

Statistical tools for high-throughput data analysis



Search...

Q

Home

Basics

Data

Visualize

Analyze

Product

Contribut

Suppor

About

Home / Easy Guides / R software / Data Visualization / ggplot2 - Essentials / ggplot2 scatter plots : Quick start guide - R software and data visualization

Actions menu for module Wiki



↑ ggplot2 scatter plots : Quick start guide - R software and data visualization

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Data Software

R Programming

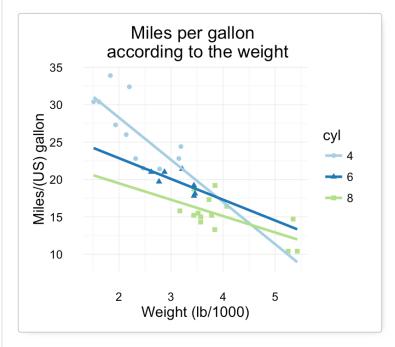
Color R

Create a Graph

≡Tools

- Prepare the data
- Basic scatter plots
- Label points in the scatter plot
 - Add regression lines
 - Change the appearance of points and lines
- Scatter plots with multiple groups
 - Change the point color/shape/size automatically
 - Add regression lines
 - Change the point color/shape/size manually
- Add marginal rugs to a scatter plot
- Scatter plots with the 2d density estimation
- Scatter plots with ellipses
- Scatter plots with rectangular bins
- Scatter plot with marginal density distribution plot
- Customized scatter plots
- Infos

This article describes how create a **scatter plot** using **R software** and **ggplot2** package. The function **geom_point()** is used.



Prepare the data

mtcars data sets are used in the examples below.

```
# Convert cyl column from a numeric to a factor variable
mtcars$cyl <- as.factor(mtcars$cyl)
head(mtcars)</pre>
```

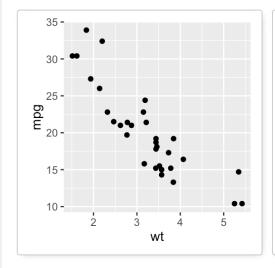
```
##
                      mpg cyl disp hp drat
                                                wt
                                                   qsec vs am gear carb
## Mazda RX4
                               160 110 3.90 2.620 16.46
## Mazda RX4 Wag
                     21.0
                               160 110 3.90 2.875 17.02
                                                             1
                                                                        4
## Datsun 710
                     22.8
                               108
                                    93 3.85 2.320 18.61
                                                                        1
                                                                        1
## Hornet 4 Drive
                               258 110 3.08 3.215 19.44
                     21.4
                            6
## Hornet Sportabout 18.7
                               360 175 3.15 3.440 17.02
                                                                        2
                            8
                                                                  3
                               225 105 2.76 3.460 20.22
## Valiant
                     18.1
                                                                        1
```

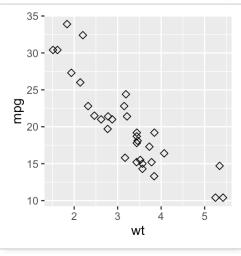
Basic scatter plots

Simple scatter plots are created using the R code below. The color, the size and the shape of points can be changed using the function **geom_point()** as follow:

```
geom_point(size, color, shape)
```

```
library(ggplot2)
# Basic scatter plot
ggplot(mtcars, aes(x=wt, y=mpg)) + geom_point()
# Change the point size, and shape
ggplot(mtcars, aes(x=wt, y=mpg)) +
  geom_point(size=2, shape=23)
```

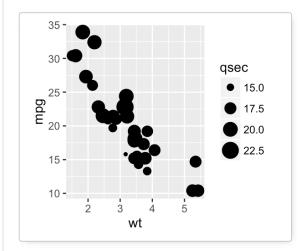






Note that, the size of the points can be controlled by the values of a continuous variable as in the example below.

```
# Change the point size
ggplot(mtcars, aes(x=wt, y=mpg)) +
geom_point(aes(size=qsec))
```

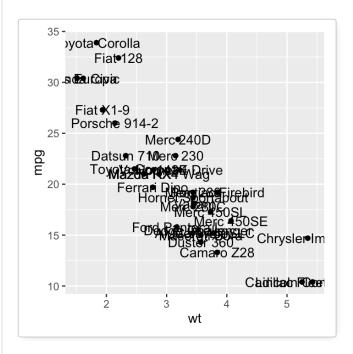


Read more on point shapes: ggplot2 point shapes

Label points in the scatter plot

The function **geom_text()** can be used:

```
ggplot(mtcars, aes(x=wt, y=mpg)) +
  geom_point() +
  geom_text(label=rownames(mtcars))
```



Read more on text annotations: ggplot2 - add texts to a plot

Add regression lines

The functions below can be used to add regression lines to a scatter plot:

- geom_smooth() and stat_smooth()
- geom_abline()

geom_abline() has been already described at this link: ggplot2 add straight lines to a plot.

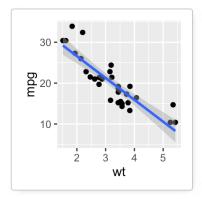
Only the function **geom_smooth()** is covered in this section.

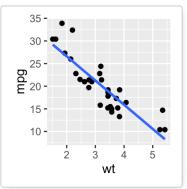
A simplified format is:

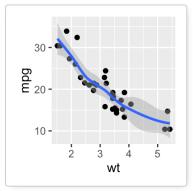
```
geom_smooth(method="auto", se=TRUE, fullrange=FALSE, level=0.95)
```

- **method**: smoothing method to be used. Possible values are lm, glm, gam, loess, rlm.
 - **method = "loess"**: This is the default value for small number of observations. It computes a smooth local regression. You can read more about **loess** using the R code **?loess**.
 - method ="Im": It fits a linear model. Note that, it's also possible to indicate the formula as formula = y ~ poly(x, 3) to specify a degree 3 polynomial.
- **se**: logical value. If TRUE, confidence interval is displayed around smooth.
- fullrange: logical value. If TRUE, the fit spans the full range of the plot
- level: level of confidence interval to use. Default value is 0.95

```
# Add the regression line
ggplot(mtcars, aes(x=wt, y=mpg)) +
    geom_point()+
    geom_smooth(method=lm)
# Remove the confidence interval
ggplot(mtcars, aes(x=wt, y=mpg)) +
    geom_point()+
    geom_smooth(method=lm, se=FALSE)
# Loess method
ggplot(mtcars, aes(x=wt, y=mpg)) +
    geom_point()+
    geom_point()+
    geom_smooth()
```



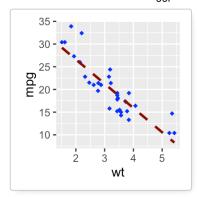


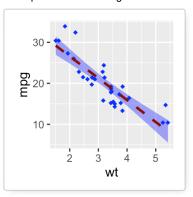


Change the appearance of points and lines

This section describes how to change:

- the color and the shape of points
- the line type and color of the regression line
- the fill color of the confidence interval







Note that a transparent color is used, by default, for the confidence band. This can be changed by using the argument alpha: geom_smooth(fill="blue", alpha=1)

Read more on point shapes: ggplot2 point shapes

Read more on line types: ggplot2 line types

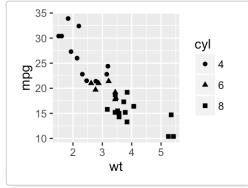
Scatter plots with multiple groups

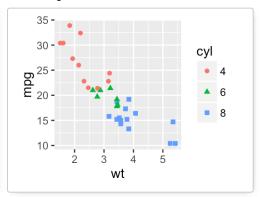
This section describes how to change point colors and shapes automatically and manually.

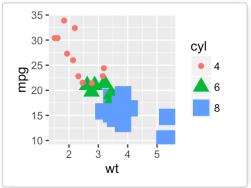
Change the point color/shape/size automatically

In the R code below, point shapes, colors and sizes are controlled by the levels of the factor variable cyl:

```
# Change point shapes by the levels of cyl
ggplot(mtcars, aes(x=wt, y=mpg, shape=cyl)) +
  geom_point()
# Change point shapes and colors
ggplot(mtcars, aes(x=wt, y=mpg, shape=cyl, color=cyl)) +
  geom_point()
# Change point shapes, colors and sizes
ggplot(mtcars, aes(x=wt, y=mpg, shape=cyl, color=cyl, size=cyl)) +
  geom_point()
```



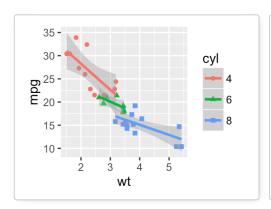


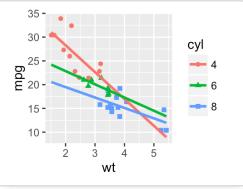


Add regression lines

Regression lines can be added as follow:

```
# Add regression lines
ggplot(mtcars, aes(x=wt, y=mpg, color=cyl, shape=cyl)) +
  geom_point() +
  geom_smooth(method=lm)
# Remove confidence intervals
# Extend the regression lines
ggplot(mtcars, aes(x=wt, y=mpg, color=cyl, shape=cyl)) +
  geom_point() +
  geom_smooth(method=lm, se=FALSE, fullrange=TRUE)
```



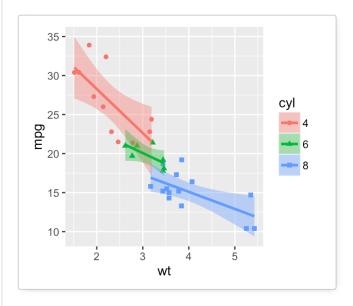




 \blacksquare Note that, you can also change the line type of the regression lines by using the aesthetic *linetype* = cyl.

The fill color of confidence bands can be changed as follow:

```
ggplot(mtcars, aes(x=wt, y=mpg, color=cyl, shape=cyl)) +
geom_point() +
geom_smooth(method=lm, aes(fill=cyl))
```



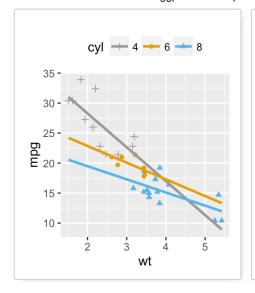
Change the point color/shape/size manually

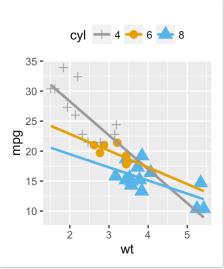
The functions below are used:

- scale_shape_manual() for point shapes
- scale_color_manual() for point colors
- scale_size_manual() for point sizes

```
# Change point shapes and colors manually
ggplot(mtcars, aes(x=wt, y=mpg, color=cyl, shape=cyl)) +
geom_point() +
geom_smooth(method=lm, se=FALSE, fullrange=TRUE)+
scale_shape_manual(values=c(3, 16, 17))+
scale_color_manual(values=c('#999999', '#E69F00', '#56B4E9'))+
theme(legend.position="top")

# Change the point sizes manually
ggplot(mtcars, aes(x=wt, y=mpg, color=cyl, shape=cyl))+
geom_point(aes(size=cyl)) +
geom_smooth(method=lm, se=FALSE, fullrange=TRUE)+
scale_shape_manual(values=c(3, 16, 17))+
scale_color_manual(values=c(3, 16, 17))+
scale_size_manual(values=c(2,3,4))+
theme(legend.position="top")
```

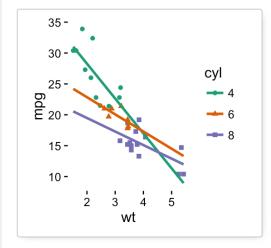


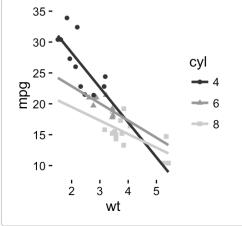


It is also possible to *change manually point and line colors* using the functions:

- scale_color_brewer(): to use color palettes from RColorBrewer package
- *scale_color_grey()* : to use grey color palettes

```
p <- ggplot(mtcars, aes(x=wt, y=mpg, color=cyl, shape=cyl)) +
    geom_point() +
    geom_smooth(method=lm, se=FALSE, fullrange=TRUE)+
    theme_classic()
# Use brewer color palettes
p+scale_color_brewer(palette="Dark2")
# Use grey scale
p + scale_color_grey()</pre>
```





Read more on ggplot2 colors here: ggplot2 colors

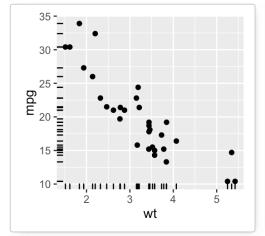
Add marginal rugs to a scatter plot

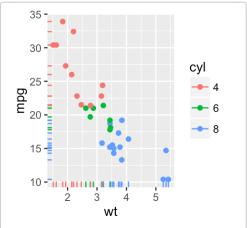
The function **geom_rug()** can be used:

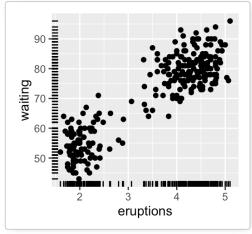
```
geom_rug(sides ="bl")
```

sides: a string that controls which sides of the plot the rugs appear on. Allowed value is a string containing any of "trbl", for top, right, bottom, and left.

```
# Add marginal rugs
ggplot(mtcars, aes(x=wt, y=mpg)) +
   geom_point() + geom_rug()
# Change colors
ggplot(mtcars, aes(x=wt, y=mpg, color=cyl)) +
   geom_point() + geom_rug()
# Add marginal rugs using faithful data
ggplot(faithful, aes(x=eruptions, y=waiting)) +
   geom_point() + geom_rug()
```





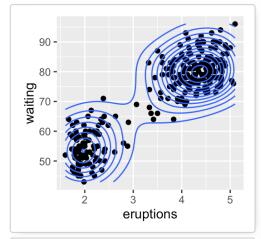


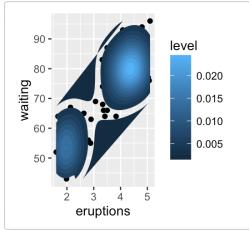
Scatter plots with the 2d density estimation

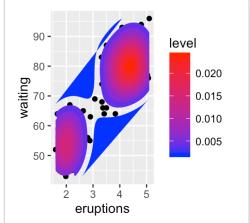
The functions **geom_density_2d()** or **stat_density_2d()** can be used:

```
# Scatter plot with the 2d density estimation
sp <- ggplot(faithful, aes(x=eruptions, y=waiting)) +
   geom_point()
sp + geom_density_2d()</pre>
```

```
# Gradient color
sp + stat_density_2d(aes(fill = ..level..), geom="polygon")
# Change the gradient color
sp + stat_density_2d(aes(fill = ..level..), geom="polygon")+
    scale_fill_gradient(low="blue", high="red")
```





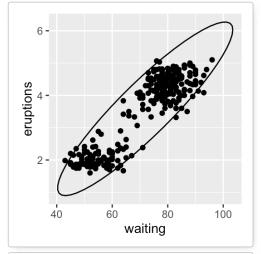


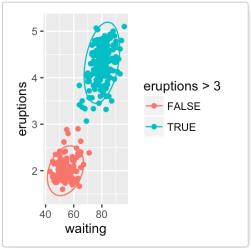
Read more on ggplot2 colors here: ggplot2 colors

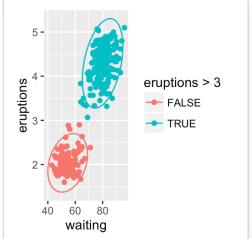
Scatter plots with ellipses

The function **stat_ellipse()** can be used as follow:

```
# One ellipse arround all points
ggplot(faithful, aes(waiting, eruptions))+
  geom_point()+
  stat_ellipse()
# Ellipse by groups
p <- ggplot(faithful, aes(waiting, eruptions, color = eruptions > 3))+
  geom_point()
p + stat_ellipse()
# Change the type of ellipses: possible values are "t", "norm", "euclid"
p + stat_ellipse(type = "norm")
```







Scatter plots with rectangular bins

The number of observations is counted in each bins and displayed using any of the functions below:

- **geom_bin2d()** for adding a heatmap of 2d bin counts
- stat_bin_2d() for counting the number of observation in rectangular bins
- stat_summary_2d() to apply function for 2D rectangular bins

The simplified formats of these functions are:

```
plot + geom_bin2d(...)
plot+stat_bin_2d(geom=NULL, bins=30)
plot + stat_summary_2d(geom = NULL, bins = 30, fun = mean)
```

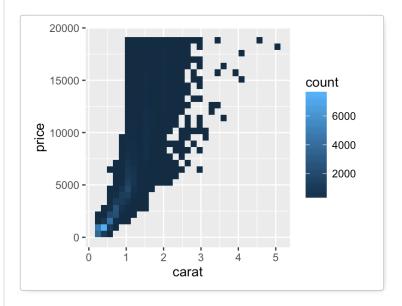
- **geom**: geometrical object to display the data
- bins: Number of bins in both vertical and horizontal directions. The default value is 30
- **fun** : function for summary

The data sets diamonds from ggplot2 package is used:

```
head(diamonds)
```

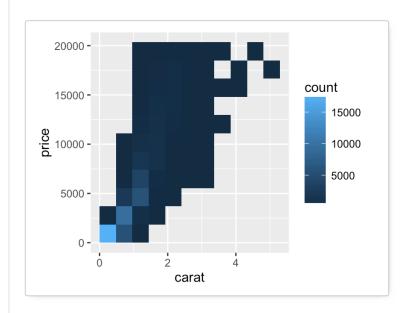
```
cut color clarity depth table price
##
     carat
                                                         Х
                                                               У
      0.23
                         Ε
                                SI2
                                    61.5
                                             55
                                                  326 3.95 3.98 2.43
## 1
               Ideal
## 2
      0.21
             Premium
                         Ε
                                SI1
                                     59.8
                                             61
                                                  326 3.89 3.84 2.31
## 3
      0.23
                Good
                         Ε
                                VS1
                                    56.9
                                             65
                                                  327 4.05 4.07 2.31
      0.29
             Premium
                         Ι
                                VS2
                                    62.4
                                             58
                                                  334 4.20 4.23 2.63
##
  4
      0.31
                Good
                         J
                                SI2 63.3
                                             58
                                                  335 4.34 4.35 2.75
## 5
                         J
     0.24 Very Good
                               VVS2 62.8
                                             57
                                                  336 3.94 3.96 2.48
```

```
# Plot
p <- ggplot(diamonds, aes(carat, price))
p + geom_bin2d()</pre>
```



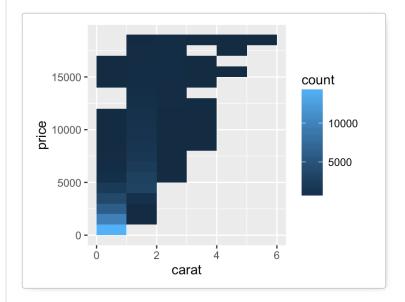
Change the number of bins:

```
# Change the number of bins
p + geom_bin2d(bins=10)
```



Or specify the width of bins:

```
# Or specify the width of bins
p + geom_bin2d(binwidth=c(1, 1000))
```



Scatter plot with marginal density distribution plot

Step 1/3. Create some data:

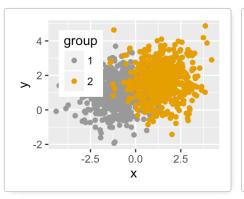
```
set.seed(1234)
x <- c(rnorm(500, mean = -1), rnorm(500, mean = 1.5))
y <- c(rnorm(500, mean = 1), rnorm(500, mean = 1.7))
group <- as.factor(rep(c(1,2), each=500))
df <- data.frame(x, y, group)
head(df)</pre>
```

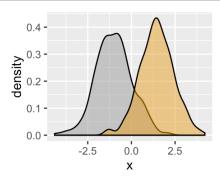
```
## 1 -2.20706575 -0.2053334 1
## 2 -0.72257076 1.3014667 1
## 3 0.08444118 -0.5391452 1
## 4 -3.34569770 1.6353707 1
## 5 -0.57087531 1.7029518 1
## 6 -0.49394411 -0.9058829 1
```

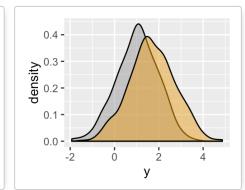
Step 2/3. Create the plots :

```
# scatter plot of x and y variables
# color by groups
scatterPlot <- ggplot(df,aes(x, y, color=group)) +
   geom_point() +
   scale_color_manual(values = c('#999999','#E69F00')) +</pre>
```

```
theme(legend.position=c(0,1), legend.justification=c(0,1))
scatterPlot
# Marginal density plot of x (top panel)
xdensity <- ggplot(df, aes(x, fill=group)) +
    geom_density(alpha=.5) +
    scale_fill_manual(values = c('#9999999','#E69F00')) +
    theme(legend.position = "none")
xdensity
# Marginal density plot of y (right panel)
ydensity <- ggplot(df, aes(y, fill=group)) +
    geom_density(alpha=.5) +
    scale_fill_manual(values = c('#9999999','#E69F00')) +
    theme(legend.position = "none")
ydensity</pre>
```







Create a blank placeholder plot:

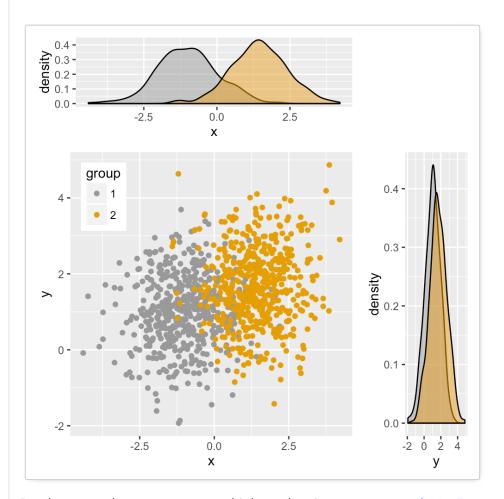
```
blankPlot <- ggplot()+geom_blank(aes(1,1))+
    theme(plot.background = element_blank(),
    panel.grid.major = element_blank(),
    panel.grid.minor = element_blank(),
    panel.border = element_blank(),
    panel.background = element_blank(),
    axis.title.x = element_blank(),
    axis.title.y = element_blank(),
    axis.text.x = element_blank(),
    axis.text.y = element_blank(),
    axis.ticks = element_blank())</pre>
```

Step 3/3. Put the plots together:

To put multiple plots on the same page, the package **gridExtra** can be used. Install the package as follow:

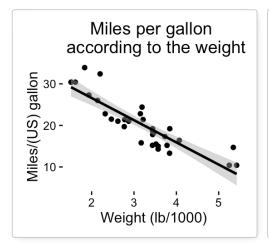
```
install.packages("gridExtra")
```

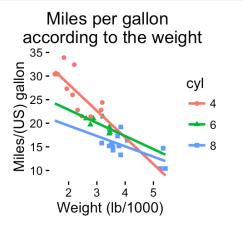
Arrange ggplot2 with adapted height and width for each row and column:



Read more on how to arrange multiple ggplots in one page : ggplot2 - Easy way to mix multiple graphs on the same page

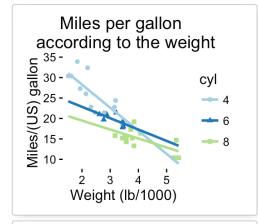
Customized scatter plots

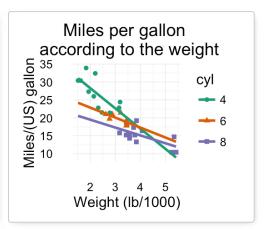


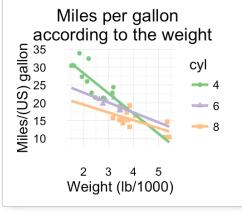


Change colors manually:

```
# Continuous colors
p + scale_color_brewer(palette="Paired") + theme_classic()
# Discrete colors
p + scale_color_brewer(palette="Dark2") + theme_minimal()
# Gradient colors
p + scale_color_brewer(palette="Accent") + theme_minimal()
```







Read more on ggplot2 colors here: ggplot2 colors

Infos



This analysis has been performed using **R software** (ver. 3.2.4) and **ggplot2** (ver. 2.1.0)



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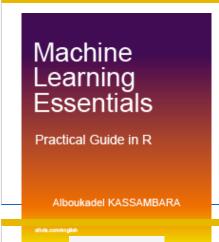
Show me some love with the like buttons below... Thank you and please don't forget to share and comment below!!

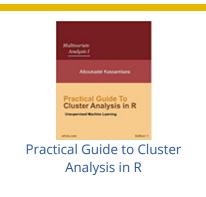
(i) Ads by Google Create a Graph 3D Chart Excel Plot Map Excel Data Graph

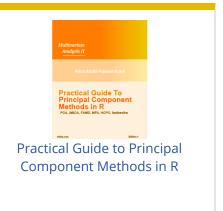
Share 10 Like 10 Tweet Share G+ Save Share 57



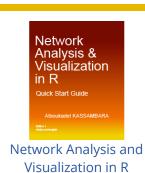
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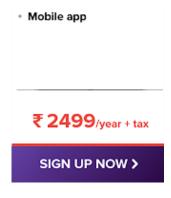
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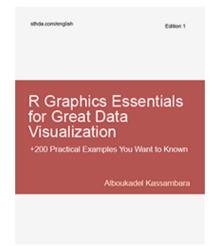


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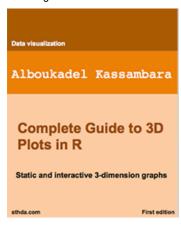
Practical Guide to Cluster Analysis in R



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I have neer used any machine learning things before, so this website was extremely useful in introducing the basics of working with clustering in R Studio! What a nice and "down to earth" way of expla... [Read more]

By Kate

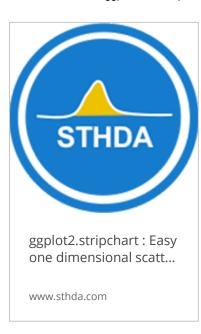
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