

# Assignment 5: Data Visualization

Student Name

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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.

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## 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 Project setup
```

```
##Load packages
```

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

```
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr      1.1.4      v readr      2.1.5
## v forcats    1.0.0      v stringr   1.5.1
## v ggplot2     3.5.1      v tibble    3.2.1
## v lubridate  1.9.3      v tidyr     1.3.1
## 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
## here() starts at /home/guest/EDE
##
##
## Attaching package: 'cowplot'
##
##
## The following object is masked from 'package:lubridate':
##
##     stamp
```

```
## Check working directory
here()
```

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

```
##Load datasets
NTL_tidy <- read.csv(
  here('Data/Processed_KEY/NTL-LTER_Lake_Chemistry_Nutrients_PeterPaul_Processed.csv'),
  stringsAsFactors = T)
Trap_data <- read.csv(
  here('Data/Processed_KEY/NEON_NIWO_Litter_mass_trap_Processed.csv'),
  stringsAsFactors = T)

#2 Check/convert date columns
NTL_tidy$sampldate <- ymd(NTL_tidy$sampldate)
Trap_data$collectDate <- ymd(Trap_data$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 Create my theme
my_theme = theme_bw() +
  theme(
    plot.background = element_rect(
      fill = 'Light gray',
      color = 'Black'
    ),
    axis.line = element_line(
      linewidth = 1,
      color='Blue'
    ),
    axis.text = element_text(
      family='serif'
    ),
```

```

    plot.title = element_text(
      face = 'bold',
      color='Blue',
      family='serif'
    )
  )
)

#Set my theme to the default
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_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()`).

```

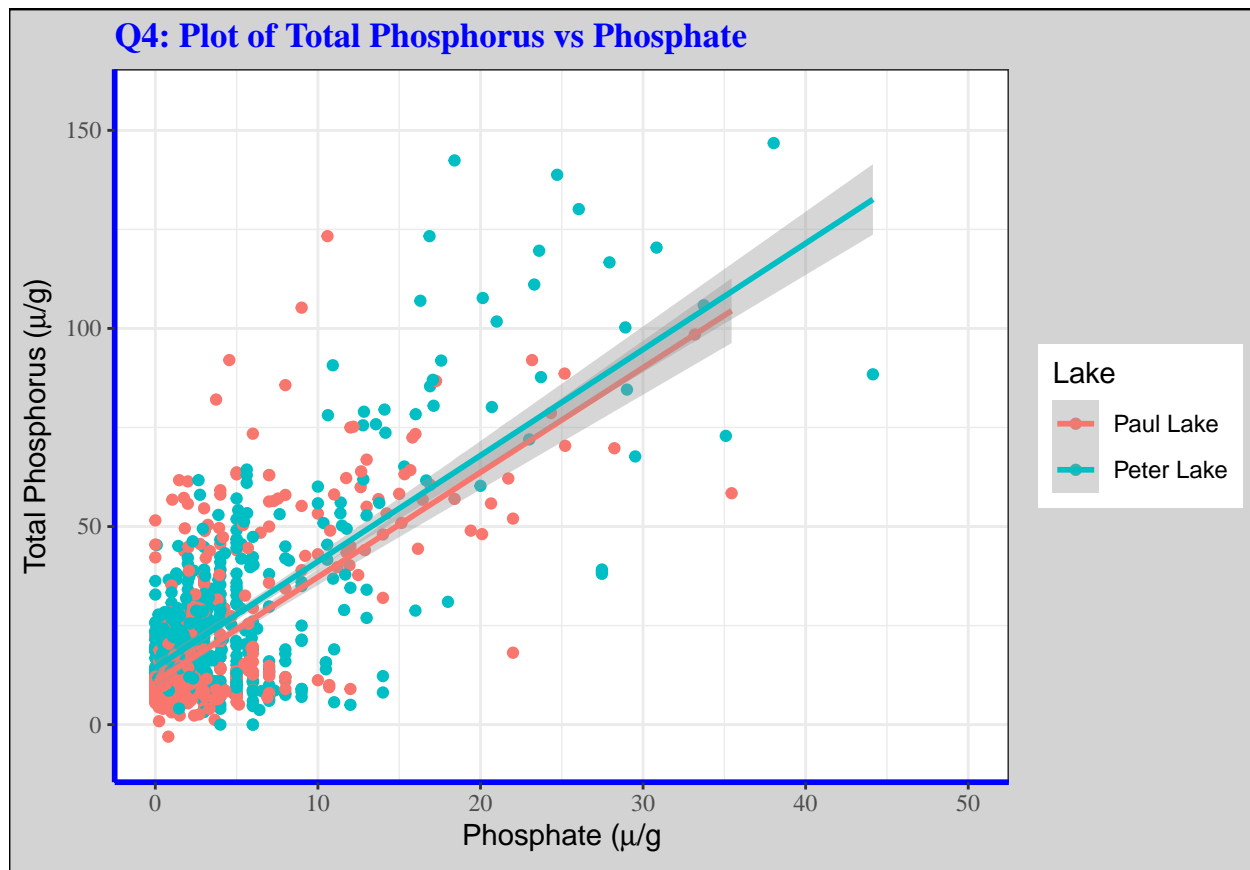
#4 Plot total P vs PO4
ggplot(NTL_tidy, aes(x=po4, y=tp_ug, color=lakename)) +
  geom_point() +
  geom_smooth(method='lm') +
  xlim(0, 50) +
  labs(
    x = expression(paste('Phosphate (', mu, '/g)'),
    y = expression(paste('Total Phosphorus (', mu, '/g)'),
    title = 'Q4: Plot of Total Phosphorus vs Phosphate',
    color='Lake'
  )

```

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

```
## Warning: Removed 21947 rows containing non-finite outside the scale range
## ('stat_smooth()').
```

```
## Warning: Removed 21947 rows containing missing values or values outside the scale range
## ('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>

```
#5 Boxlots
base_plt <- NTL_tidy %>%
  mutate(the_month = factor(month, levels = 1:12, labels = month.abb)) %>%
  ggplot(aes(x=the_month, color=lakename)) #+
  scale_x_discrete(labels=month.abb)

## <ggproto object: Class ScaleDiscretePosition, ScaleDiscrete, Scale, gg>
##   aesthetics: x xmin xmax xend
##   axis_order: function
##   break_info: function
##   break_positions: function
##   breaks: waiver
##   call: call
##   clone: function
##   dimension: function
##   drop: TRUE
##   expand: waiver
##   get_breaks: function
```

```
##      get_breaks_minor: function
##      get_labels: function
##      get_limits: function
##      get_transformation: function
##      guide: waiver
##      is_discrete: function
##      is_empty: function
##      labels: Jan Feb Mar Apr May Jun Jul Aug Sep Oct Nov Dec
##      limits: NULL
##      make_sec_title: function
##      make_title: function
##      map: function
##      map_df: function
##      n.breaks.cache: NULL
##      na.translate: TRUE
##      na.value: NA
##      name: waiver
##      palette: function
##      palette.cache: NULL
##      position: bottom
##      range: environment
##      range_c: environment
##      rescale: function
##      reset: function
##      train: function
##      train_df: function
##      transform: function
##      transform_df: function
##      super: <ggproto object: Class ScaleDiscretePosition, ScaleDiscrete, Scale, gg>
```

```
plt1 <- base_plt+
  geom_boxplot(aes(y=temperature_C),show.legend = T) +
  labs(
    x='',
    y='Temperature (C)',
    title='Temperature'
  ) +
  theme(
    legend.position = "top"
  )
```

```
plt2 <- base_plt+
  geom_boxplot(aes(y=tp_ug),show.legend = F) +
  labs(
    x='',
    y='Total P',
    title='Total P'
  )
```

```
plt3 <- base_plt+
  geom_boxplot(aes(y=tn_ug),show.legend = F) +
  labs(
    x='Month',
    y='Total N',
```

```

    title='Total N',
    color='Lake'
  )

plot_grid(plt1,plt2,plt3,
          nrow = 3,
          align = 'h',
          rel_heights = c(1.3,1,1))

```

```

## Warning: Removed 3566 rows containing non-finite outside the scale range
## ('stat_boxplot()').

```

```

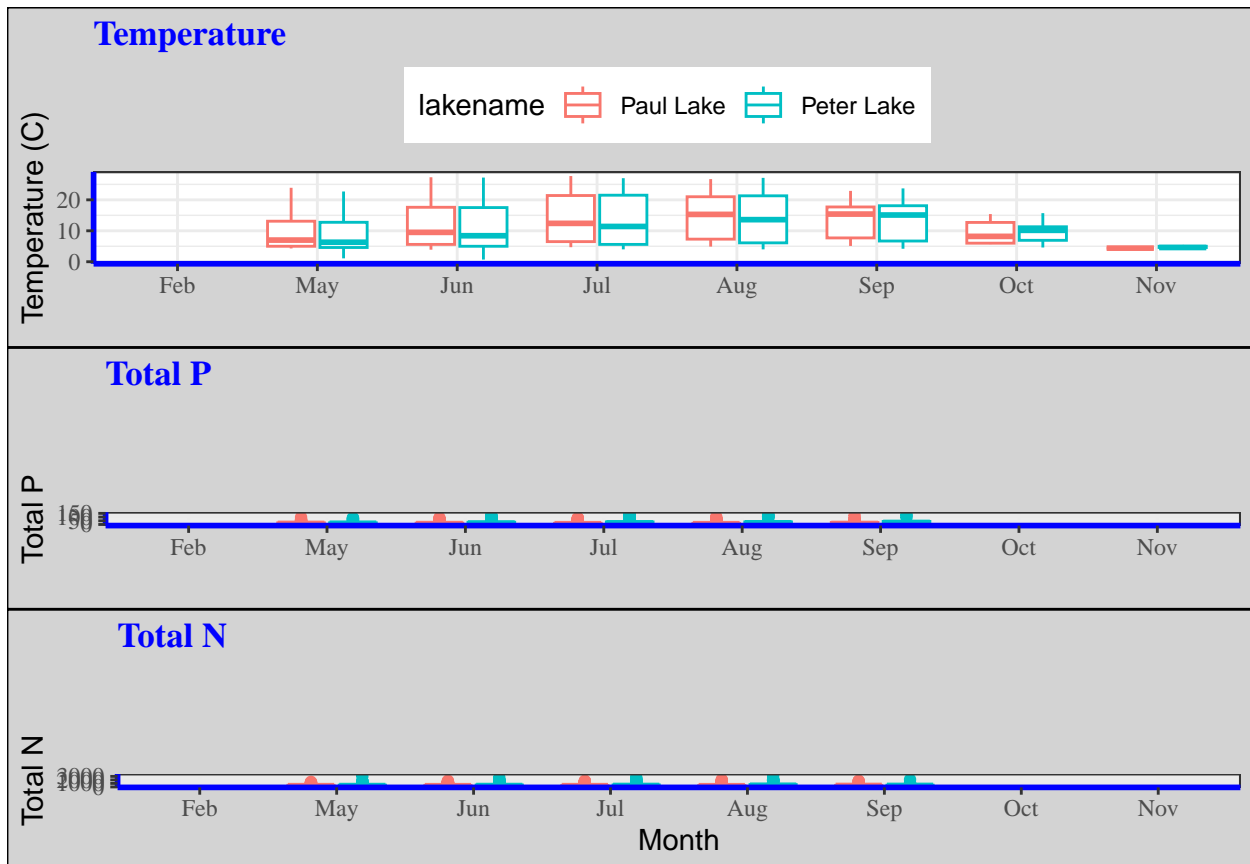
## Warning: Removed 20729 rows containing non-finite outside the scale range
## ('stat_boxplot()').

```

