



# Quick-R

accessing the power of R

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## R Interface

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## R in Action



## Line Charts

Line charts are created with the function `lines(x, y, type=)` where `x` and `y` are numeric vectors of `(x,y)` points to connect. `type=` can take the following values:

type	description
<code>p</code>	points
<code>l</code>	lines
<code>o</code>	overplotted points and lines
<code>b, c</code>	points (empty if "c") joined by lines
<code>s, S</code>	stair steps
<code>h</code>	histogram-like vertical lines
<code>n</code>	does not produce any points or lines

The `lines( )` function *adds* information to a graph. It can not produce a graph on its own. Usually it follows a `plot(x, y)` command that produces a graph.

By default, `plot( )` plots the `(x,y)` points. Use the `type="n"` option in the `plot( )` command, to create the graph with axes, titles, etc., but *without* plotting the points.

[R in Action](#) (2nd ed) significantly expands upon this material. Use promo code **ria38** for a 38% discount.

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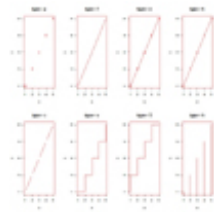
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In the following code each of the **type=** options is applied to the same dataset. The **plot( )** command sets up the graph, but *does not* plot the points.

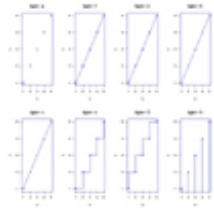
```
x <- c(1:5); y <- x # create some data
par(pch=22, col="red") # plotting symbol and color
par(mfrow=c(2,4)) # all plots on one page
opts = c("p","l","o","b","c","s","S","h")
for(i in 1:length(opts)){
  heading = paste("type=",opts[i])
  plot(x, y, type="n", main=heading)
  lines(x, y, type=opts[i])
}
```



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Next, we demonstrate each of the **type=** options when **plot( )** sets up the graph and *does* plot the points.

```
x <- c(1:5); y <- x # create some data
par(pch=22, col="blue") # plotting symbol and color
par(mfrow=c(2,4)) # all plots on one page
opts = c("p","l","o","b","c","s","S","h")
for(i in 1:length(opts)){
  heading = paste("type=",opts[i])
  plot(x, y, main=heading)
  lines(x, y, type=opts[i])
}
```



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As you can see, the **type="c"** option only looks different from the **type="b"** option if the plotting of points is suppressed in the **plot( )** command.

To demonstrate the creation of a more complex line chart, let's plot the growth of 5 orange trees over time. Each tree will have its own distinctive line. The data come from the dataset **Orange**.

```
# Create Line Chart

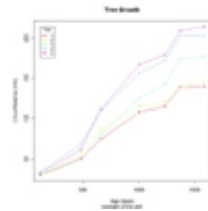
# convert factor to numeric for convenience
Orange$Tree <- as.numeric(Orange$Tree)
ntrees <- max(Orange$Tree)

# get the range for the x and y axis
xrange <- range(Orange$age)
yrange <- range(Orange$circumference)

# set up the plot
plot(xrange, yrange, type="n", xlab="Age (days)",
     ylab="Circumference (mm)" )
colors <- rainbow(ntrees)
linetype <- c(1:ntrees)
plotchar <- seq(18,18+ntrees,1)

# add lines
for (i in 1:ntrees) {
  tree <- subset(Orange, Tree==i)
  lines(tree$age, tree$circumference, type="b", lwd=1.5,
       lty=linetype[i], col=colors[i], pch=plotchar[i])
}
```

```
}  
  
# add a title and subtitle  
title("Tree Growth", "example of line plot")  
  
# add a legend  
legend(xrange[1], yrange[2], 1:ntrees, cex=0.8, col=colors,  
       pch=plotchar, lty=linetype, title="Tree")
```



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