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.

```
library(tidyverse)
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr
             1.1.4
                       v readr
                                  2.1.5
## v forcats
              1.0.0
                                  1.5.1
                       v stringr
## v ggplot2
             3.4.4
                       v tibble
                                  3.2.1
## v lubridate 1.9.3
                       v tidyr
                                  1.3.0
## v purrr
                            ## -- Conflicts -----
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                   masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
```

```
library(lubridate)
library(here)
## here() starts at /Users/sarah/Documents/872_EDA/EDA_Spring2024
library(cowplot)
##
## Attaching package: 'cowplot'
## The following object is masked from 'package:lubridate':
##
##
       stamp
library(ggplot2)
library(ggthemes)
##
## Attaching package: 'ggthemes'
## The following object is masked from 'package:cowplot':
##
##
       theme_map
processed_data = "Data/Processed_KEY"
PeterPaul.chem.nutrients <- read.csv(</pre>
    processed_data,
    "NTL-LTER_Lake_Chemistry_Nutrients_PeterPaul_Processed.csv"),
  stringsAsFactors = TRUE
Niwot.Ridge.Litter <- read.csv(</pre>
    processed_data, "NEON_NIWO_Litter_mass_trap_Processed.csv"),
  stringsAsFactors = TRUE
```

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
A05_theme <- theme_base() + theme(
   plot.title = element_text( #Updating Title
      color= 'darkslategray',
     size = 14
   ),
   axis.text = element_text( #Updating axis text
      color = "black",
     size = 10
   ),
   axis.title.x = element_text( #Updating x-axis
     color = "black",
     size = 12
   ),
   axis.title.y = element_text( #Updating y-axis
     color = "black",
     size = 12
   legend.position = "top", #Putting legend to top
    legend.title = element_text( #Updating legend
      color='black',
      size = 12
   ),
  legend.text = element_text( #Updating legend text
     size = 12
   plot.background = element_blank() #takeaway plot edge
```

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.

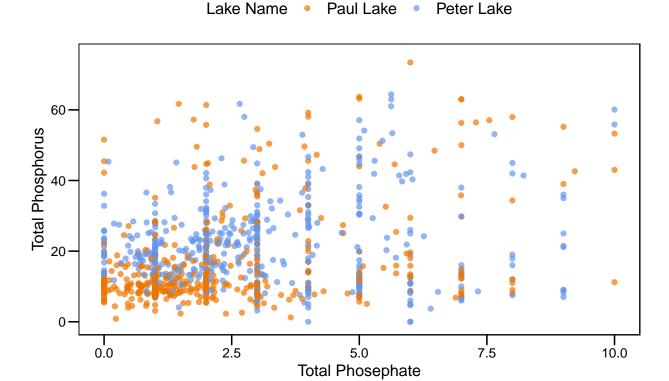
4. [NTL-LTER] Plot total phosphorus (tp_ug) by phosphate (po4), with separate aesthetics for Peter and Paul lakes. Add line(s) of best fit using the lm method. Adjust your axes to hide extreme values (hint: change the limits using xlim() and/or ylim()).

```
#4
plot4 <- PeterPaul.chem.nutrients %>%
    ggplot(aes(
        x=po4,
        y=tp_ug,
        color=lakename)
    ) +
    geom_point(alpha=0.70) +
    xlim(0, 10) + # is this the correct way to hide the extreme values? should i keep it at 5, 10, 25?
    ylim(0, 75) +
    ylab("Total Phosphorus") +
    xlab("Total Phosphorus by Phosphate at Paul Lake and Peter Lake") +
    ggtitle("Phosphorus by Phosphate at Paul Lake and Peter Lake") +
    scale_color_manual(values = c("darkorange2", "cornflowerblue"),
```

