

There are six pre-described line types available in base R. You can use those for any type of graphics, like plotting for line charts or developing simple shapes.

In R base plot functions, two options are available **lty** and **lwd**, **lty** stands for line types, and **lwd** for line width. The type of line you can be specified based on a number or a string. In R the default line type is “solid”.

In the case of **ggplot2** package, the parameters **linetype** and **size** are used to decide the type and the size of lines, respectively.

In this tutorial describes how to change line types in R for plots created using either the R base plot or from the ggplot2 package.

[Visualization Graphs-ggside with ggplot »](#)

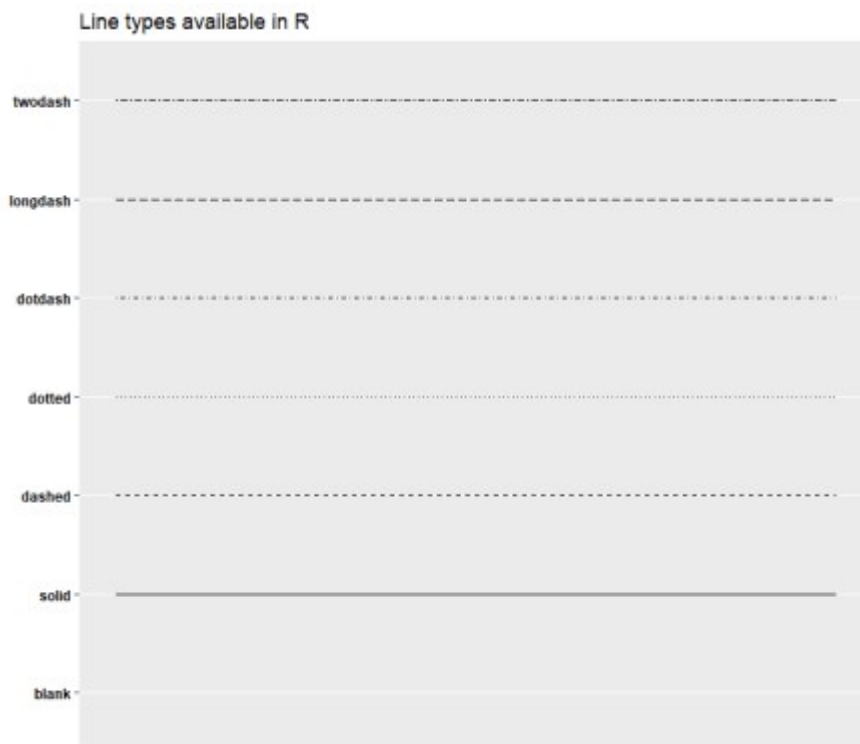
You will understand how to:

1. Use the different types line graphs in R.
2. Plot two lines and modify the line style for base plots and ggplot
3. Adjust the R line thickness by specifying the options **lwd** and **size**.
4. Change manually the appearance of **linetype**, color and size

Different line types in R

From ggpubr package with single line of code we can show the list of line types available in R.

```
library(ggpubr)
show_line_types()
```



