

We shall be using one built-in dataset called the **air quality** dataset, which pertains to the daily air quality measurements in New York from May to September 1973.

This dataset consists of more than 100 observations on **6 variables**, i.e., **Ozone**(mean parts per billion), **Solar.R**(Solar Radiation), **Wind**(Average wind speed), **Temp**(maximum daily temperature in Fahrenheit), **Month**(month of observation) and **Day**(Day of the month).

Details:

Daily readings of the following air quality values for May 1, 1973 (a Tuesday) to September 30, 1973.

Ozone: Mean ozone in parts per billion from 1300 to 1500 hours at Roosevelt Island

Solar.R: Solar radiation in Langleys in the frequency band 4000--7700 Angstroms from 0800 to 1200 hours at Central Park

Wind: Average wind speed in miles per hour at 0700 and 1000 hours at LaGuardia Airport

Temp: Maximum daily temperature in degrees Fahrenheit at La Guardia Airport.

- data(airquality)
- We learnt head(airquality) shows first 6 observations.
- You can also use head(airquality,n) and tail(airquality,n)
 The head outputs the top n elements in the dataset while the tail method outputs the bottom n.

- #Example
- # suppose we need to print first 20 outputs from airquality data
- data(airquality)
- n = 20
- # for the first 20 observations
- head(airquality,n)
- # for the last 20 observations
- tail(airquality,n)

- summary(airquality)
- #save ozone data
- data.ozone = airquality\$Ozone
- #save temp data
- data.temp = airquality\$Temp
- #save wind data
- data.wind = airquality\$Wind

- # if there is any missing value
- is.na(data.ozone)
- is.na(data.temp)
- is.na(data.wind)
- # If there is TRUE in the output, it means there is missing value. So there is missing value in ozone data.
- # Now we remove the missing value from the ozone data
- data.ozone = data.ozone[!is.na(data.ozone)]
- data.ozone



We start with loading the required packages.

Data visualization



Package ggplot2 is included in the tidyverse package.



Package ggplot2 is a plotting package that makes it simple to create complex plots from data in a data frame.

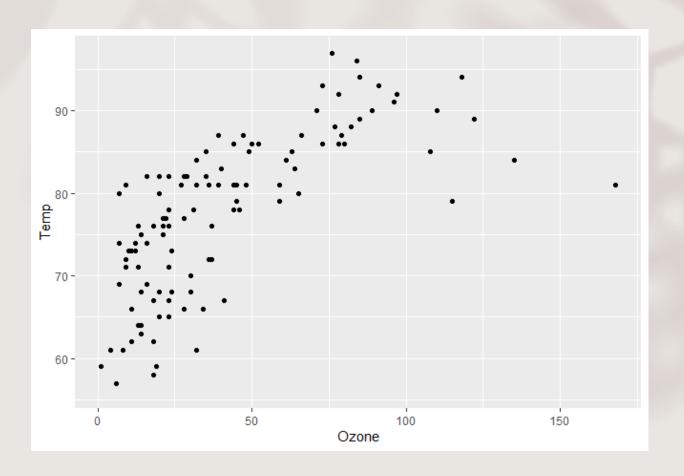
- To build a ggplot, we will use the following basic template that can be used for different types of plots:
- #ggplot(data = <DATA>, mapping = aes(<MAPPINGS>)) + <GEOM_FUNCTION>()
- #use the ggplot() function and bind the plot to a specific data frame using the data argument
- ggplot(data = airquality)

• Define a mapping using aes() function, by selecting variables and specifying how to present them in the graph.

• Add **'geoms'** - this function decides the graphical representations of the data in the plot (points, lines, bars).

- To add a geom to the plot use '+' operator.
- Let's use geom_point() first to create a scatter plot.

- ggplot(data = airquality, mapping = aes(x = Ozone, y = Temp))+geom_point()
- Temperature vs. Mean Ozone parts scatter plot looks like →



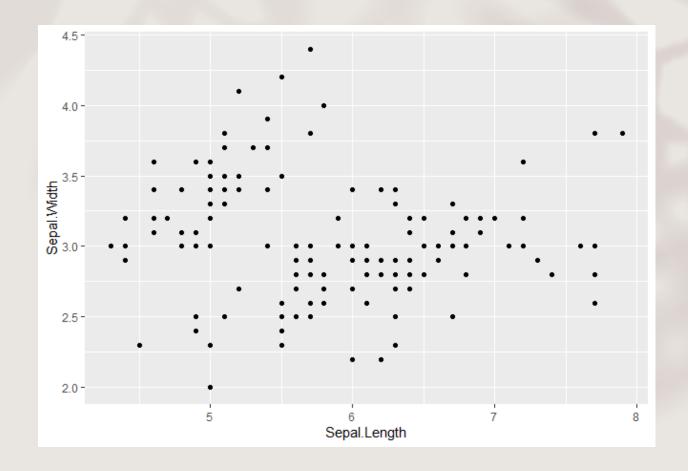
- What do we conclude by a quick look at the plot?
- Average ozone parts and temperature correlated.

- # Let us use the Iris data.
- # The **iris** dataset is provided natively by R, we call it a built-in dataset.
- # So, we do not need to use a package to get this data.

head(iris)

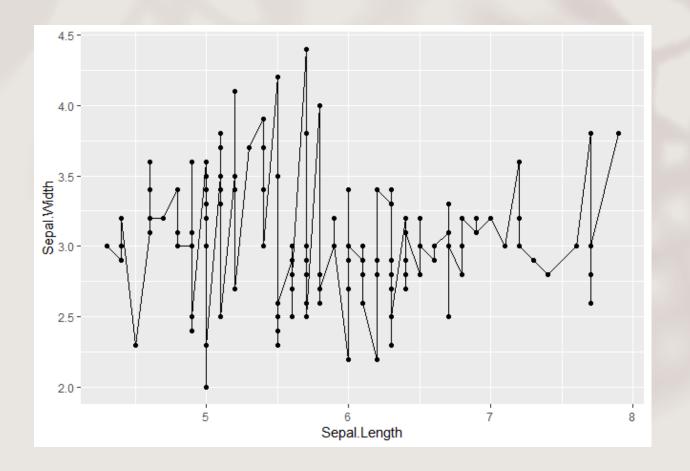
- # basic scatterplot
- ggplot(iris, aes(x=Sepal.Length, y=Sepal.Width)) +
- geom_point()

 Sepal.width vs. Sepal.Length scatter plot looks like →

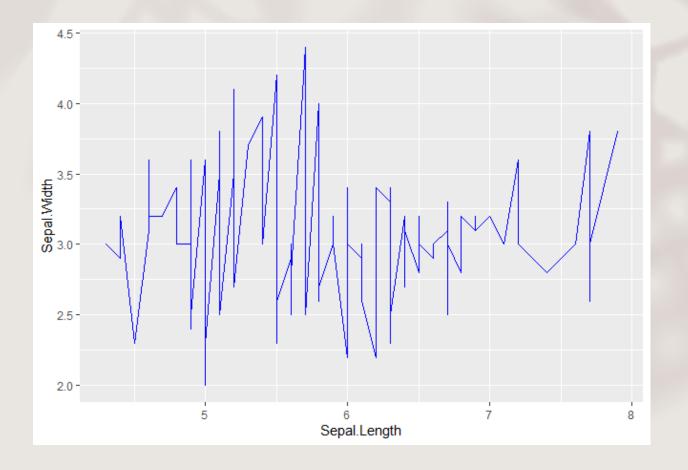


- #To create a connected scatter plot
- ggplot(iris, aes(x=Sepal.Length, y=Sepal.Width)) +
- geom_point()+geom_line()

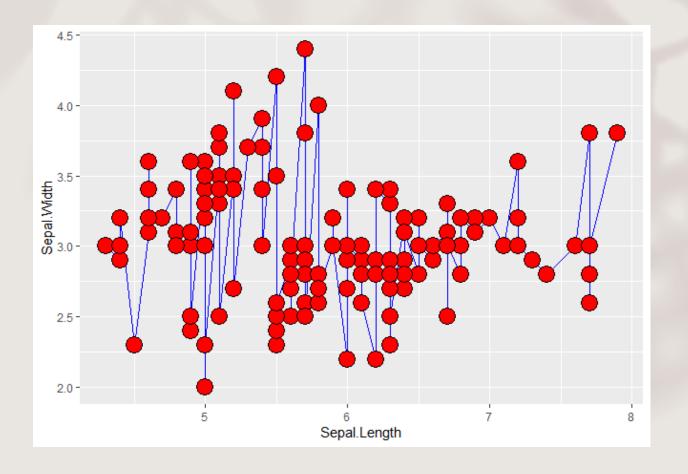
 Sepal.width vs. Sepal.Length connected scatter plot looks like →