```
name = "Lake Name") +
A05_theme
print(plot4)
```

Warning: Removed 22064 rows containing missing values ('geom_point()').

Phosphorus by Phosphate at Paul Lake and Peter Lake



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.

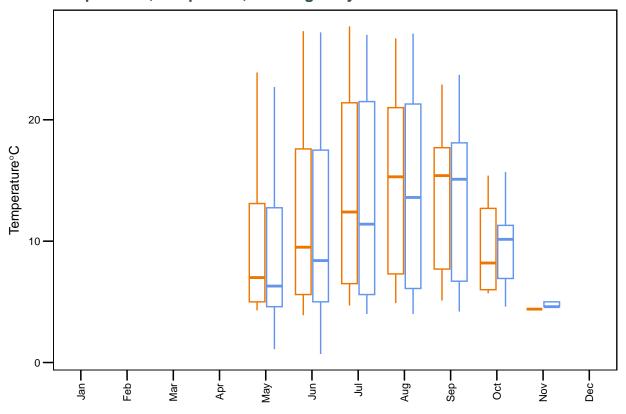
Tips: * Recall the discussion on factors in the lab section as it may be helpful here. * Setting an axis title in your theme to element_blank() removes the axis title (useful when multiple, aligned plots use the same axis values) * Setting a legend's position to "none" will remove the legend from a plot. * Individual plots can have different sizes when combined using cowplot.

```
#5
A05_theme1 <- theme_base() + theme(
    plot.title = element_text(
        color= 'darkslategray',
        size = 12
),
    axis.text = element_text(
        color = "black",
        size = 8</pre>
```

```
axis.title.x = element_blank(), #leaving x-axis blank
   axis.title.y = element_text(
    color = "black",
     size = 10
     legend.title = element_text(
      color='black',
     size = 8
    ),
   legend.text = element_text(
    size = 8
     ),
   axis.text.x = element_text(
     angle = 90,
     vjust = 0.5,
    hjust=1),
   legend.position = "top",
   plot.background = element_blank()
PeterPaul.chem.nutrients$month_f <- factor(</pre>
  PeterPaul.chem.nutrients$month,
  levels=1:12,
  labels = month.abb
temp <- PeterPaul.chem.nutrients %>%
  ggplot(aes(
     x=month_f,
      y=temperature_C,
      color=lakename)
    ) +
  geom_boxplot() +
  ylab(expression(paste("Temperature", degree, "C"))) +
  ggtitle("Temperature, Phophorus, & Nitrogen by Month") +
  scale_color_manual(values = c("darkorange2", "cornflowerblue"),
                   name = "Lake Name") +
  A05_theme1 +
  theme(legend.position = "NONE") +
  scale_x_discrete(drop=FALSE)
temp
```

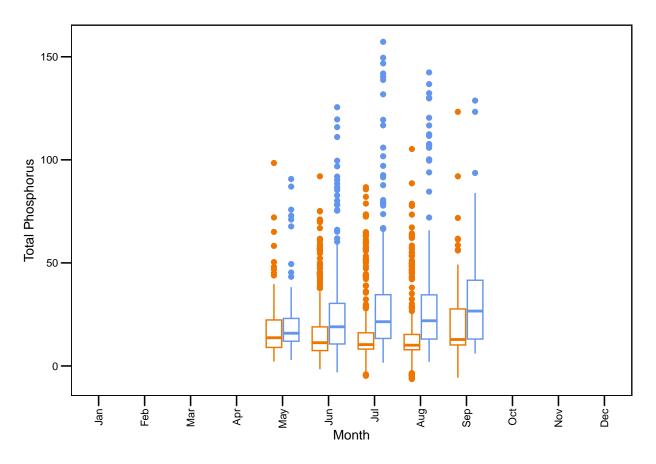
Warning: Removed 3566 rows containing non-finite values ('stat_boxplot()').

Temperature, Phophorus, & Nitrogen by Month

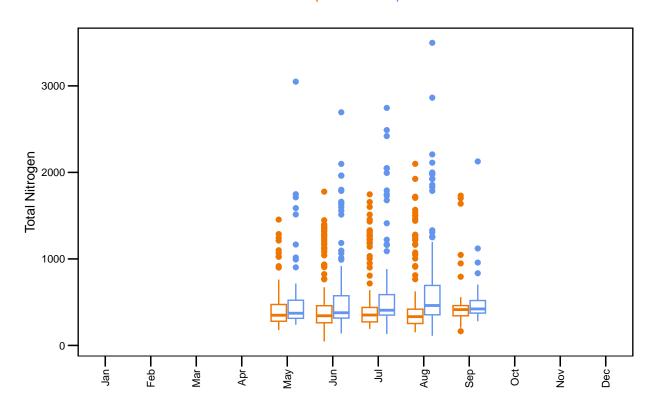


```
tp <- PeterPaul.chem.nutrients %>%
 ggplot(aes(
     x=month_f,
     y=tp_ug,
      color=lakename)
   ) +
 geom_boxplot() +
 ylab("Total Phosphorus") +
   xlab("Month") +
  scale_color_manual(values = c("darkorange2", "cornflowerblue"),
                     name = "Lake Name") +
 A05_theme1 +
 theme(legend.position = "NONE",
       axis.title.x = element_text(
     color = "black",
     size = 10)) +
 scale_x_discrete(drop=FALSE)
```

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



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

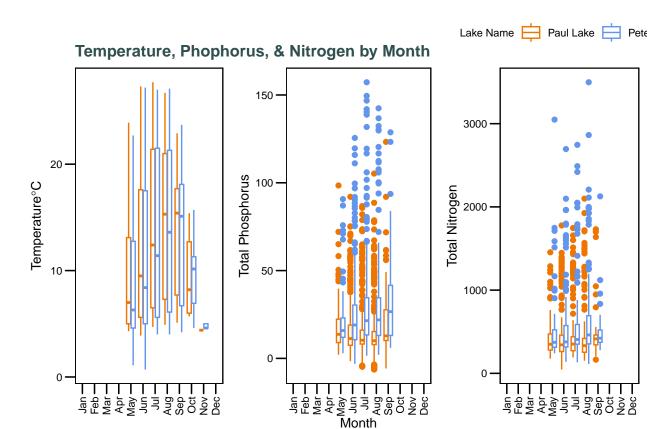


```
## Warning: Removed 3566 rows containing non-finite values ('stat_boxplot()').
```

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

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

print(plot5)



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

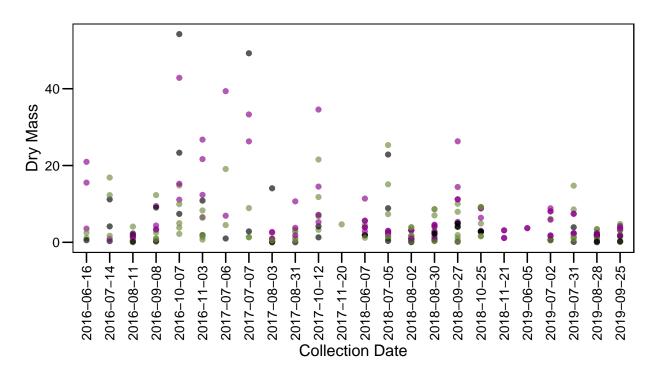
Answer: Temp is fairly the same between both lakes throughout the year.

- 6. [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)
- 7. [Niwot Ridge] Now, plot the same plot but with NLCD classes separated into three facets rather than separated by color.

```
theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))
print(plot6)
```

Needle Dry Mass by Collection Date

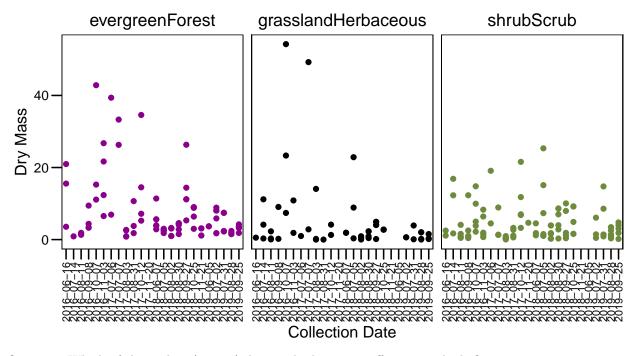
NLCD Class • evergreenForest • grasslandHerbaceous • shrubScrub



```
plot7 <- Niwot.Ridge.Litter %>%
  filter(functionalGroup == "Needles") %>%
  ggplot(aes(
      x=collectDate,
     y=dryMass,
      color=nlcdClass)) +
  geom_point() +
  ylab("Dry Mass") +
  xlab("Collection Date") +
  ggtitle("Needle Dry Mass by Collection Date") +
  facet_wrap(vars(nlcdClass)) +
  A05_theme +
  scale_color_manual(values = c("darkmagenta", "black", "darkolivegreen4"),
   name = "NLCD Class") +
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1, size = 8))
print(plot7)
```

Needle Dry Mass by Collection Date

NLCD Class • evergreenForest • grasslandHerbaceous • shrubScrub



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

Answer: