



Studio<sup>®</sup>

# Lecture 3: Introduction to R Part II

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# AGENDA

**01** Conditions and Loops

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**02** Functions

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**03** R Graphics

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**04** Documentation

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# Conditions and Loops

- Understanding conditions and loops are necessary for efficient data analysis
  - ✓ Example of conditions
    - Want to remove instances whose value is greater than 3 standard deviations
    - Want to remove variables with zero variance
    - Want to replace NULL with a constant value
  - ✓ Example of loops
    - Want to make a histogram for each variable in a dataframe
    - Want to compare various machine learning algorithms for the same dataset

# Conditions

- if-else condition

```
if (condition) {  
  statement 1  
} else {  
  statement 2  
}
```

- ✓ condition can be a simple logical comparison to a complex function
- ✓ statement 1: run if the condition is met
- ✓ statement 2: run if the condition is not met

```
> r <- 1  
> if (r==4) {  
+   printf("The value of r is 4")  
+ } else {  
+   print("The value of r is not 4")  
+ }  
[1] "The value of r is not 4"
```

```
> carbon <- c(10, 12, 15, 19, 20)  
> if (mean(carbon) > median(carbon)) {  
+   print ("Mean > Median")  
+ } else {  
+   print ("Median <= Mean")  
+ }  
[1] "Mean > Median"
```

```
> if (mean(carbon) > median(carbon)) {  
+   print ("Mean > Median")  
+ }  
[1] "Mean > Median"  
> else {  
Error: unexpected 'else' in "else"  
>   print ("Median <= Mean")  
[1] "Median <= Mean"  
> }  
Error: unexpected '}' in "}"
```

# Conditions

- ifelse: a vectorized condition

ifelse (condition, statement 1, statement 2)

- ✓ condition: Boolean vector
- ✓ statement 1: run if the condition is met
- ✓ statement 2: run if the condition is not met

```
> x <- 1:10  
> y <- ifelse(x%%2 == 0, "even", "odd")  
> y  
[1] "odd"  "even" "odd"  "even" "odd"  "even" "odd"  "even" "odd"  "even"
```

# Loops

- for loop

```
for (i in x) {  
  statement  
}
```

- ✓ i: index of loop
- ✓ x: a set of element for which the loop runs
- ✓ statement: running part

```
> for (i in n) {  
+   print(i^2)  
+ }  
[1] 25  
[1] 100  
[1] 225
```

# Loop

- while loop

```
while (condition) {  
  statement  
}
```

✓ run the statement until the condition is not met

```
> i <- 1  
> while (i <= 10) {  
+   i <- i+4  
+   print(i)  
+ }  
[1] 5  
[1] 9  
[1] 13
```

- repeat-break loop

```
repeat {  
  statement  
  condition break  
}
```

✓ run the statement first, check the condition, stop if the condition is met

```
> i <- 1  
> repeat {  
+   i <- i+4  
+   print(i)  
+   if (i > 10) break  
+ }  
[1] 5  
[1] 9  
[1] 13
```

# AGENDA

**01** Conditions and Loops

---

**02** **Functions**

---

**03** R Graphics

---

**04** Documentation

---



# Function

- Why functions?
- An incidental advantage of putting code into functions is that **the workspace is not then cluttered with objects that are local to the function**

```
all()      # returns TRUE if all values are TRUE
any()      # returns TRUE if any values are TRUE
args()     # information on the arguments to a function
cat()      # prints multiple objects, one after the other
cumprod()  # cumulative product
cumsum()   # cumulative sum
diff()     # form vector of first differences
           # N. B. diff(x) has one less element than x
history()  # displays previous commands used
is.factor() # returns TRUE if the argument is a factor
is.na()    # returns TRUE if the argument is an NA
           # NB also is.logical(), is.matrix(), etc.
length()   # number of elements in a vector or of a list
ls()       # list names of objects in the workspace
```

# Function

- Why functions?
- An incidental advantage of putting code into functions is that the workspace is not then cluttered with objects that are local to the function

```
mean()      # mean of the elements of a vector
median()    # median of the elements of a vector
order()     # x[order(x)] sorts x (by default, NAs are last)
print()     # prints a single R object
range()     # minimum and maximum value elements of vector
sort()      # sort elements into order, by default omitting NAs
rev()       # reverse the order of vector elements
str()       # information on an R object
unique()    # form the vector of distinct values
which()     # locates 'TRUE' indices of logical vectors
which.max() # locates (first) maximum of a numeric vector
which.min() # locates (first) minimum of a numeric vector
with()      # do computation using columns of specified data frame
```

# Function

- Writing a function

```
function_name <- function(arguments) {  
  statement 1  
  statement 2  
  ...  
  return(object)  
}
```

- ✓ function\_name: name that the function is referred to
- ✓ arguments: inputs that a user should provide to run the function
- ✓ statements: operations running inside the function
- ✓ object: function output