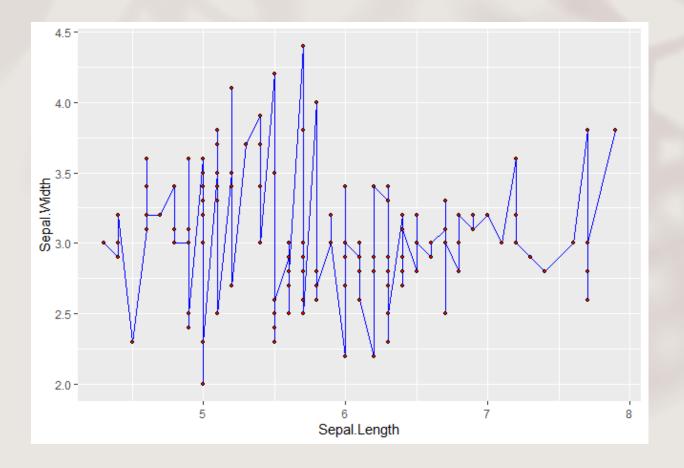
- Customize the connected scatterplot
- You can add a color:
- ggplot(iris, aes(x=Sepal.Length, y=Sepal.Width)) + geom_line(color="blue")



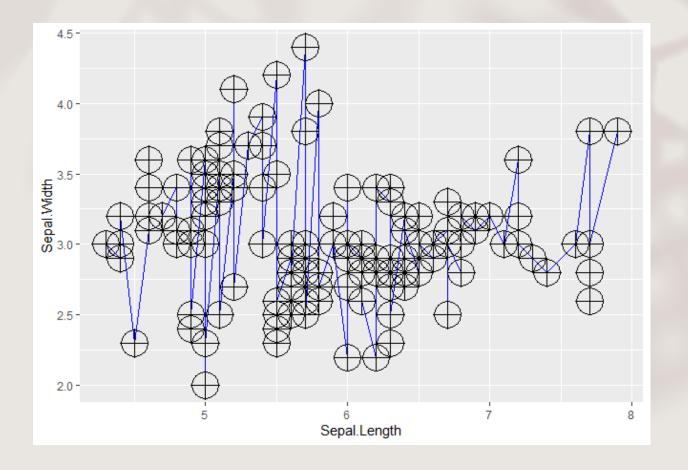
ggplot(iris, aes(x=Sepal.Length, y=Sepal.Width)) + geom_line(color="blue") + geom_point(shape=21, color="black", fill="red", size=6)



- Change the size of the points
- ggplot(iris, aes(x=Sepal.Length, y=Sepal.Width)) + geom_line(color="blue") + geom_point(shape=21, color="black", fill="red", size=1)

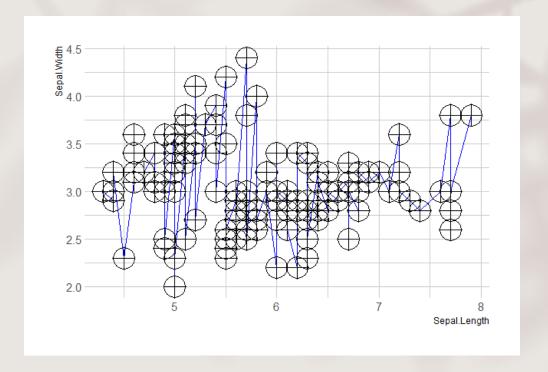


- Change the size of the points
- ggplot(iris, aes(x=Sepal.Length, y=Sepal.Width)) + geom_line(color="blue") + geom_point(shape=21, color="black", fill="red", size=1)



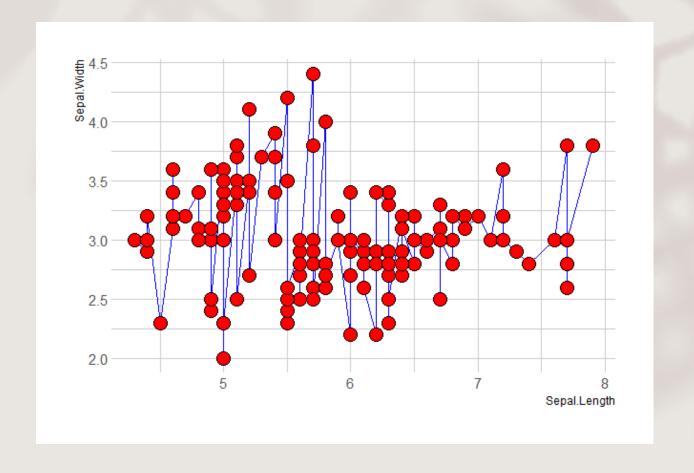
- Custom the general theme with the theme_ipsum() function of the hrbrthemes package.
- Add a title with ggtitle().
- Custom circle and line with arguments like shape, size, color and more.

- # Libraries
- library(tidyverse)
- library(hrbrthemes)
- ggplot(iris, aes(x=Sepal.Length, y=Sepal.Width)) +
 geom_line(color="blue") +
 geom_point(shape=10, color="black", fill="red", size=10) +
 theme_ipsum()



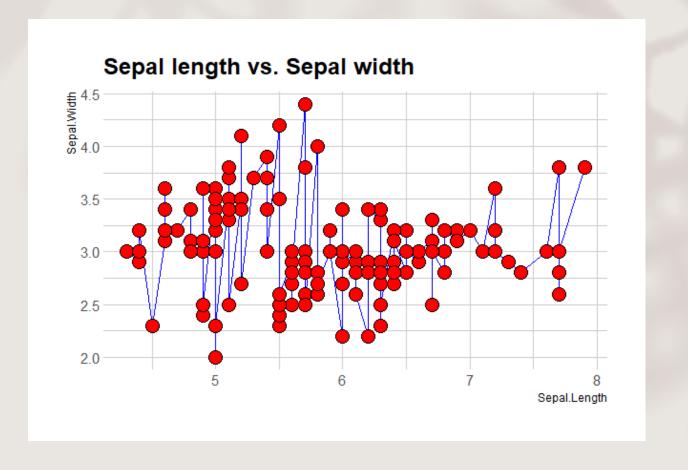
- Change size and shape with the general theme as the background:
- ggplot(iris, aes(x=Sepal.Length, y=Sepal.Width)) +

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geom_line( color="blue") +
  geom_point(shape=21, color="black",
fill="red", size=5) +
  theme_ipsum()
```



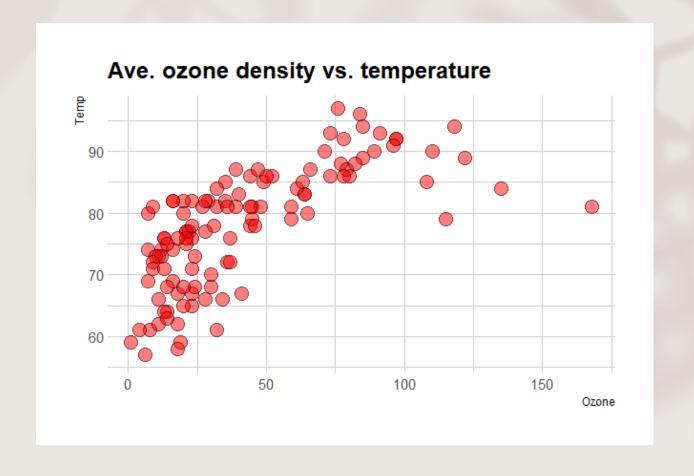
- Add title with the ggtitle() function
- ggplot(iris, aes(x=Sepal.Length, y=Sepal.Width)) +