```

## Warning: Removed 21583 rows containing non-finite outside the scale range
## ('stat_boxplot()').

```



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

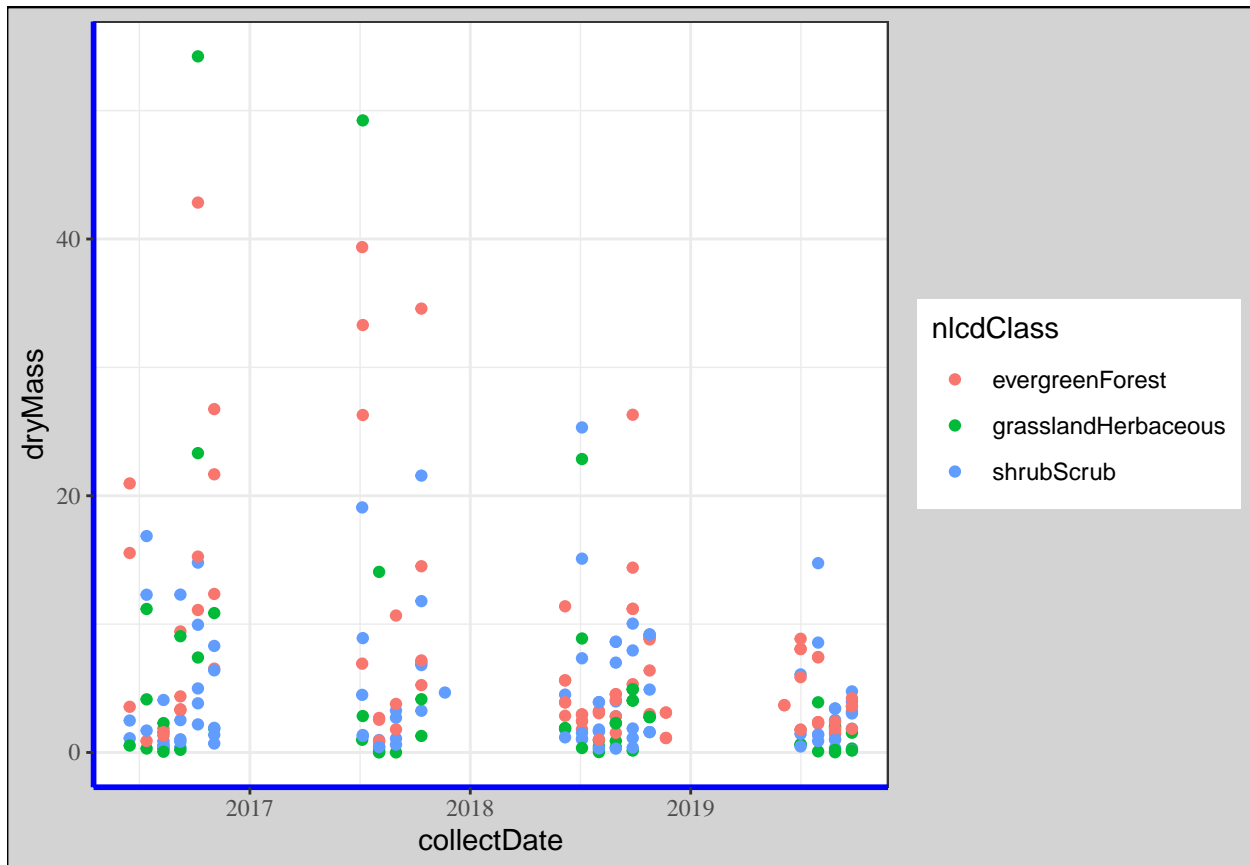
Answer:

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.

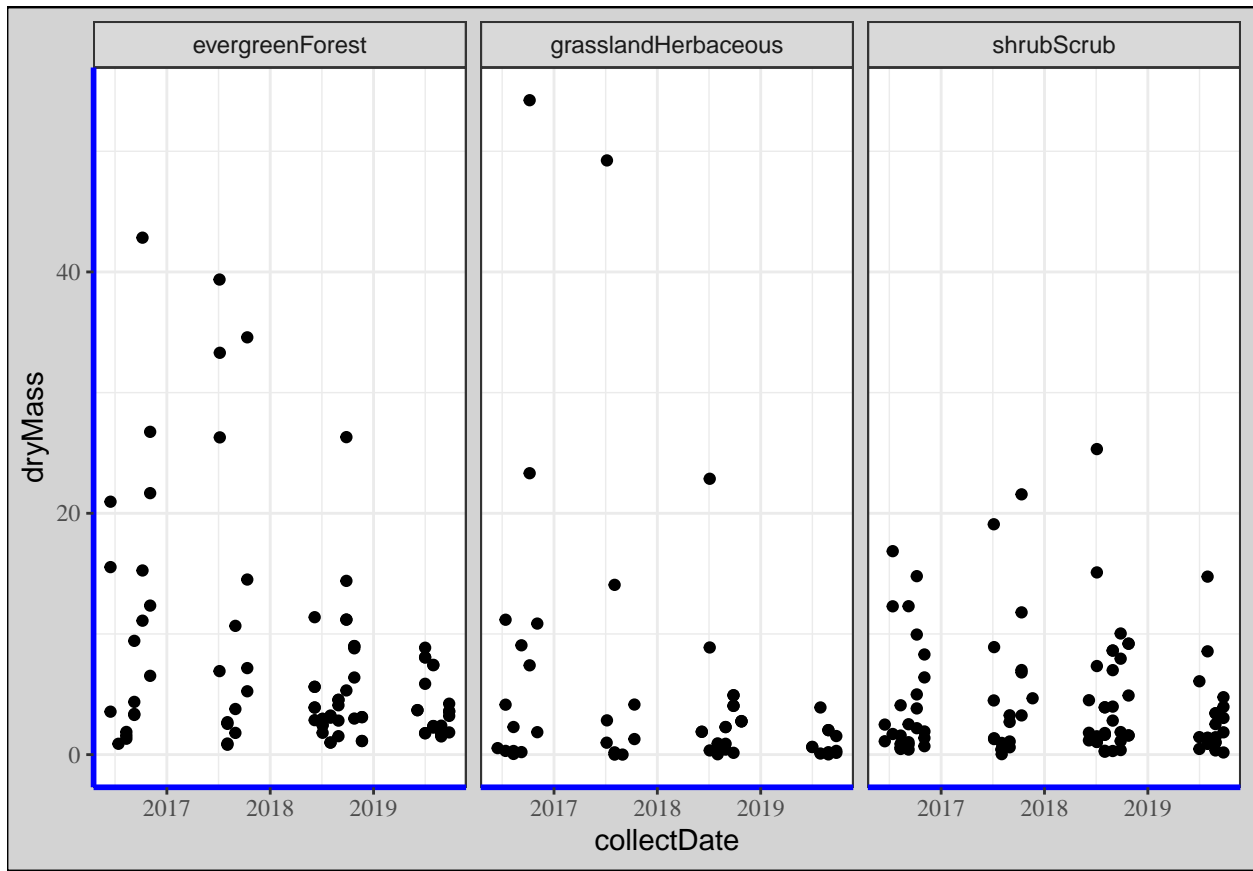
#6

```
Trap_data %>%  
  filter(functionalGroup == 'Needles') %>%  
  ggplot(aes(x=collectDate,y=dryMass,color=nlcdClass)) +  
  geom_point()
```



#7

```
Trap_data %>%  
  filter(functionalGroup == 'Needles') %>%  
  ggplot(aes(x=collectDate,y=dryMass)) +  
  geom_point() +  
  facet_wrap(vars(nlcdClass))
```



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

Answer: