



Base Graphics #1

(Functions and Parameters)

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Graphics Packages

There are three main graphics packages for R:

- `graphics` (base graphics)
 - Don't need to load
- `lattice` (grid graphics)
 - Need to load `library(lattice)`
- `ggplot2` (grid graphics)
 - Need to install and load

Graphics Demos

demo (graphics)

Graphics Commands

Plotting commands in R are separated into three basic groups:

- **High-level commands:** plotting functions that produce a new graphical device with a plot. High-level commands generally produce a full graphic with axes, labels, and titles
- **Low-level commands:** plotting functions that add elements to graphics such as lines, points, legends
- **Interactive commands:** graphical functions that allow you to interact with the graphic (e.g., using the mouse)

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The `plot()` Function

plot()

The `plot()` function is the primary high level command for producing graphics in R

```
x <- 1:100  
y <- rnorm(100)
```

```
plot(x, y)  
plot(y ~ x)
```

plot () is a Generic Function

```
# Dataframe (scatterplot matrix)  
plot(iris)
```

```
# Single numeric variable (scatterplot)  
plot(iris$Sepal.Length)
```

```
# Single factor variable (barplot)  
plot(iris$Species)
```

```
# Two numeric variables (scatterplot)  
plot(iris$Sepal.Length, iris$Petal.Length)
```

```
# Numeric and factor variable (dotplot)  
plot(iris$Sepal.Length, iris$Species)
```

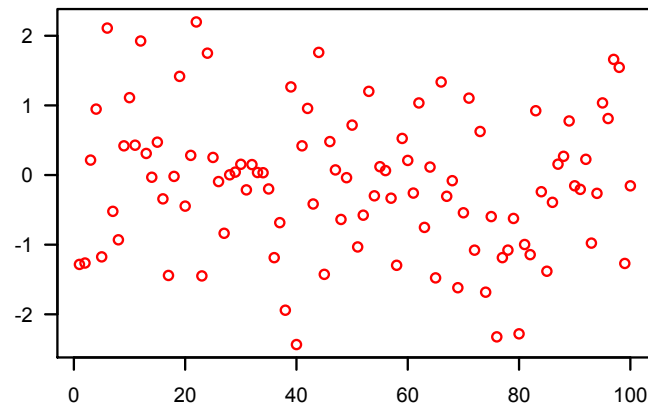
```
# Factor and numeric variable (boxplot)  
plot(iris$Species, iris$Sepal.Length)
```


plot() Arguments

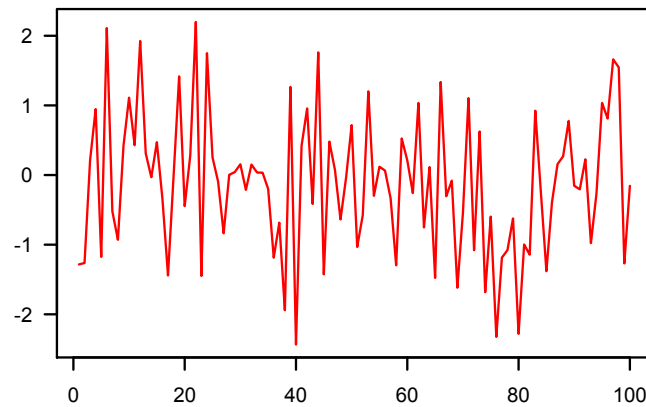
```
plot(x, y,  
     type = "p",  
     main = "Main-title",  
     sub = "Sub-title",  
     xlab = "X-lab",  
     ylab = "Y-lab",  
     asp = 0)
```

plot(x, y, type = ?)

