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

- Set up your session. Load the tidyverse, lubridate, here & cowplot packages, and verify your home directory. Upload 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) and the processed data file for the Niwot Ridge litter dataset (use the NEON_NIWO_Litter_mass_trap_Processed.csv version).
- 2. Make sure R is reading dates as date format; if not change the format to date.

```
library(tidyverse)
## — Attaching core tidyverse packages —
rse 2.0.0 --
## ✔ dplyr
             1.1.0
                        ✓ readr
                                    2.1.4
## ✓ forcats 1.0.0

✓ stringr

                                    1.5.0
## ✓ ggplot2 3.4.1

✓ tibble
                                    3.1.8
## ✔ lubridate 1.9.2

✓ tidyr

                                    1.3.0
## 🗸 purrr
              1.0.1
## — Conflicts ——
                                                      – tidyverse co
```

```
nflicts() —
## # dplyr::filter() masks stats::filter()
## # dplyr::lag()
                      masks stats::lag()
## i Use the |8;;http://conflicted.r-lib.org/conflicted package|8;; to
force all conflicts to become errors
library(lubridate)
library(here)
## here() starts at C:/Users/wwwla/Documents/EDA-Spring2023
library(cowplot)
##
## 载入程辑包: 'cowplot'
## The following object is masked from 'package:lubridate':
##
       stamp
getwd()
## [1] "C:/Users/wwwla/Documents/EDA-Spring2023"
lake.chem.nut <- read.csv("~/EDA-Spring2023/Data/Processed KEY/NTL-LTER</pre>
_Lake_Chemistry_Nutrients_PeterPaul_Processed.csv",stringsAsFactors = T
RUE)
litter.mass <- read.csv("~/EDA-Spring2023/Data/Processed/NEON_NIWO_Litt</pre>
er_mass_trap_Processed.csv",stringsAsFactors = TRUE)
#2
is.Date(lake.chem.nut$sampledate)
## [1] FALSE
is.Date(litter.mass$collectDate)
## [1] FALSE
lake.chem.nut$sampledate <- ymd(lake.chem.nut$sampledate)</pre>
litter.mass$collectDate <- ymd(litter.mass$collectDate)</pre>
is.Date(lake.chem.nut$sampledate)
## [1] TRUE
is.Date(litter.mass$collectDate)
## [1] 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

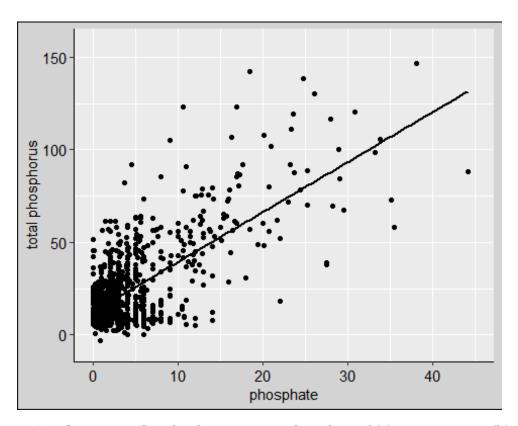
- Plot title
- Axis labels
- Axis ticks/gridlines
- Legend

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 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()).

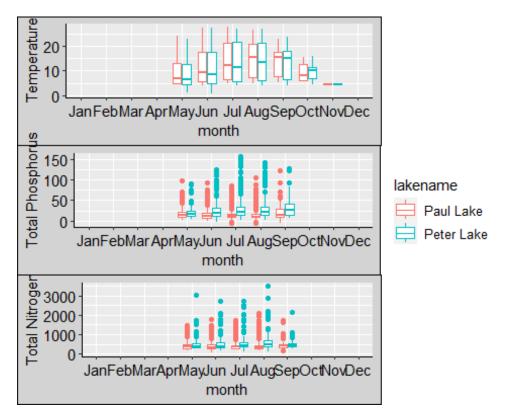
```
#4
ggplot(lake.chem.nut, aes(x = po4, y = tp_ug)) +
    geom_point() +
    xlim(0,45) +
    labs(x = "phosphate", y = "total phosphorus") +
    geom_smooth(method = lm, se = FALSE, colour = "black")
## `geom_smooth()` using formula = 'y ~ x'
## Warning: Removed 21947 rows containing non-finite values (`stat_smooth()`).
## Warning: Removed 21947 rows containing missing values (`geom_point()`).
```



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: R has a build in variable called month.abb that returns a list of months;see https://r-lang.com/month-abb-in-r-with-example

