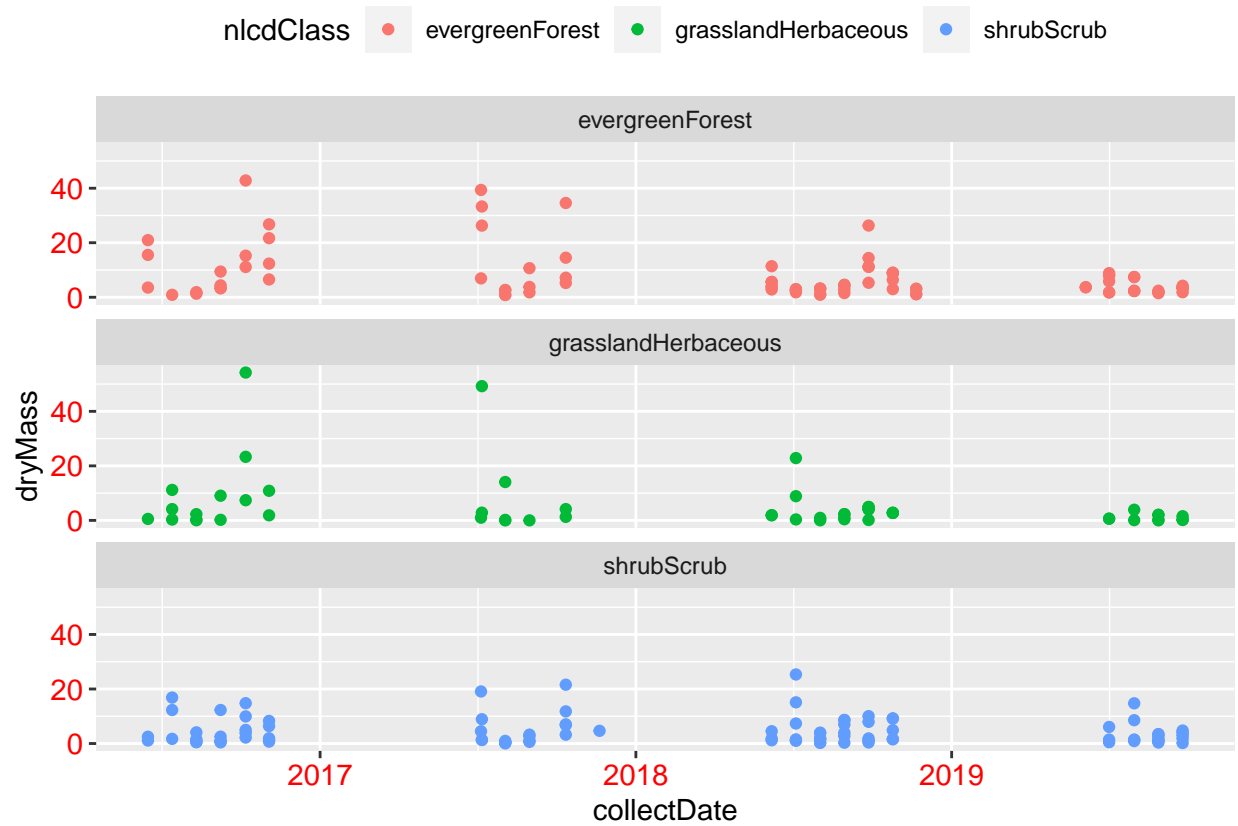
```
#6 plot histogram
litter$collectDate <- as.Date(litter$collectDate, format = "%Y/%m/%d")

needles_graph<- ggplot(subset(litter, functionalGroup == "Needles"),
  (aes(x = collectDate, y = dryMass, color = nlcdClass)))+
  geom_point()
print(needles_graph)
```





```
#7 plot
facet_plot <- ggplot(subset(litter, functionalGroup == "Needles"), (aes(x = collectDate, y = dryMass, color = nlcdClass)) +
  geom_point() +
  facet_wrap(vars(nlcdClass), nrow = 3))
print(facet_plot)
```



Question: Which of these plots (6 vs. 7) do you think is more effective, and why?

Answer: 7 is easier to understand because the data is separated into 3 graphs but aligned.