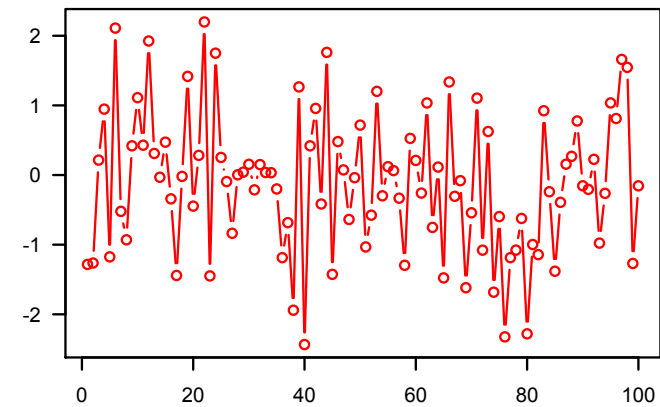
type = 'p'



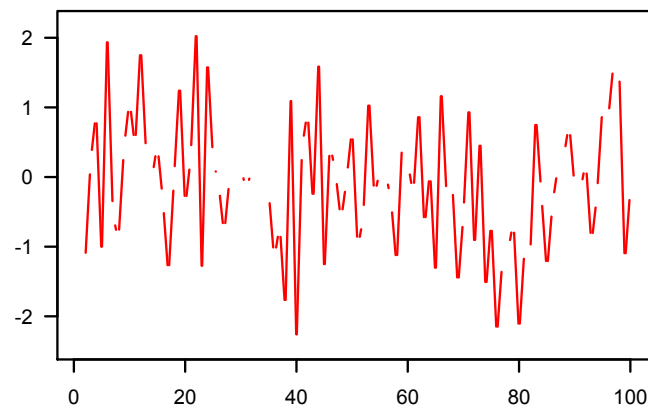
type = 'l'



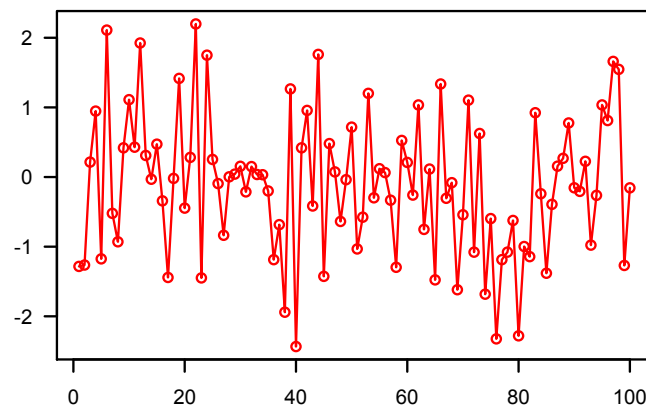
type = 'b'



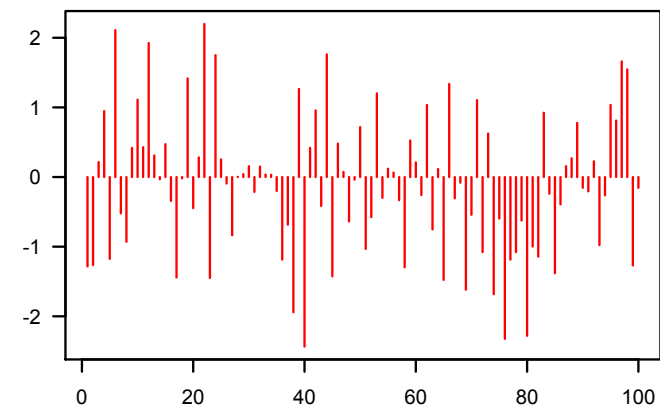
type = 'c'



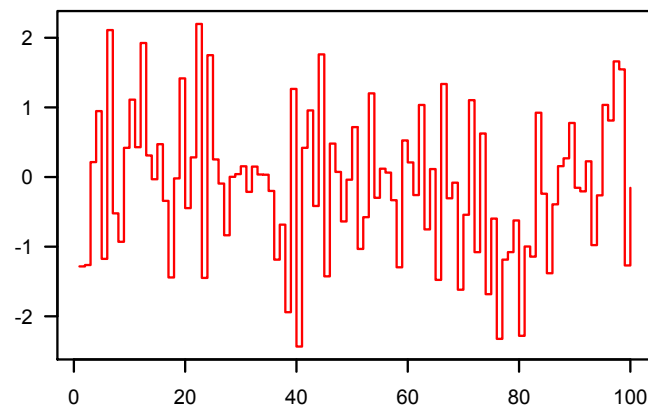
type = 'o'



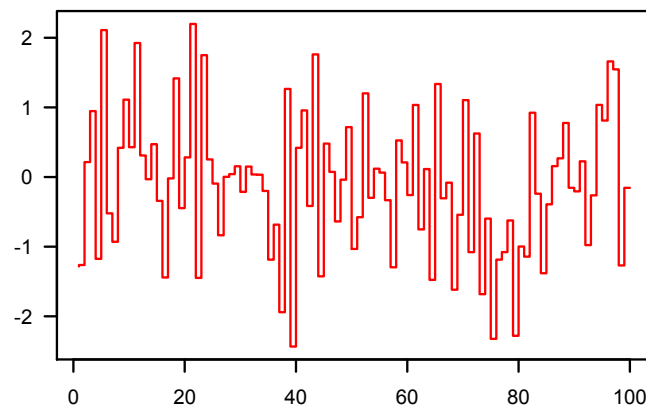
type = 'h'



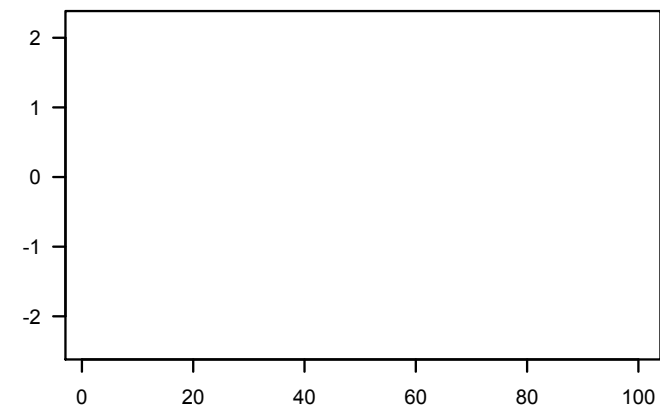
type = 's'



type = 'S'



type = 'n'



xlim and ylim

You can limit the scales of the x and y axis using `xlim` and `ylim`

```
x <- rnorm(100)
```

```
y <- rnorm(100)
```

```
plot(x, y)
```

```
plot(x, y, xlim = c(-10, 10))
```

```
plot(x, y, xlim = c(-10, 10),  
      ylim = c(-10, 10))
```

Graphics Functions

Base Graphics	Description	Lattice
barplot	Bargraph	barchart
biplot	Represents rows and columns of a matrix	
boxplot	Box and whisker plot	bwplot
contour	Contour plot	contourplot
coplot	Conditional plot	xyplot
dotchart	Cleveland dot chart	dotplot
filled.contour	Contour plot filled with colors	contourplot
hist	Histogram	histogram
image	High-density image plot	levelplot
interaction.plot	Interactions plot for two-factors	
matplot	Multiple plots specified by columns of a matrix	
pairs	All pairwise plots between variables	splo
persp	3D perspective plot	wireframe, cloud
pie	Pie chart	
plot	Plot	xyplot
qqnorm	Normal QQ plot	qqmath
qqplot	Quantile-Quantile plot	qq
scatter.smooth	Scatterplot with a smooth curve	
stars	Star plots for multivariate data	

You Try...

1. Using the `mtcars` dataset plot `mpg` vs. `cyl`
2. Create a histogram of the `mpg` variable in the `mtcars` dataset
3. Create boxplots for each variable in the `mtcars` dataset

Graphics Devices

Graphics Devices

All graphics have to be printed on a graphics device

Any call to a high level graphics command automatically opens a new graphics device if one is not already open

```
dev.new()  
plot(rnorm(100))
```

```
dev.new(height = 8, width = 8)  
  plot(runif(100))  
dev.off()
```

```
graphics.off()
```

Print to a File

R can easily print graphics to different file types (devices)

- `pdf()`
- `jpeg()`
- `png()`
- `tiff()`
- `bmp()`

Print to a PDF File

```
pdf(file = "graph.pdf")  
  plot(x, y, main = "Graph 1")  
  plot(y, x, main = "Graph 2")  
dev.off()
```

```
pdf(file = "graph.pdf", height = 4, width = 4)  
  plot(x, y)  
dev.off()
```

Print to a JPEG File

```
# Only one graph per file
jpeg(file = "graph.jpg")
  plot(x, y)
dev.off()
```

```
jpeg(file = "graph.jpg", width = 900,
      height = 900)
  plot(x, y)
dev.off()
```

You Try...