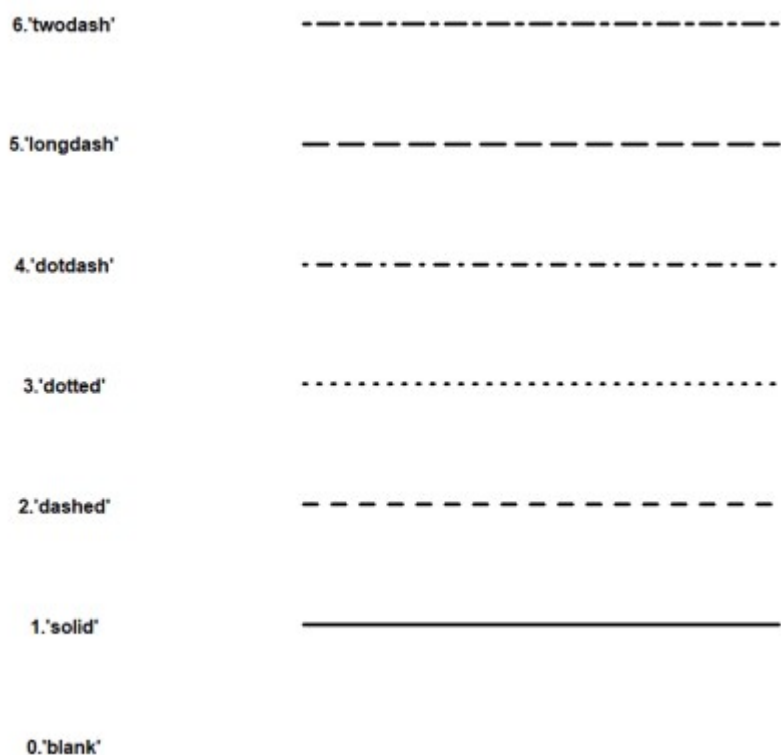
We can make use of the below function and write clear labels with number and string.

```
LineTypes<-function() {
```

```

oldPar<-par()  par(font=2, mar=c(0,0,0,0))
plot(1, pch="", ylim=c(0,6), xlim=c(0,0.7), axes=FALSE,xlab="",
ylab="")
for(i in 0:6) lines(c(0.3,0.7), c(i,i), lty=i, lwd=3)
text(rep(0.1,6), 0:6, labels=c("0.'blank'", "1.'solid'",
"2.'dashed'", "3.'dotted'",
                                "4.'dotdash'", "5.'longdash'",
                                "6.'twodash'"))
par(mar=oldPar$mar,font=oldPar$font )}
LineTypes()

```



Change R base plot line types

R lines functions:-

`plot(x, y, type = "l", lty = 1)`. Create the main plot
`lines(x, y, type = "l", lty = 1)`. Add lines onto the plot.

Key options:

- x, y: variables to be used for the x and y axes.
- type: display the data as line and/or point. Lowed values: l (display line only), p (show point only), and b (show both).
- pch and cex: setpoints shape and size, respectively.
- lty, lwd: set line types and thickness.
- col: change the color of point and line.
- xlab and ylab: for x and y-axis labels, respectively.

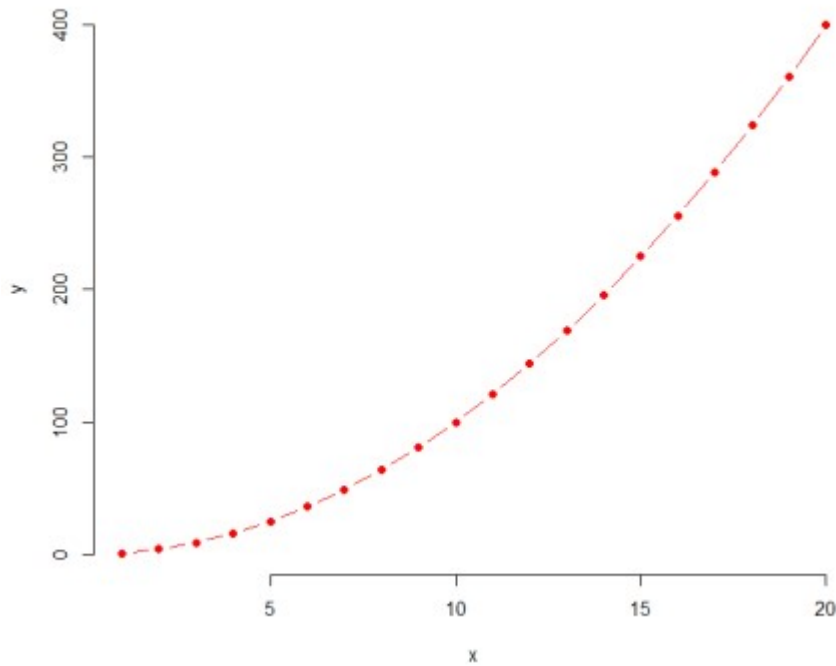
Create some variables for visualization,

[Principal component analysis \(PCA\) in R »](#)

```
x <- 1:20  
y1 <- x*x  
y2 <- 1*y1
```