```
lake.TN <- ggplot(lake.chem.nut,</pre>
                 aes(x = factor(month, levels = 1:12, labels = month.ab
b),
                     y = tn_ug, color = lakename)) +
  scale_x_discrete(drop = FALSE) +
  geom_boxplot() +
  labs(x = "month", y = "Total Nitrogen") +
 theme(legend.position = "none")
legend <- get_legend(</pre>
  lake.T +
  guides(color = guide legend(nrow = 2)) +
 theme(legend.position = "right")
## Warning: Removed 3566 rows containing non-finite values (`stat_boxpl
ot()`).
plot1 <- plot_grid(lake.T + theme(legend.position = "none"),</pre>
          lake.TP + theme(legend.position = "none"),
          lake.TN + theme(legend.position = "none"),
          nrow = 3)
## Warning: Removed 3566 rows containing non-finite values (`stat_boxpl
ot()`).
## Warning: Removed 20729 rows containing non-finite values (`stat boxp
lot()`).
## Warning: Removed 21583 rows containing non-finite values (`stat_boxp
lot()`).
plot_grid(plot1, legend, nrow = 1, rel_widths = c(3, 1))
```

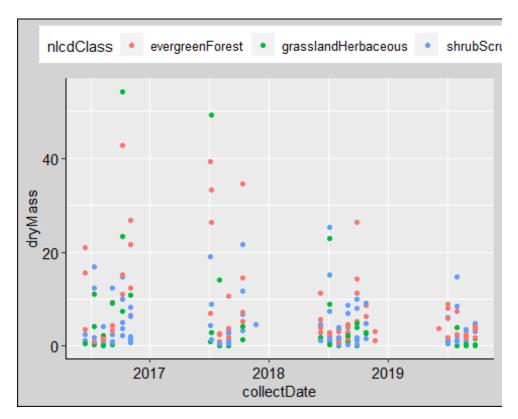


Question: What do you observe about the variables of interest over seasons and between lakes?

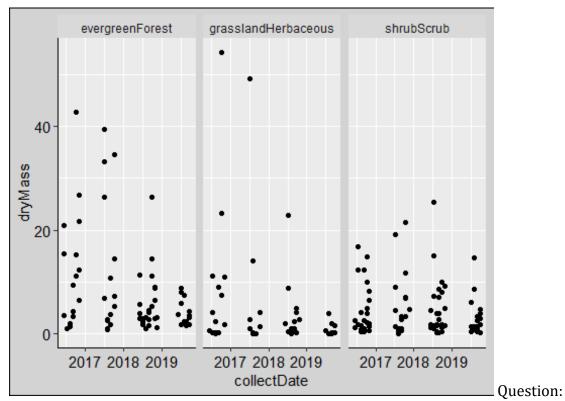
Answer: during the summer, the temperature of two lakes are similar, but TP and TN in Peter Lake are generally higher than them in Paul Lake.

- 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.

```
#6
litter.mass %>%
  filter(functionalGroup == "Needles") %>%
  ggplot(aes(x = collectDate, y = dryMass, color = nlcdClass)) +
  geom_point()
```



```
#7
litter.mass %>%
  filter(functionalGroup == "Needles") %>%
  ggplot(aes(x = collectDate, y = dryMass)) +
  geom_point() +
  facet_wrap(vars(nlcdClass))
```



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

Answer: the second one, because it shows the trend of drymass more clearly than the 6th, there are too many points appear in a single grapth that make me hard to catch informations.