1. Using the `mtcars` dataset plot `mpg` vs. `cyl`
2. Save the plot you just created as a pdf file

Graphical Parameters

Graphics Parameters

Graphics parameters can be set to change almost any part of a graphic (over 70 parameters can be set)

- background
- labels
- colors
- margins of the plot
- font type
- font size
- orientation
- plotting characters
- aspect ratios
- tick marks
- axis label rotation
- number of plots
- line type
- line width
- plot dimensions
- box type ...

Temporary Changes to Parameters

Most graphical parameters can be set within the plot call (the exception being parameters that deal with the plot layout)

```
plot(x, y)
```

```
plot(x, y, col = 2, las = 1, pch = 19)
```

```
plot(x, y, col = 4, las = 1, pch = 12)
```

- Setting graphical parameters within the plot call only changes the parameters for that particular graph

Lasting Changes to Parameters

The `par()` function is used to set graphic parameters

- Setting graphical parameters with `par()` changes the parameters for the device (until the device is closed)

```
graphics.off()  
par(pch = 19, las = 1)
```

```
plot(x, y, col = 4)
```

```
plot(x, y, col = 6)
```

Example: Multiple Plots on a Device

```
par(mfrow = c(2, 2), mar = c(3, 3, 3, 2))
```

```
plot(iris$Sepal.Length, col = 2, main = "Sepal Length",  
     ylab = "Sepal length")
```

```
plot(iris$Petal.Length, col = 3, main = "Petal Length",  
     ylab = "Petal length")
```

```
plot(iris$Sepal.Width, col = 5, main = "Sepal Width",  
     ylab = "Sepal width")
```

```
plot(iris$Petal.Width, col = 6, main = "Petal Width",  
     ylab = "Petal width")
```

```
dev.off()
```


Example: Colored Plot

```
par(bg = "black",          # background color of device
    fg = "white",          # color of axis lines and ticks
    bty = "l",             # box type around plot
    las = 1,               # axis labels are horizontal
    col.axis = "white",    # color of axis
    col.lab = "white",     # color of labels
    col.main = "white",    # color of title
    mar = c(4, 4, 2, 1),   # margins of the plot
    tck = -0.03,           # length of tick marks
    font.main = 2,         # bold font for title
    font.lab = 2,          # bold font for axis labels
    family = "sans",       # use sans-serif font
    cex.lab = 1.2,         # make axis labels larger
    cex.axis = 1.1,        # make axis annotations larger
    pch = 19)              # change plotting character

plot(iris$Sepal.Length, col = 6, main = "Sepal Length",
     ylab = "Sepal length")
```

You Try...

1. Using the `mtcars` dataset plot `mpg` vs. `cyl`
2. Save the plot you just created as a pdf file
3. Change the color, plotting character, axis labels, and create a title for the plot



Base Graphics #2

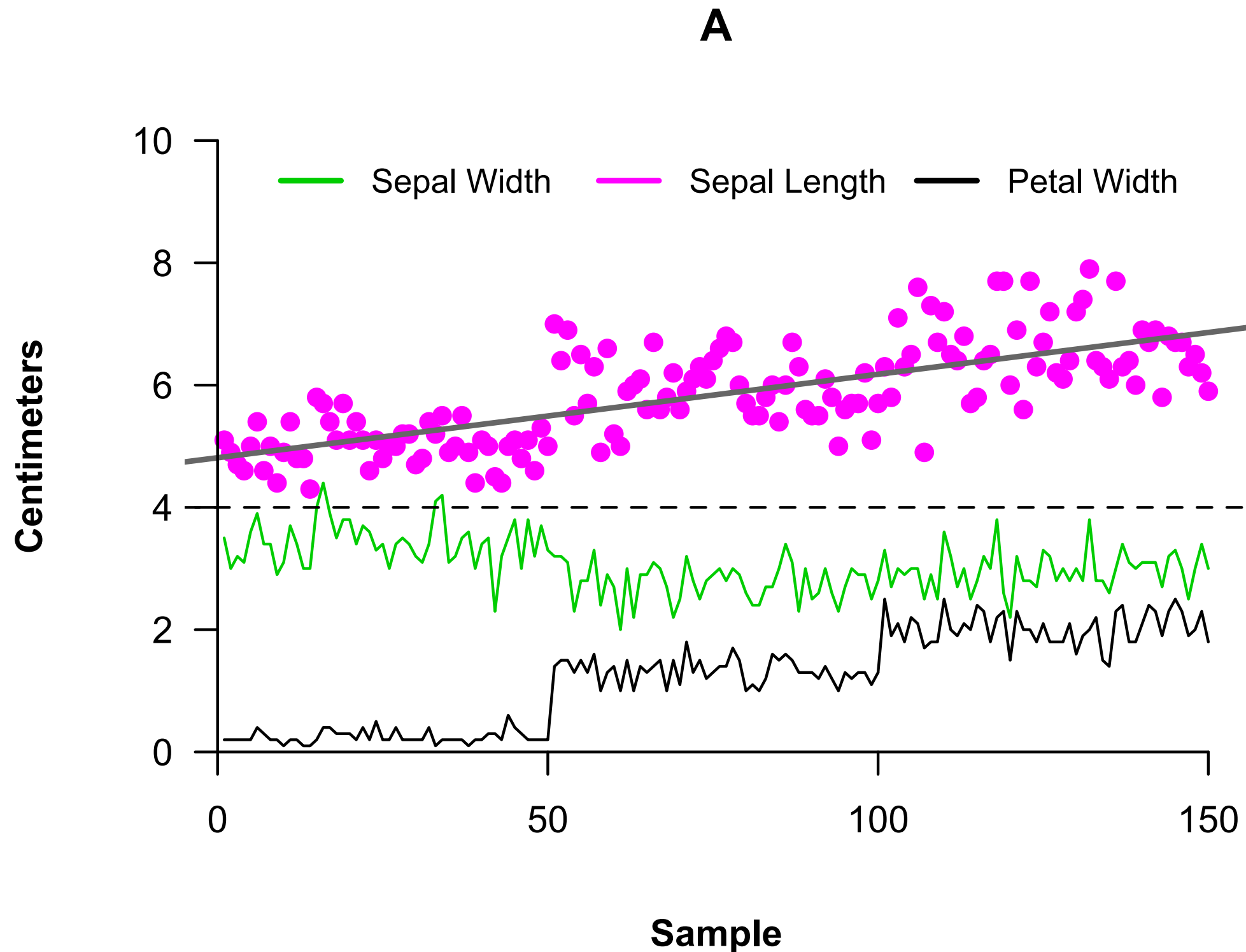
(Lines, Points, Legends, Axis)

Graphics Commands

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Graphic to Create ...



Functions We Need

- `plot()`
- `par()`
- `points()`
- `lines()`
- `abline()`
- `axis()`
- `legend()`

Step #1: Call `plot()`

Need to call a high-level plotting command first (`plot()`)

```
attach(iris)
```

```
dev.new(width = 5, height = 4)
```

```
par(pch = 19, las = 1, font.lab = 2)
```

```
plot(Sepal.Width, type = "l", ylim = c(0, 10), col = 3,  
     ylab = "Centimeters", axes = FALSE, xlab =  
     "Sample", main = "A")
```

Step #2: Add `axis()`

The `axis()` function adds axes to a plot

```
dev.new(width = 5, height = 4)
par(pch = 19, las = 1, font.lab = 2)

plot(Sepal.Width, type = "l", ylim = c(0, 10), col = 3,
      ylab = "Centimeters", axes = FALSE, xlab =
        "Sample", main = "A")

axis(1, pos = 0) # x-axis
axis(2, pos = 0) # y-axis
```


Step #3: Add `lines()`

The `lines()` function adds lines to an existing plot

```
dev.new(width = 5, height = 4)
par(pch = 19, las = 1, font.lab = 2)

plot(Sepal.Width, type = "l", ylim = c(0, 10), col = 3,
     ylab = "Centimeters", axes = FALSE, xlab =
     "Sample", main = "A")

axis(1, pos = 0) # x-axis
axis(2, pos = 0) # y-axis

lines(Petal.Width)
```

Step #4: Add `points()`

The `points()` function adds points to an existing plot

```
dev.new(width = 5, height = 4)
par(pch = 19, las = 1, font.lab = 2)

plot(Sepal.Width, type = "l", ylim = c(0, 10), col = 3,
     ylab = "Centimeters", axes = FALSE, xlab =
       "Sample", main = "A")

axis(1, pos = 0) # x-axis
axis(2, pos = 0) # y-axis

lines(Petal.Width)
points(Sepal.Length, col = 6)
```

Step #5: Add Reference Line

The `abline()` function adds vertical or horizontal reference lines

```
dev.new(width = 5, height = 4)
par(pch = 19, las = 1, font.lab = 2)

plot(Sepal.Width, type = "l", ylim = c(0, 10), col = 3,
     ylab = "Centimeters", axes = FALSE, xlab =
     "Sample", main = "A")

axis(1, pos = 0) # x-axis
axis(2, pos = 0) # y-axis

lines(Petal.Width)
points(Sepal.Length, col = 6)

abline(h = 4, lty = 2)
```

Step #6: Add Regression Line

The `abline()` function can also add regression lines to an existing plot

```
dev.new(width = 5, height = 4)
par(pch = 19, las = 1, font.lab = 2)

plot(Sepal.Width, type = "l", ylim = c(0, 10), col = 3,
     ylab = "Centimeters", axes = FALSE, xlab = "Sample",
     main = "A")

axis(1, pos = 0) # x-axis
axis(2, pos = 0) # y-axis

lines(Petal.Width)
points(Sepal.Length, col = 6)

abline(h = 4, lty = 2)

x <- 1:150
abline(lm(Sepal.Length ~ x), lwd = 2, col = "grey40")
```

Step #7: Add legend()

The legend() function creates a legend for the plot

```
dev.new(width = 5, height = 4)
par(pch = 19, las = 1, font.lab = 2)

plot(Sepal.Width, type = "l", ylim = c(0, 10), col = 3,
     ylab = "Centimeters", axes = FALSE, xlab = "Sample", main = "A")

axis(1, pos = 0)          # x-axis
axis(2, pos = 0, las = 1) # y-axis

lines(Petal.Width)
points(Sepal.Length, col = 6)

abline(h = 4, lty = 2)

x <- 1:150
abline(lm(Sepal.Length ~ x), lwd = 2, col = "grey40")

legend(x = 5, y = 10, legend = c("Sepal Width", "Sepal Length",
  "Petal Width"), col = c(3, 6, 1), lwd = 2, bty = "n",
  horiz = TRUE, cex = 0.9)
```

Entire Code

```
attach(iris)
dev.new(width = 5, height = 4)
par(pch = 19, las = 1, font.lab = 2)

plot(Sepal.Width, type = "l", ylim = c(0, 10), col = 3,
     ylab = "Centimeters", axes = FALSE, xlab = "Sample", main = "A")

axis(1, pos = 0) # x-axis
axis(2, pos = 0) # y-axis

lines(Petal.Width)
points(Sepal.Length, col = 6)

abline(h = 4, lty = 2)

x <- 1:150
abline(lm(Sepal.Length ~ x), lwd = 2, col = "grey40")

legend(x = 5, y = 10, legend = c("Sepal Width", "Sepal Length",
  "Petal Width"), col = c(3, 6, 1), lwd = 2, bty = "n",
  horiz = TRUE, cex = 0.9)
```

You Try...

1. Using the `mtcars` dataset plot `mpg vs. cyl`
2. Save the plot you just created as a pdf file
3. Change the color, plotting character, axis labels, and create a title for the plot
4. Add to the plot the `qsec vs. cyl` using a different color
 - Hint: use the `points()` function
5. Add a legend to the plot to differentiate between `mpg` and `qsec`