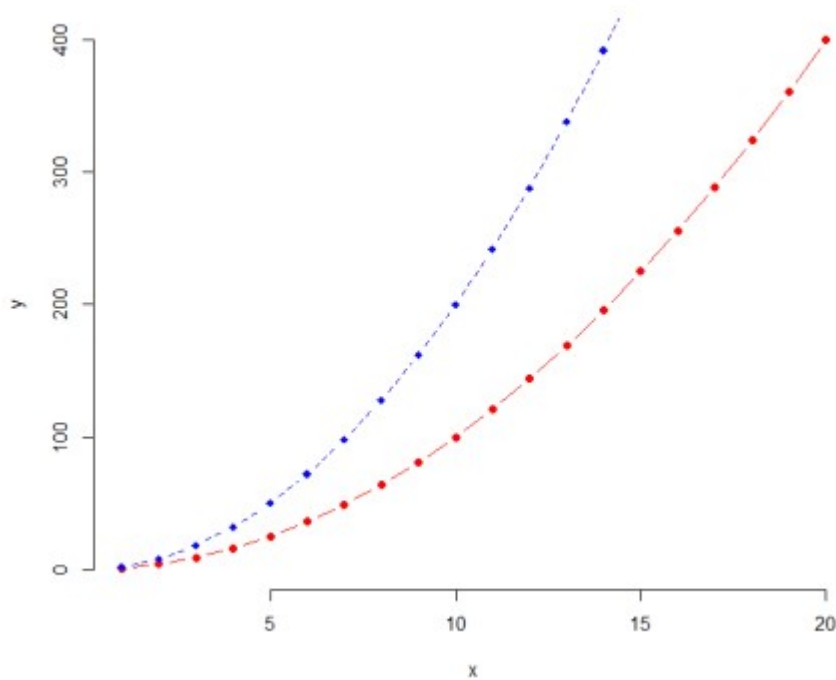
Just plot a first line based on plot function in R,

```
plot(x, y1, type = "b", frame = FALSE, pch = 19,  
     col = "red", xlab = "x", ylab = "y",  
     lty = 1, lwd = 1)
```



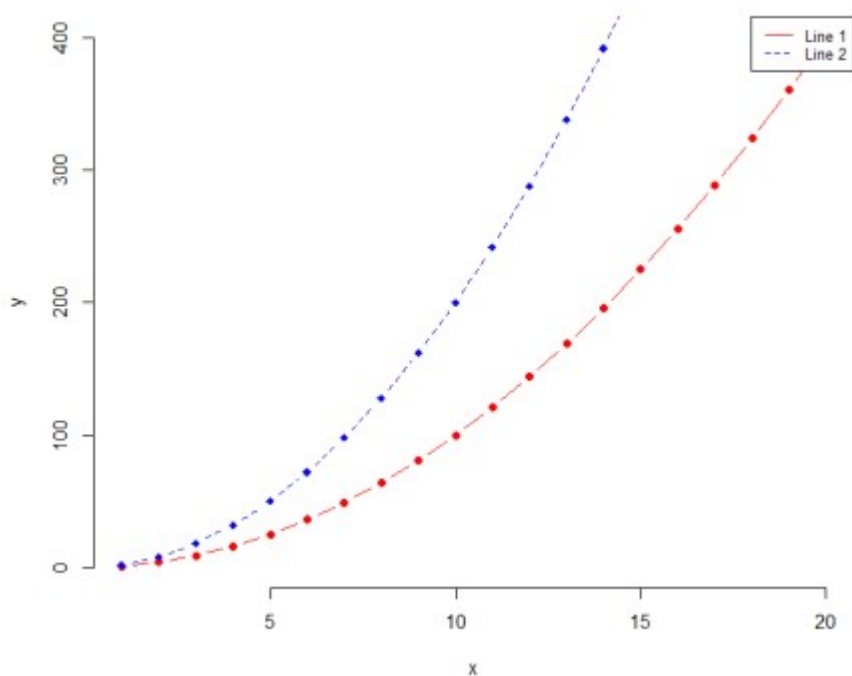
Now Add a second line

```
lines(x, y2, pch = 18, col = "blue", type = "b", lty = 2, lwd = 1)
```



If you want you can add a legend to the plot and set legend lty

```
legend("topright", legend = c("Line 1", "Line 2"),  
      col = c("red", "blue"), lty = 1:2, cex = 0.8)
```



Line Types in ggplot

First, let's load the data set.

[Decision Trees in R » Classification & Regression »](#)

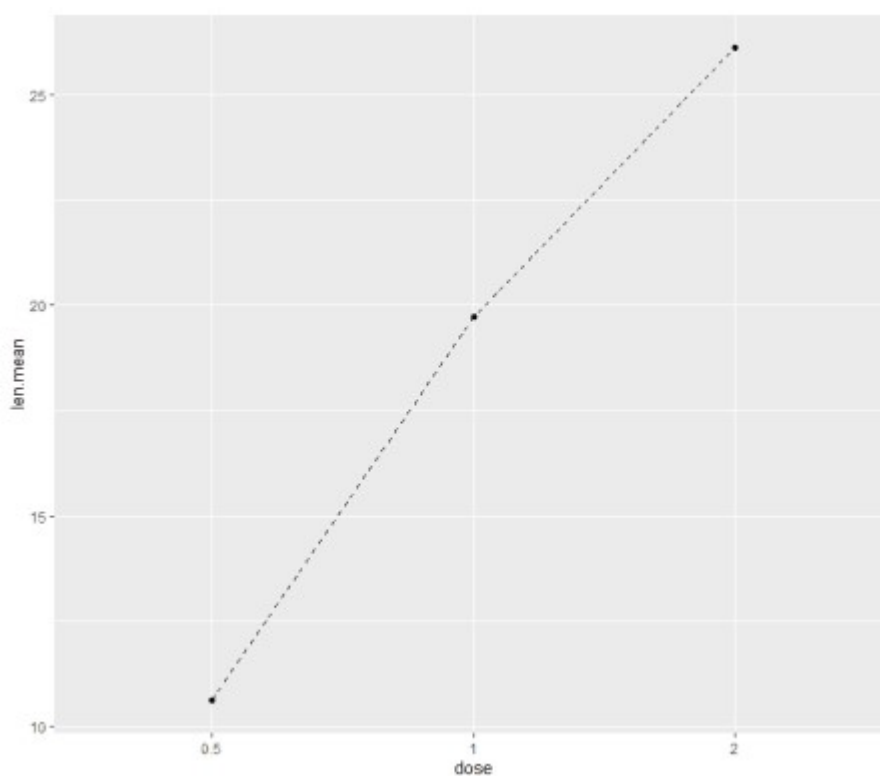
```
ToothGrowth$dose <- as.factor(ToothGrowth$dose)
head(ToothGrowth)
  len supp dose
1  4.2   VC  0.5
2 11.5   VC  0.5
3  7.3   VC  0.5
4  5.8   VC  0.5
5  6.4   VC  0.5
6 10.0   VC  0.5
library(dplyr)
df <- ToothGrowth %>%
  group_by(dose) %>%
  summarise(len.mean = mean(len))
df
```

Now created average values based on group by dose.

```
  dose len.mean
1  0.5    10.605
2    1    19.735
3    2    26.100
```

Let's plot the same

```
library(ggplot2)
ggplot(data = df, aes(x = dose, y = len.mean, group = 1)) +
  geom_line(linetype = "dashed") +
  geom_point()
```