```
geom_line( color="blue") +
geom_point(shape=21, color="black",
fill="red", size=5) +
theme_ipsum() +
ggtitle("Sepal length vs. Sepal width")
```



- ggplot(data = airquality, mapping = aes(x = Ozone, y = Temp))+
- geom_point(alpha=0.5,shape=21, color="black", fill="red", size=5) +

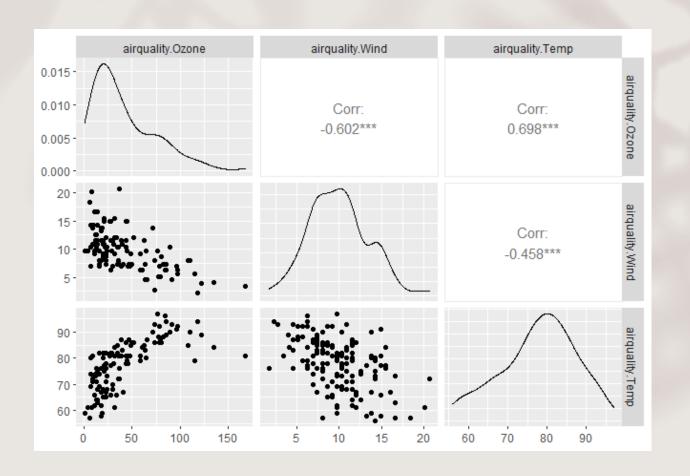
theme_ipsum() + ggtitle("Ave. ozone density vs. temperature")



• ## package GGally is has R functions for great correlation plots

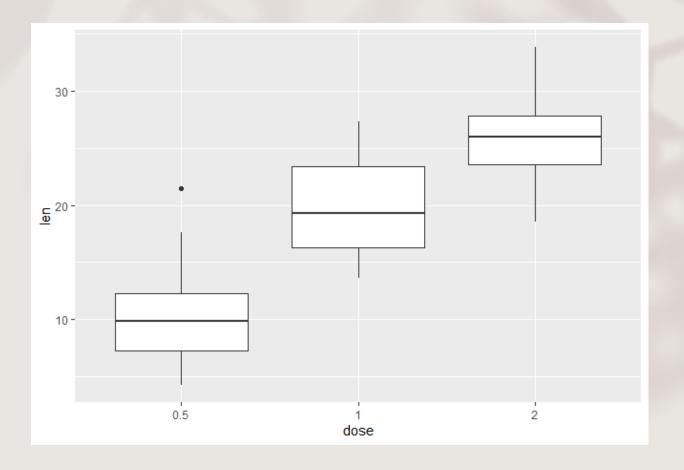
- library(GGally)
- data = data.frame(airquality\$Ozone,airquality\$Wind, airquality\$Temp)
- ggpairs(data)

- Scatterplots of each pair of numeric variable are drawn on the left part of the figure.
- Pearson correlation is displayed on the right.
- Variable distribution is available on the diagonal.



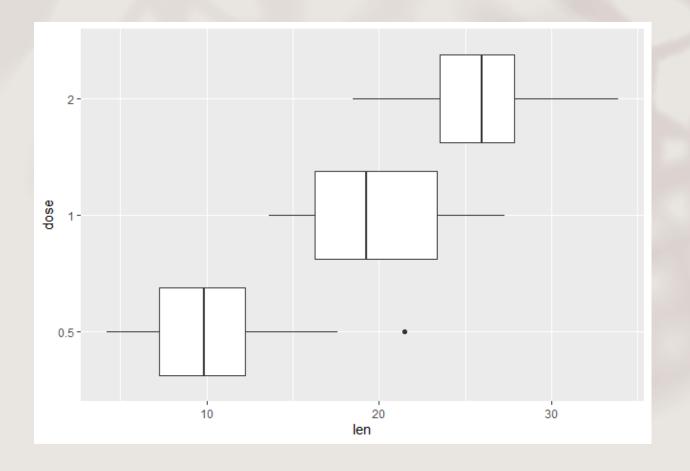
- To create a boxplot we use a dataset called ToothGrowth.
- Convert the variable dose from a numeric to a factor variable ToothGrowth\$dose <- as.factor(ToothGrowth\$dose)
 head(ToothGrowth)

ggplot(ToothGrowth, aes(x=dose, y=len)) + geom_boxplot()



- # Rotate the box plot
- ggplot(ToothGrowth, aes(x=dose, y=len)) +

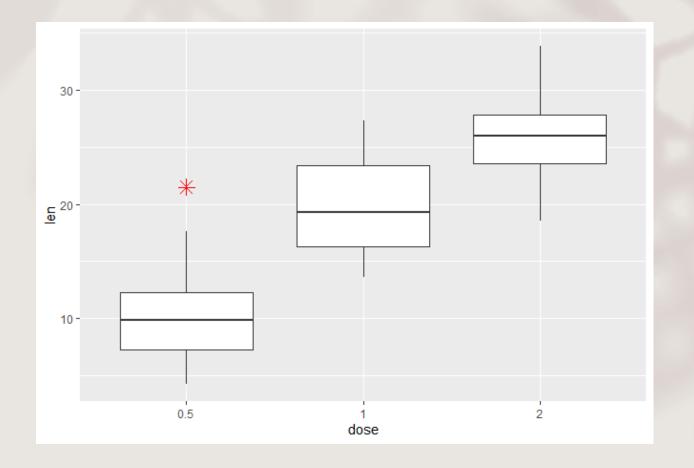
geom_boxplot() + coord_flip()



- # Notched box plot
- ggplot(ToothGrowth, aes(x=dose, y=len)) +
- geom_boxplot(notch=TRUE)



- # Change outlier, color, shape and size
- ggplot(ToothGrowth, aes(x=dose, y=len)) +
- geom_boxplot(outlier.colour="red", outlier.shape=8,
- outlier.size=4)



- # Change outlier, color, shape and size
- ggplot(ToothGrowth, aes(x=dose, y=len)) + geom_boxplot(outlier.colour="red", outlier.shape=11, outlier.size=9)

