

BrookHemphill_A05_DataVisualization.Rmd

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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.

```
install.packages('formatR')
```

```
## Installing package into '/home/guest/R/x86_64-pc-linux-gnu-library/4.3'  
## (as 'lib' is unspecified)
```

```
library(knitr)
opts_chunk$set(tidy.opts=list(width.cutoff=50), tidy=TRUE)

#1
getwd()
```

```
## [1] "/home/guest/EDA_Spring2024_New"
```

```
library(cowplot); library(tidyverse); library(lubridate); library(here)
```

```
## -- 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()
## x lubridate::stamp()   masks cowplot::stamp()
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors
## here() starts at /home/guest/EDA_Spring2024_New
```

```
here()
```

```
## [1] "/home/guest/EDA_Spring2024_New"
```

```
PeterPaul.chem.nutrients_processed <-
  read.csv(here("Data/Processed_KEY/NTL-LTER_Lake_Chemistry_Nutrients_PeterPaul_Processed.csv"),
           stringsAsFactors = T)
Niwot_Ridge_Litter_processed <-
  read.csv(here("Data/Processed_KEY/NEON_NIWO_Litter_mass_trap_Processed.csv"),
           stringsAsFactors = T)
```

```
#2
```

```
PeterPaul.chem.nutrients_processed$sampldate <- as.Date(PeterPaul.chem.nutrients_processed$sampldate)
#PeterPaul.chem.nutrients_processed$month <- as.Date(PeterPaul.chem.nutrients_processed$month)
Niwot_Ridge_Litter_processed$collectDate <- as.Date(Niwot_Ridge_Litter_processed$collectDate)
```

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
```

```
mytheme <- theme_minimal(base_size = 14) + theme(axis.text = element_text(color = "black",
  size = 10), legend.position = "top", legend.text = element_text(size = 12),
  legend.title = element_text(size = 12), plot.title = element_text(hjust = 0.5),
  axis.title.x = element_text(size = 10), axis.title.y = element_text(size = 10))

mytheme2 <- theme_minimal(base_size = 14) + theme(axis.text = element_text(color = "black",
  size = 10), legend.position = "top", plot.title = element_text(hjust = 0.5),
  axis.title.x = element_blank())
```

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
```

```
PvP04 <- ggplot(PeterPaul.chem.nutrients_processed,
  aes(x = po4, y = tp_ug, color = lakename, position = "dodge",
  shape = lakename)) + geom_point(alpha = 0.8) +
  xlim(0, 35) + geom_smooth(method = "lm", se = FALSE,
  size = 1) + scale_color_manual(values = c("darkblue",
  "lightblue")) + labs(title = "Total Phosphorus vs Phosphate\n",
  y = "Total Phosphorus (ug)", x = "Phosphate") +
  mytheme
```

```
## Warning: Using 'size' aesthetic for lines was deprecated in ggplot2 3.4.0.
## i Please use 'linewidth' instead.
## This warning is displayed once every 8 hours.
## Call 'lifecycle::last_lifecycle_warnings()' to see where this warning was
## generated.
```

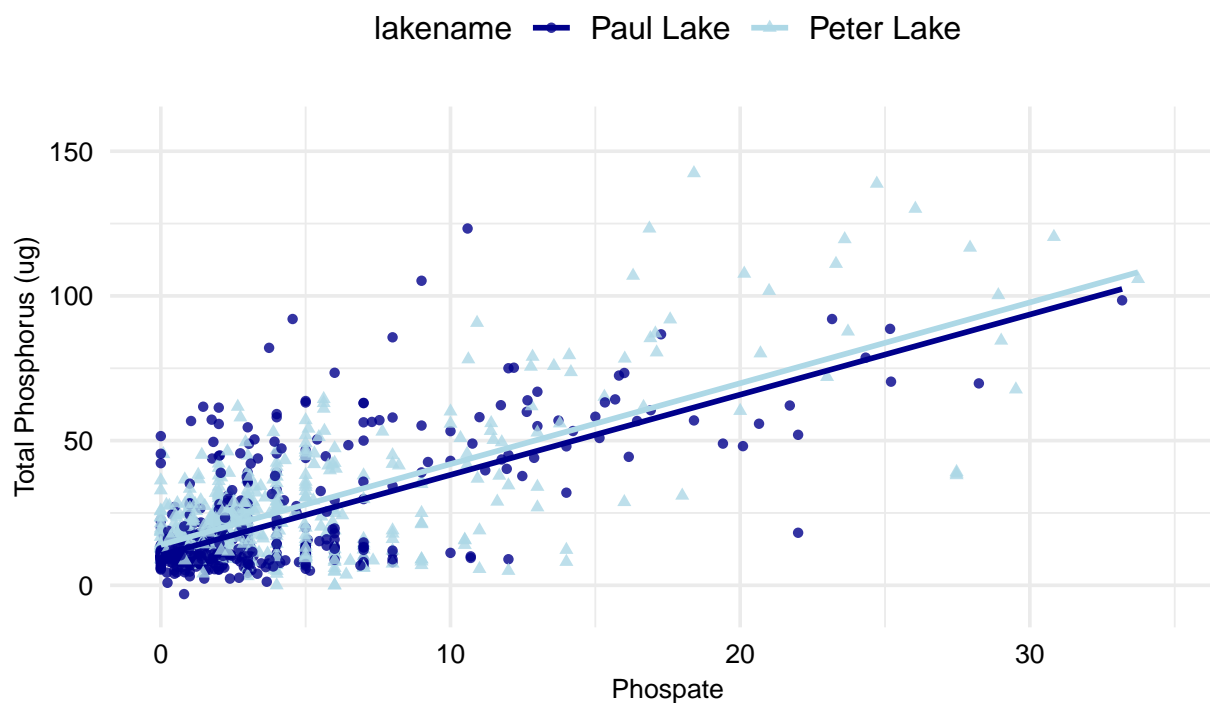
```
print(PvP04)
```

```
## 'geom_smooth()' using formula = 'y ~ x'
```

```
## Warning: Removed 21951 rows containing non-finite values ('stat_smooth()').
```

```
## Warning: Removed 21951 rows containing missing values ('geom_point()').
```

Total Phosphorus vs Phosphate



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 months in integers to factors
PeterPaul.chem.nutrients_processed$month <- factor(PeterPaul.chem.nutrients_processed$month,
  levels = 1:12, labels = month.abb)

NLT_LTER_temperature <- ggplot(PeterPaul.chem.nutrients_processed,
  aes(x = month, y = temperature_C)) + geom_boxplot(aes(color = lakename),
  alpha = 0.8) + scale_color_manual(values = c("darkblue",
  "lightblue")) + labs(title = "Temperature each Month\n",
  y = "Temperature (C)", x = "Month", color = "Lake Name") +
  mytheme2

NLT_LTER_TP <- ggplot(PeterPaul.chem.nutrients_processed,
  aes(x = month, y = tp_ug)) + geom_boxplot(aes(color = lakename),
  alpha = 0.8, show.legend = FALSE) + scale_color_manual(values = c("darkblue",
  "lightblue")) + labs(title = "Total Phosphorus each Month\n",
  y = "Total Phosphorus", x = "Month", color = "Lake Name") +
  theme(legend.position = "none", axis.title.x = element_blank()) +
  mytheme

NLT_LTER_TN <- ggplot(PeterPaul.chem.nutrients_processed,
  aes(x = month, y = tn_ug)) + geom_boxplot(aes(color = lakename),
  alpha = 0.8, show.legend = FALSE) + scale_color_manual(values = c("darkblue",
  "lightblue")) + labs(title = "Total Nitrogen each Month\n",
  y = "Total Nitrogen (ug)", x = "Month", color = "Lake Name") +
  mytheme2

print(NLT_LTER_temperature)

```

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

Temperature each Month

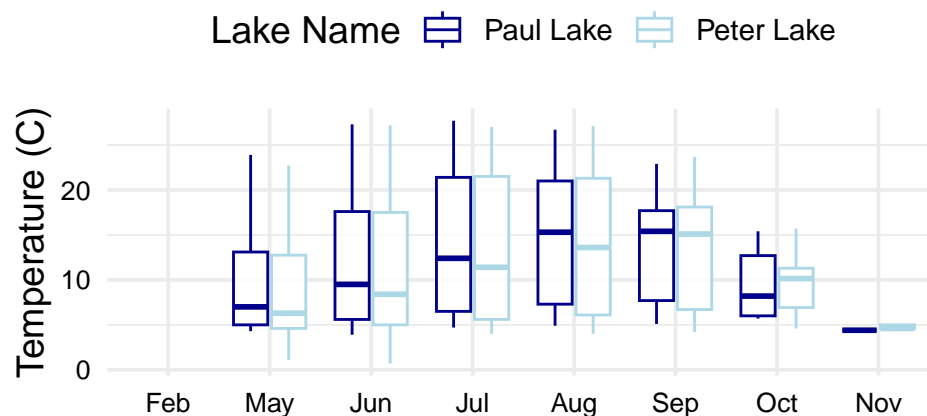


Figure 1: Figure 1

```
print(NTL_LTER_TP)
```

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

Total Phosphorus each Month

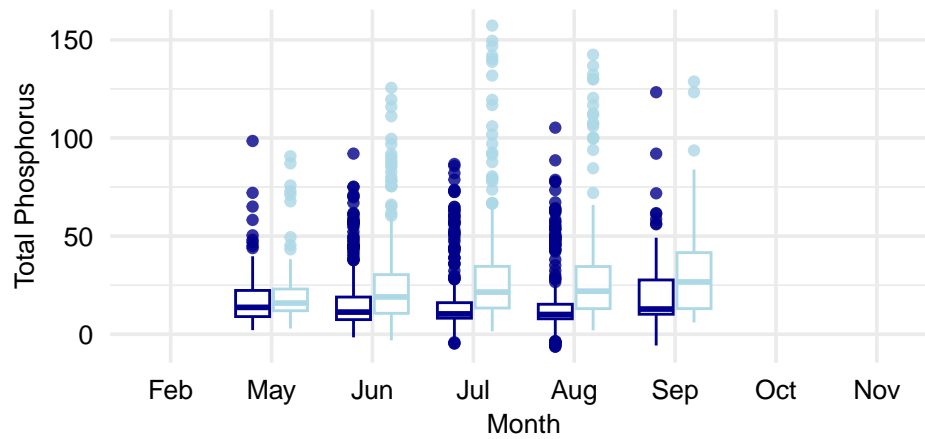


Figure 2: Figure 1

```
print(NTL_LTER_TN)
```

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

Total Nitrogen each Month

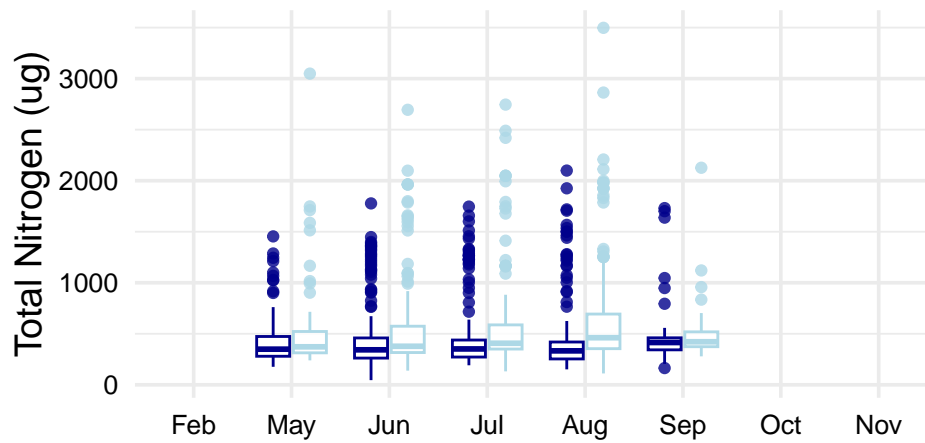


Figure 3: Figure 1

```
plot_grid(NTL_LTER_temperature, NTL_LTER_TN, NTL_LTER_TP,
  nrow = 3, align = "h", rel_heights = c(1, 1, 1))
```

```
## Warning: Removed 3566 rows containing non-finite values ('stat_boxplot()').
## Warning: Removed 21583 rows containing non-finite values ('stat_boxplot()').
## Warning: Removed 20729 rows containing non-finite values ('stat_boxplot()').
## Warning: Graphs cannot be horizontally aligned unless the axis parameter is
## set. Placing graphs unaligned.
```

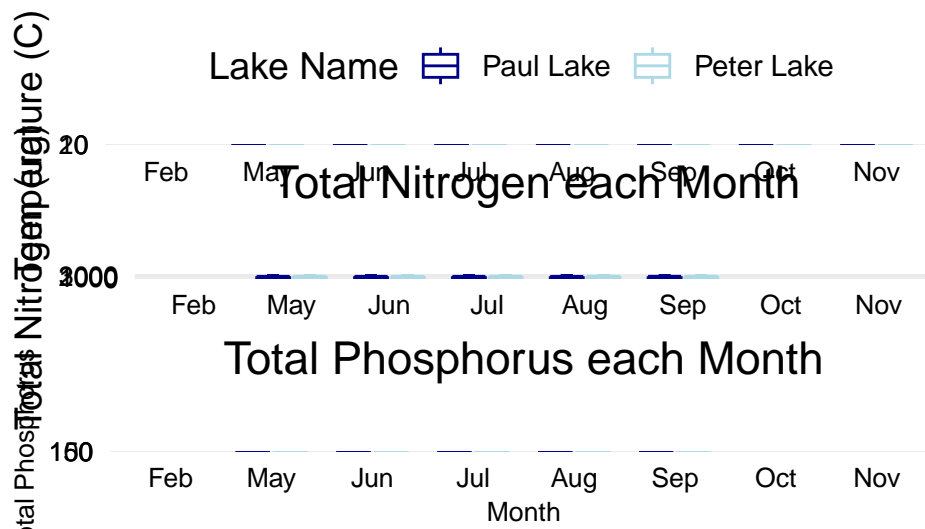


Figure 4: Figure 1

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

Answer: Over seasons and between lakes, there is generally more variability in the total phosphorus and nitrogen by month in Peter Lake than in Paul lake. As temperature rises, total P and nitrogen also slightly increase and then decrease as the temperature decreases, however, that trend is slightly harder to see without data from October, November, and December.

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
needles_subset <- Niwot_Ridge_Litter_processed %>%
  filter(functionalGroup == "Needles")
```

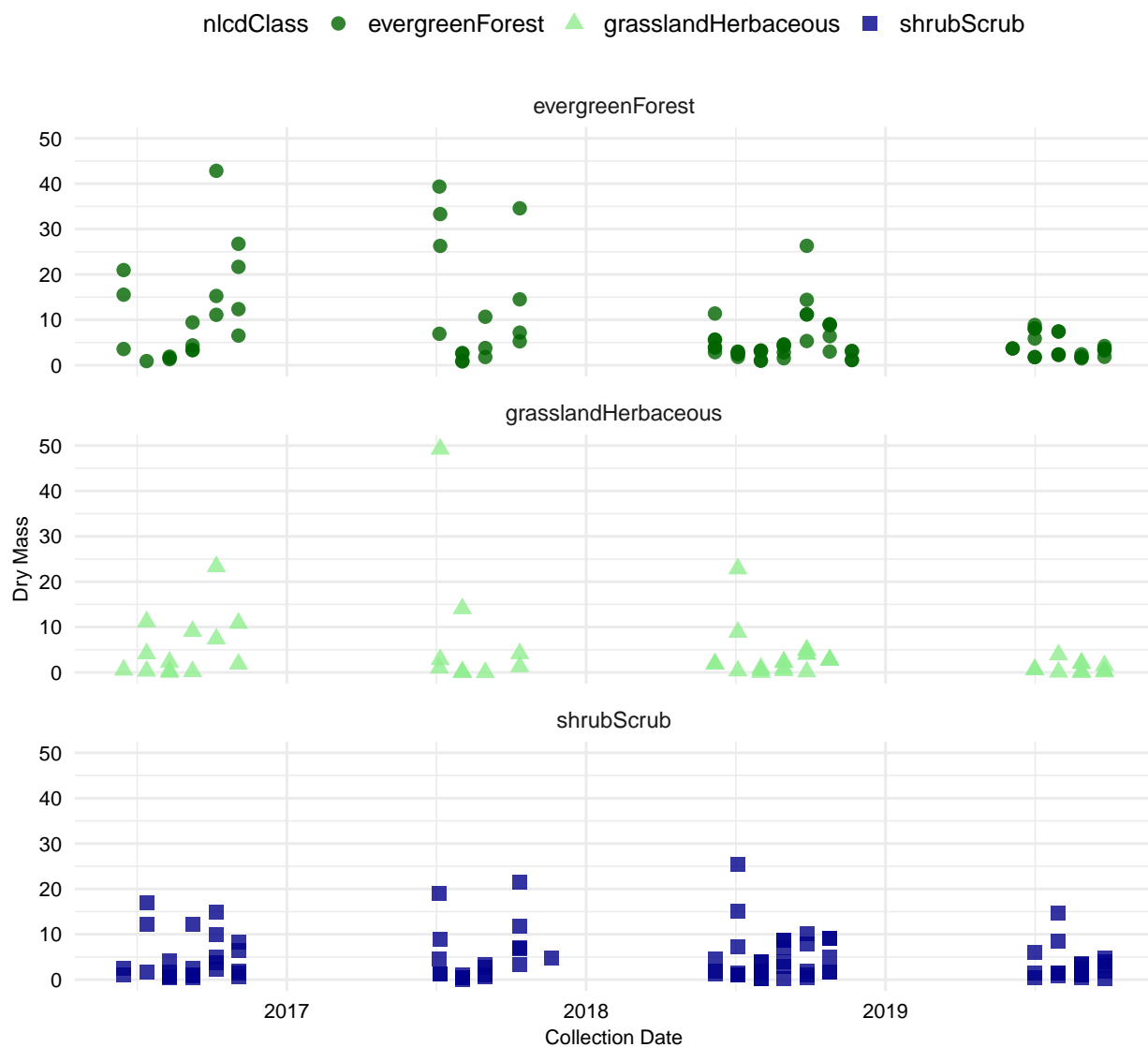


```
# 7
needles_plot2 <- ggplot(needles_subset, aes(x = collectDate,
  y = dryMass, color = nlcdClass, shape = nlcdClass)) +
  geom_point(alpha = 0.8, size = 3) + scale_color_manual(values = c("darkgreen",
    "lightgreen", "darkblue")) + labs(title = "Needle Litter Dry Mass by Year\n",
  y = "Dry Mass", x = "Collection Date", color = "nlcdClass") +
  ylim(0, 50) + facet_wrap(vars(nlcdClass), nrow = 3) +
  mytheme

print(needles_plot2)
```

Warning: Removed 1 rows containing missing values ('geom_point()').

Needle Litter Dry Mass by Year



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

Answer: I think it depends on what you are looking for, however, for general purposes 6 is more visually appealing, and the information is condensed/comparable, which makes observing trends easier.