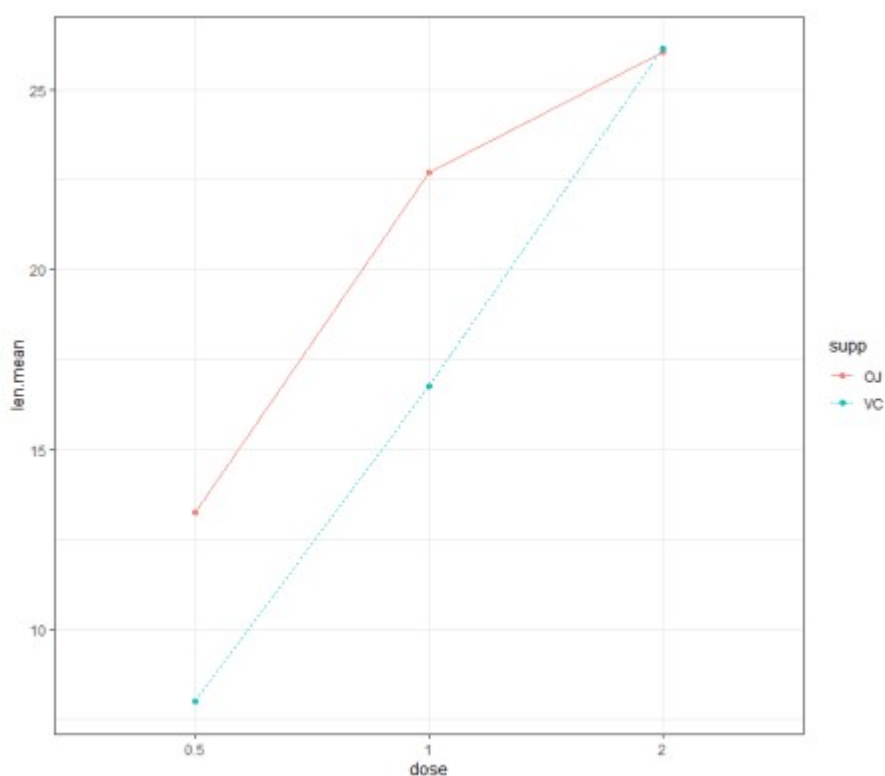


Create a line plot for multiple groups

[KNN Algorithm Machine Learning » Classification & Regression »](#)

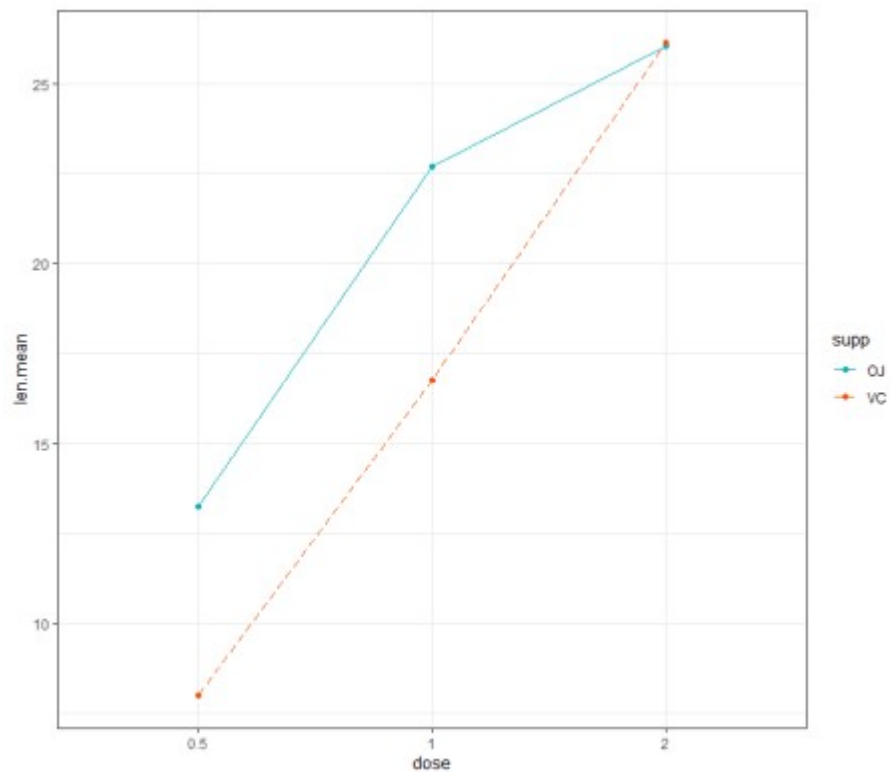
```
library(dplyr)
df2 <- ToothGrowth %>%
  group_by(dose, supp) %>%
  summarise(len.mean = mean(len))
df2
  dose supp len.mean
1  0.5   OJ    13.23
2  0.5   VC     7.98
3    1   OJ    22.70
4    1   VC    16.77
5    2   OJ    26.06
6    2   VC    26.14

ggplot(df2, aes(x = dose, y = len.mean, group = supp)) +
  geom_line(aes(linetype = supp, color = supp)) +
  geom_point(aes(color = supp)) + theme_bw()
```



Now change line type and color manually

```
ggplot(df2, aes(x = dose, y = len.mean, group = supp)) +
  geom_line(aes(linetype = supp, color = supp)) +
  geom_point(aes(color = supp)) + theme_bw() +
  scale_linetype_manual(values=c("solid", "longdash")) +
  scale_color_manual(values=c("#00AFBB", "#FC4E07"))
```



Conclusion

Use **lty** and **lwd** options, for changing lines type and thickness in R base graphics and in ggplot **linetype** and **size** are used.