# Function

- Same operations but different outputs

```
> mean.and.sd1 <- function(x) {  
+   av <- mean(x)  
+   sdev <- sd(x)  
+   return(c(mean=av, SD=sdev))  
+ }  
> distance <- c(148, 182, 173, 166, 109, 141, 166)  
> mean.and.sd1(distance)  
      mean      SD  
155.00000  24.68468
```

```
> mean.and.sd2 <- function(x) {  
+   av <- mean(x)  
+   sdev <- sd(x)  
+   c(mean=av, SD=sdev)  
+   return(av)  
+ }  
> distance <- c(148, 182, 173, 166, 109, 141, 166)  
> mean.and.sd2(distance)  
[1] 155
```

# Function

- Writing a function

- ✓ Functions can accept various types of input arguments
- ✓ It is possible to provide default arguments
  - Default arguments are used if a user calls a function but does not provide the required input arguments

```
> mean.and.sd3 <- function(x = rnorm(10)) {  
+   av <- mean(x)  
+   sdev <- sd(x)  
+   c(mean=av, SD=sdev)  
+ }  
> mean.and.sd3()  
      mean      SD  
-0.188425  0.605823  
> mean.and.sd3(distance)  
      mean      SD  
155.00000  24.68468
```


# Function


- Function arguments
  - ✓ Each argument has its own name
  - ✓ Name is used to access the corresponding argument within function
  - ✓ Three possible ways to assign the argument
    - Exact name
    - Partially matching names (not recommended)
    - Argument order

```
> addTheLog <- function(first, second) {first + log(second)}  
> addTheLog(second=exp(4),first=1)  
[1] 5  
> addTheLog(s=exp(4),first=1)  
[1] 5  
> addTheLog(1,exp(4))  
[1] 5
```

# Function


- Return object in function
  - ✓ All R object can be the return object
  - ✓ Use `return( )` function
  - ✓ If `return( )` is not used, the result of the last operation is returned (not recommended)

```
Console ~/   
> oddcount <- function(x) {  
+ k <- 0  
+ print(sprintf("odd number calculator"))  
+ for (n in 1:x) {  
+ if (n %% 2 == 1) {  
+ cat(sprintf("%d is an odd number. \n", n))  
+ k <- k+1  
+ }  
+ }  
+ return(k)  
+ }  
> oddcount(10)  
[1] "odd number calculator"  
1 is an odd number.  
3 is an odd number.  
5 is an odd number.  
7 is an odd number.  
9 is an odd number.  
[1] 5
```

```
Console ~/   
> oddcount <- function(x) {  
+ k <- 0  
+ print(sprintf("odd number calculator"))  
+ for (n in 1:x) {  
+ if (n %% 2 == 1) {  
+ cat(sprintf("%d is an odd number. \n", n))  
+ k <- k+1  
+ }  
+ }  
+ k  
+ }  
> oddcount(10)  
[1] "odd number calculator"  
1 is an odd number.  
3 is an odd number.  
5 is an odd number.  
7 is an odd number.  
9 is an odd number.  
[1] 5
```

# Function

- Return object in function
  - ✓ All R object can be the return object
  - ✓ Use return( ) function
  - ✓ If return( ) is not used, the result of the last operation is returned (not recommended)

```
Console ~/   
> # Return the result without either return() or explicit designation  
> oddcount <- function(x) {  
+ k <- 0  
+ print(sprintf("odd number calculator"))  
+ for (n in 1:x) {  
+ if (n %% 2 == 1) {  
+ cat(sprintf("%d is an odd number. \n", n))  
+ k <- k+1  
+ }  
+ }  
+ }  
> oddcount(10)  
[1] "odd number calculator"  
1 is an odd number.  
3 is an odd number.  
5 is an odd number.  
7 is an odd number.  
9 is an odd number.
```



# Function

- Function example 1

✓ Question: from a vector consisting of only 0 and 1, return the indices from which 1 repeatedly appears k times

- Ex: (1, 1, 1, 0, 0, 1, 0, 1, 1, 0, 1, 1, 1)
  - k=2: answer = (1, 2, 8, 11, 12)
  - k=3: answer = (1, 11)
  - k=4: answer = NULL

```
> findrepeats <- function(x, k) {  
+   n <- length(x)  
+   repeats <- NULL  
+   for (i in 1:(n-k+1)) {  
+     if(all(x[i:(i+k-1)] == 1)) repeats <- c(repeats, i)  
+   }  
+   return(repeats)  
+ }  
> vec <- c(0,1,1,0,0,1,1,1,0,1,1)  
> findrepeats(vec,2)  
[1] 2 6 7 10  
> findrepeats(vec,3)  
[1] 6  
> findrepeats(vec,4)  
NULL
```

# Function

- Function example 2: Kendall's tau

✓ Raw data: temperature and pressure recorded every hour

Time	10:00	11:00	12:00	13:00	14:00
Temperature	10	15	13	17	20
Pressure	900	920	890	940	920

✓ What to do

- Determine whether each indicator increases or decreases
- Return the proportion of the events in which the change directions of the two indicators are the same

```
# Example 2: Kendall's tau
findud <- function(v) {
  vud <- v[-1] - v[-length(v)]
  return(ifelse(vud > 0, 1, -1))
}
```

```
udcorr <- function(x,y) {
  ud <- lapply(list(x,y), findud)
  return(mean(ud[[1]] == ud[[2]]))
}
```

```
> temp <- c(10, 15, 13, 17, 20)
> pressure <- c(900, 920, 890, 940, 920)
> udcorr(temp,pressure)
[1] 0.75
```

# AGENDA

**01** Conditions and Loops

---

**02** Functions

---

**03** R Graphics

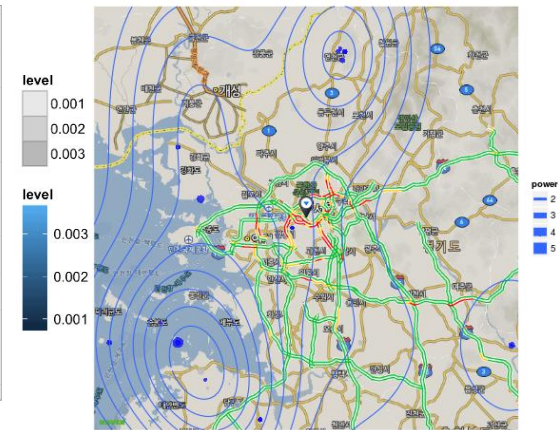
---

**04** Documentation

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# R Graphics

- What we can do with R graphics



# R Graphics

- Basic functions that are provided by “graphics” package

Graphics package function	Description
barplot	Bar and column charts
dotchart	Cleveland dot plots
hist	Histograms
density	Kernel density plots
stripchart	Strip charts
qqnorm (in stats package)	Quantile-quantile plots
xplot	Scatter plots
smoothScatter	Smooth scatter plots
qqplot (in stats package)	Quantile-quantile plots
pairs	Scatter plot matrices
image	Image plots
contour	Contour plots
persp	Perspective charts of three-dimensional data
interaction.plot	Summary of the response for two-way combinations of factors
sunflowerplot	Sunflower plots

# R Graphics

- Fisher's Iris dataset (default dataset provided by R)
  - ✓ Five variables
    - sepal length in cm, sepal width in cm, petal length in cm, petal width in cm, and
    - Species : Iris Setosa, Iris Versicolour, and Iris Virginica.



Setosa




Versicolor

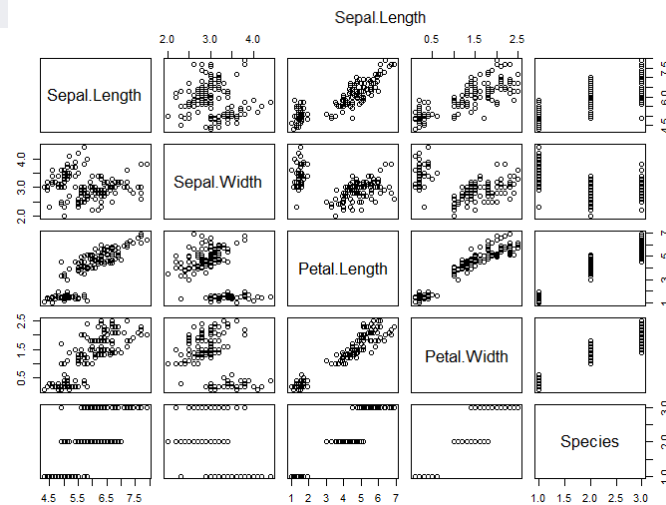
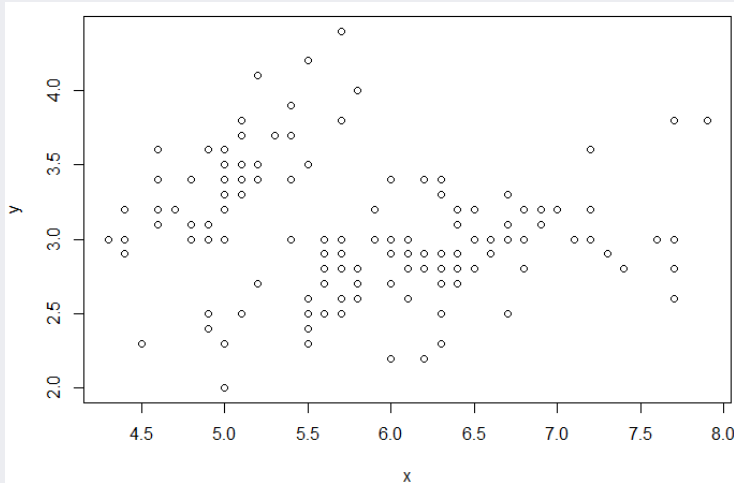
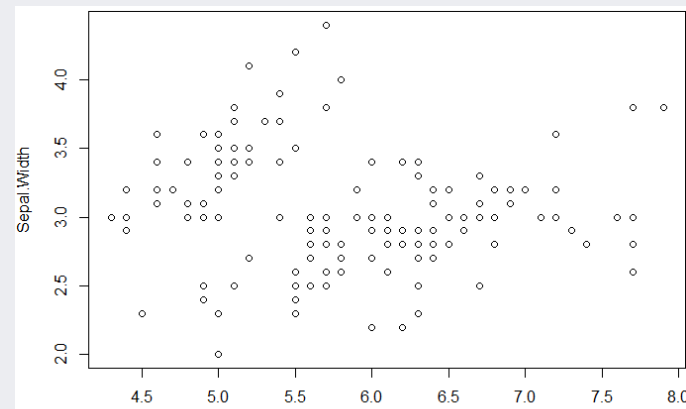


Virginica

# R Graphics

- Polymorphism of R graph functions
  - ✓ polymorphic function: has different operations for different arguments
  - ✓ ex: plot( )

```
Console ~/   
> data(iris)  
> x <- iris[,1]  
> y <- iris[,2]  
> subiris <- iris[,1:2]  
> plot(x,y)  
> plot(subiris)  
> plot(iris)
```

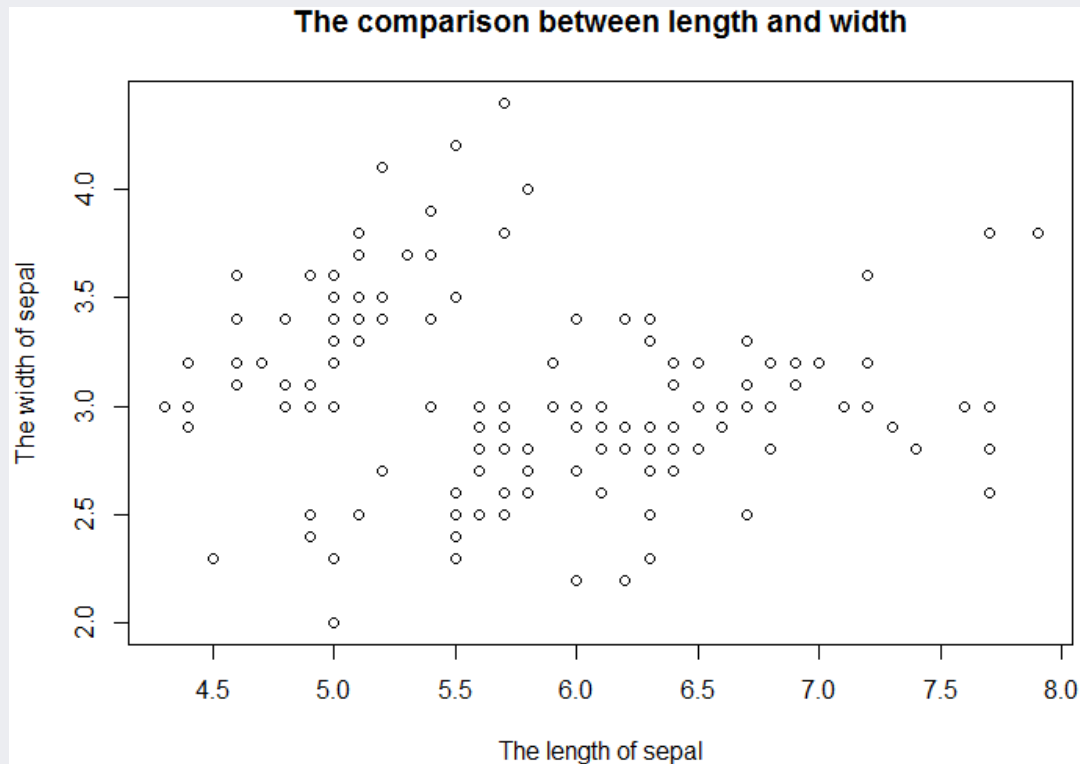


# R Graphics

- Titles and labels in a graph
  - ✓ title: main, x-axis label: xlab, y-axis label: ylab

Console ~/ ↻


```
> plot(subiris, main="The comparison between length and width",  
+ xlab = "The length of sepal",  
+ ylab = "The width of sepal")
```

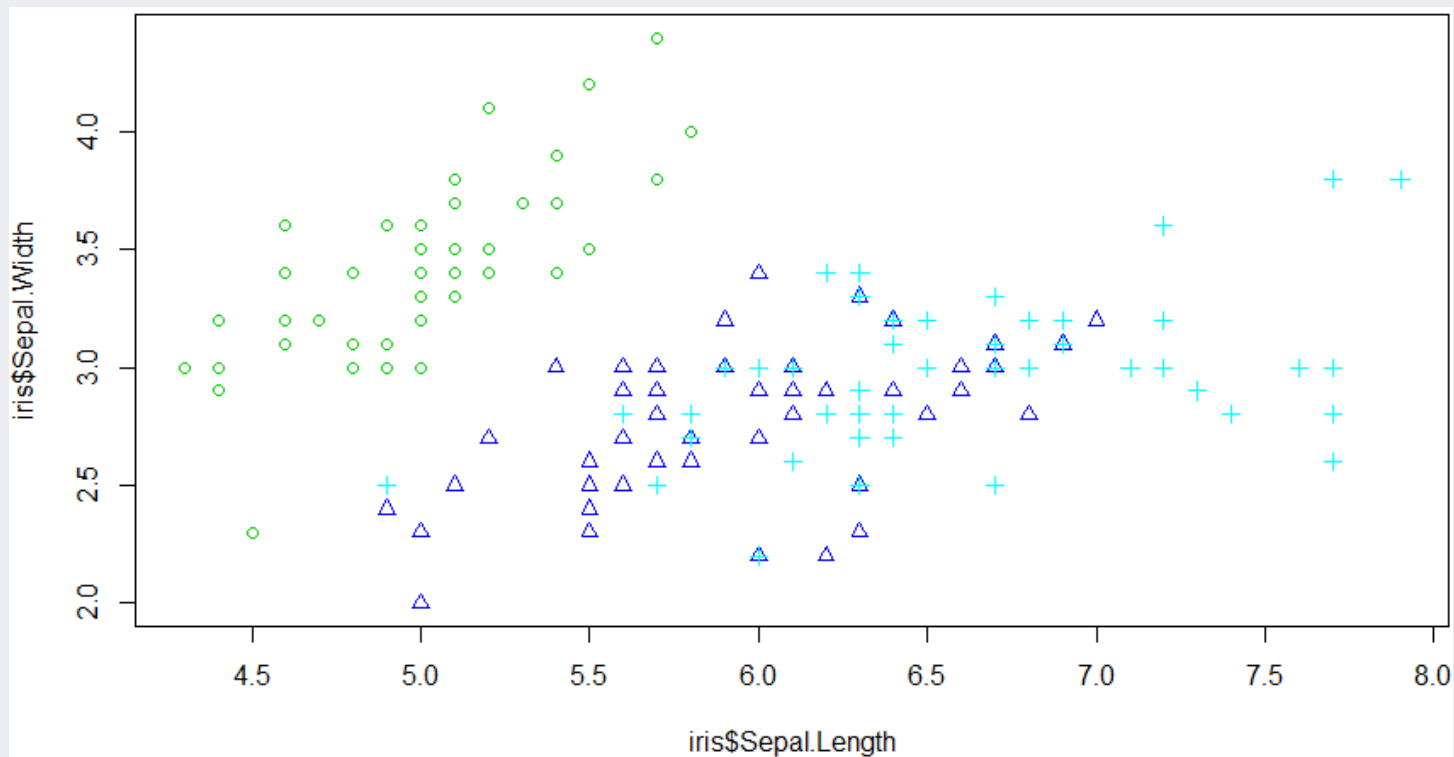




# R Graphics

- Scatter plot for various categories
  - ✓ Use `pch/col` arguments in `plot()`

```
Console ~/   
> plot(iris$Sepal.Length, iris$Sepal.Width,  
+ pch=as.integer(iris$Species), col=as.integer(iris$Species)+10)
```



# R Graphics

- Options for better readability

## A: Plot symbols and text; specify colors and/or character expansion; draw rectangle

```
par(fig=c(0, 1, 0.415, 1))
```

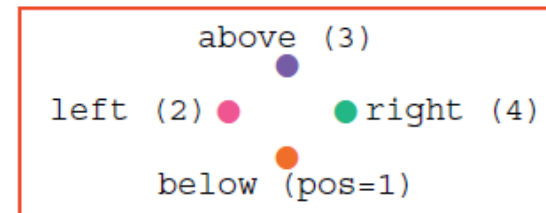
```
plot(0, 0, xlim=c(0, 13), ylim=c(0, 19), type="n")
xpos <- rep((0:12)+0.5, 2); ypos <- rep(c(14.5,12.75), c(13,13))
points(xpos, ypos, cex=2.5, col=1:26, pch=0:25)
text(xpos, ypos, labels=paste(0:25), cex=0.75)
```



```
## Plot characters, vary cex (expansion)
text((0:4)+0.5, rep(9*ht, 5), letters[1:5], cex=c(2.5,2,1,1.5,2))
```

a b c d e

```
## Position label with respect to point
xmid <- 10.5; xoff <- c(0, -0.5, 0, 0.5)
ymid <- 5.8; yoff <- c(-1,0,1,0)
col4 <- colors()[c(52, 116, 547, 610)]
points(xmid+xoff, ymid+yoff, pch=16, cex=1.5, col=col4)
posText <- c("below (pos=1)", "left (2)", "above (3)", "right (4)")
text(xmid+xoff, ymid+yoff, posText, pos=1:4)
rect(xmid-2.3, ymid-2.3, xmid+2.3, ymid+2.3, border="red")
```

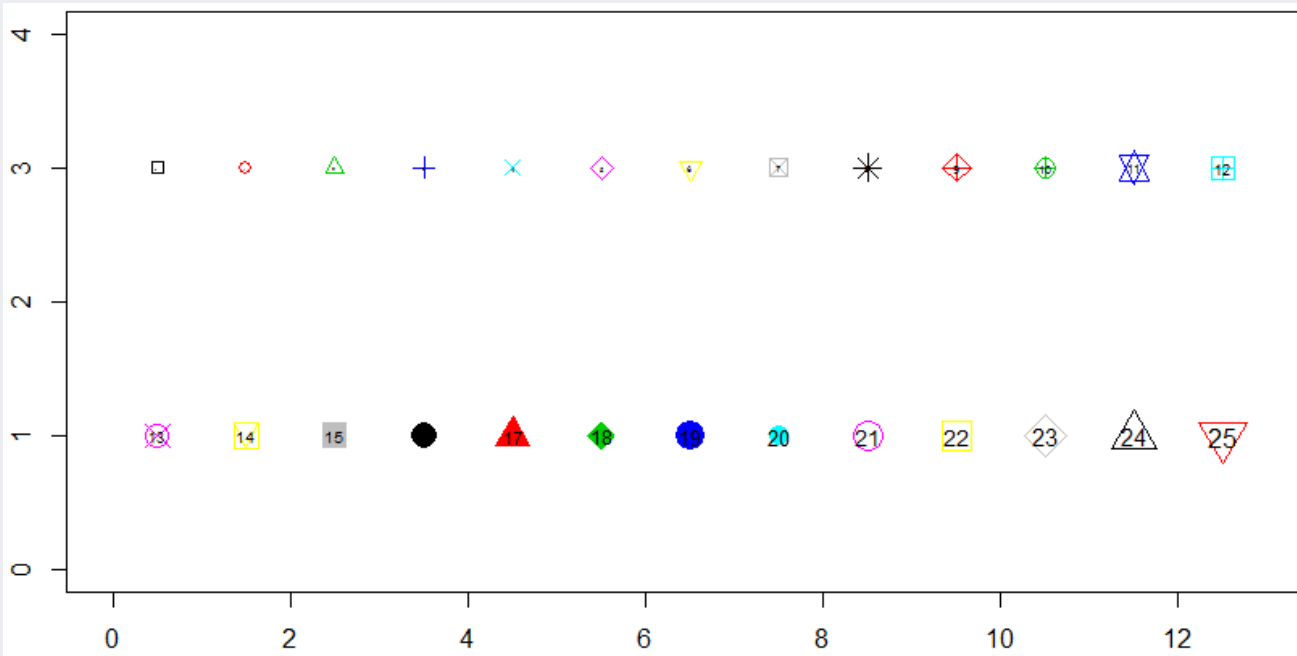


# R Graphics

- Symbols in graphs

✓ pch: shape, cex: size, col: color

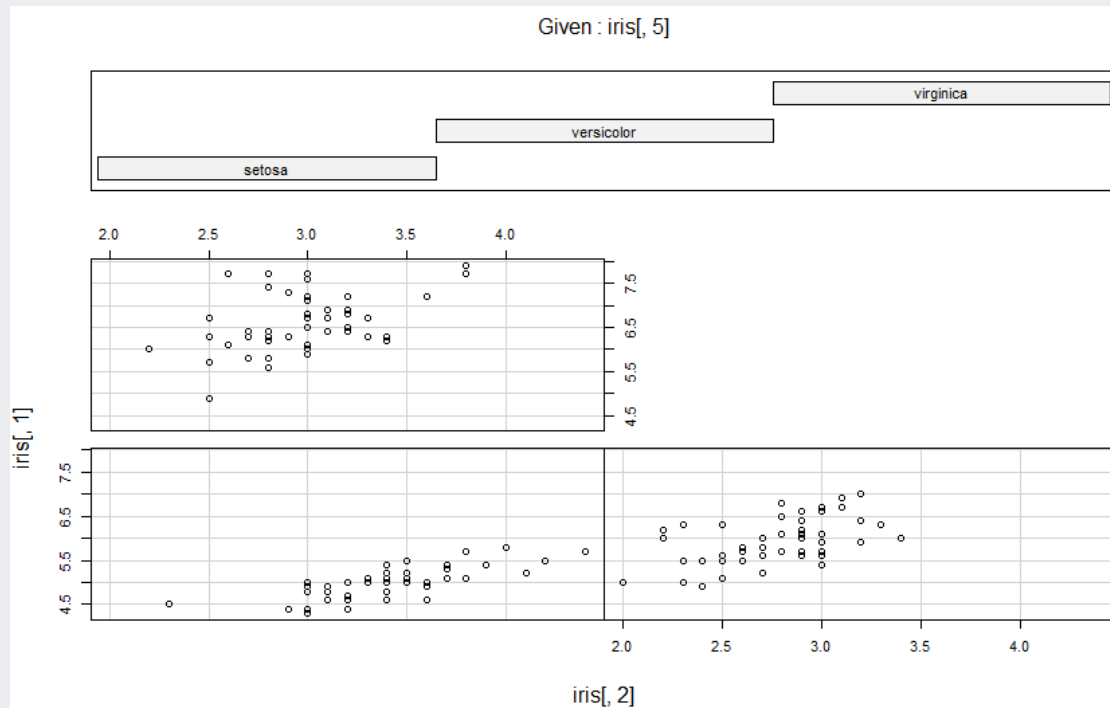
```
> # 사용 가능한 색 및 모양 예시  
> plot(0,0, xlim=c(0,13), ylim=c(0,4), type="n")  
> xpos <- rep((0:12)+0.5,2)  
> ypos <- rep(c(3,1), c(13,13))  
> points(xpos, ypos, cex=seq(from=1,to=3,length=26), col=1:26, pch=0:25)  
> text(xpos, ypos, labels = paste(0:25), cex=seq(from=0.1,to=1,length=26))
```



# R Graphics

- Conditional plot
  - ✓ `coplot(y ~ x | f)`
  - ✓ For every `f` values, draw scatter plot for `x` and `y`

```
> # 조건화 그래프  
> coplot(iris[,1]~iris[,2] | iris[,5])
```

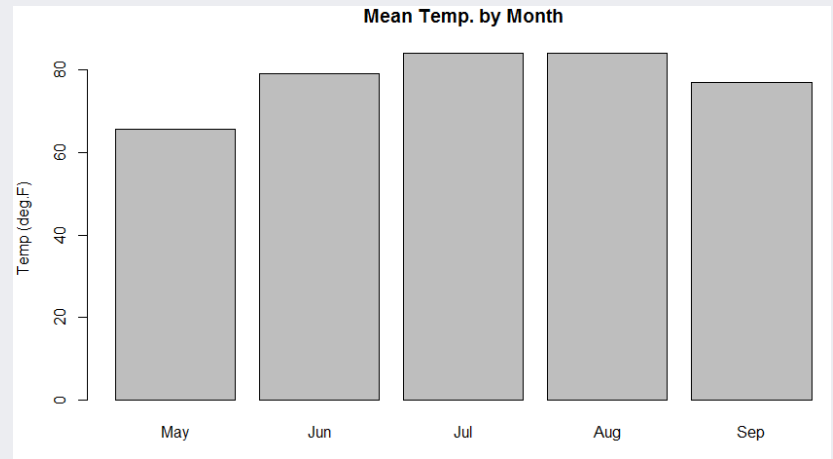
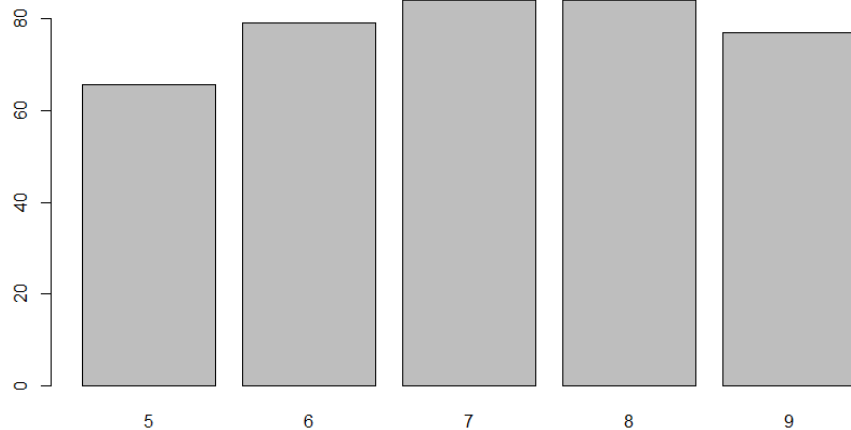


# R Graphics

- Bar plots

Console

```
> data(airquality)
> heights <- tapply(airquality$Temp, airquality$Month, mean)
> barplot(heights)
> barplot(heights, main="Mean Temp. by Month",
+ names.arg = c("May", "Jun", "Jul", "Aug", "Sep"),
+ ylab = "Temp (deg.F)")
```

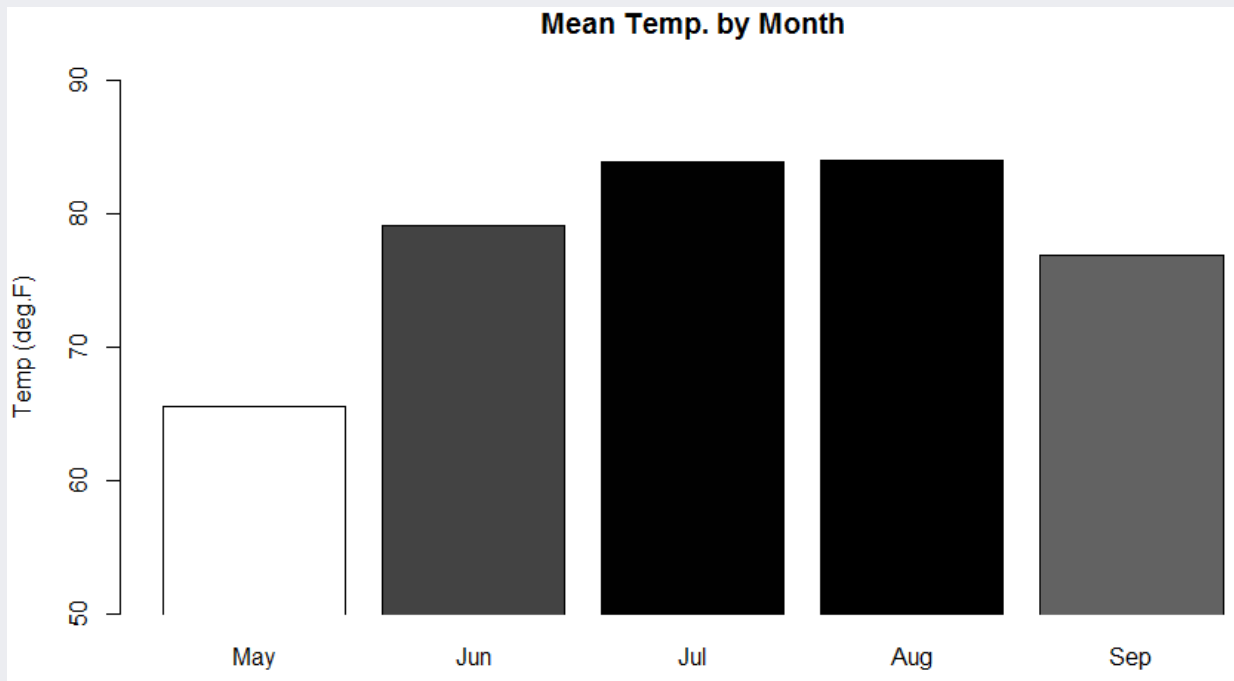


# R Graphics

- Bar plots
  - ✓ For better understanding

Console ~/

```
> rel.hts <- (heights-min(heights))/(max(heights)-min(heights))  
> grays <- gray(1-rel.hts)  
> barplot(heights, col=grays, ylim=c(50,90), xpd=FALSE,  
+ main="Mean Temp. by Month",  
+ names.arg = c("May", "Jun", "Jul", "Aug", "Sep"),  
+ ylab = "Temp (deg.F)")
```



# R Graphics

- Histogram

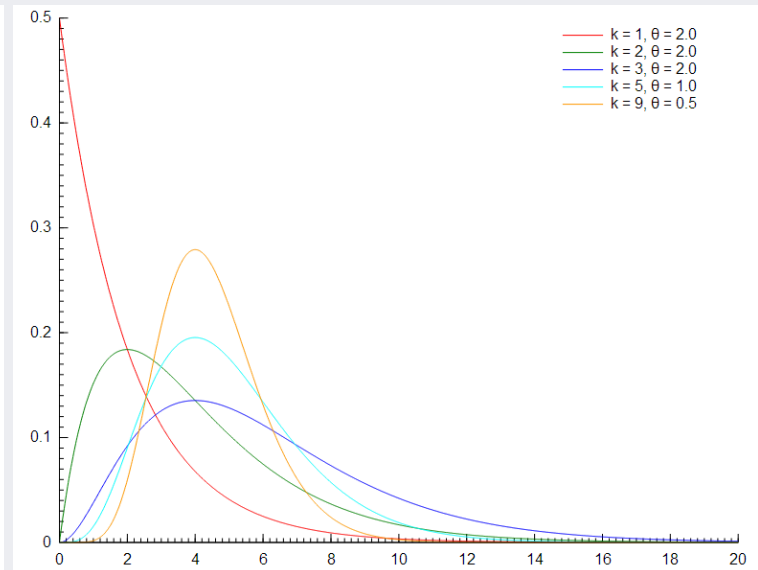
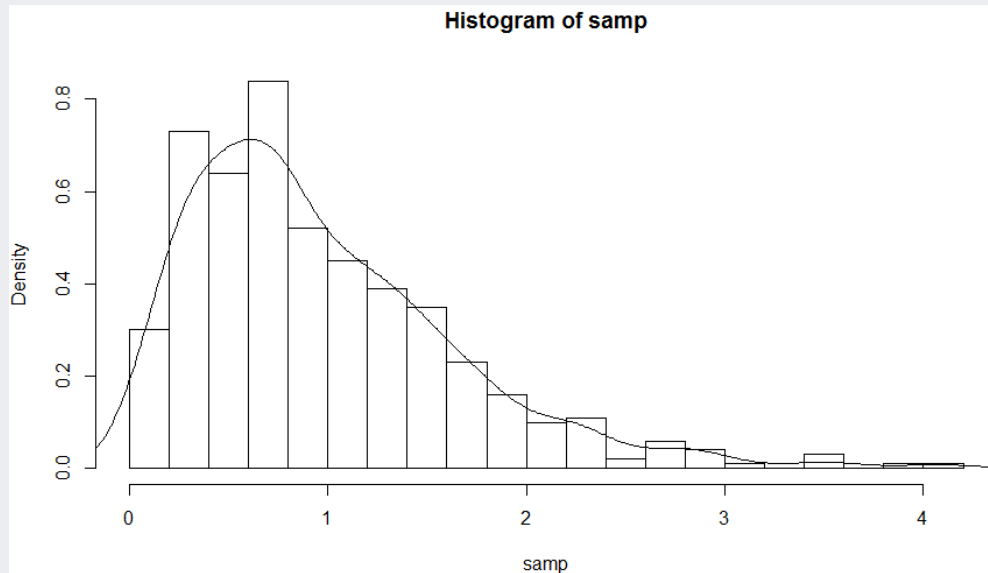
✓ Draw a histogram and an estimated distribution

Console ~/ ↻

```
> samp <- rgamma(500,2,2)
> hist(samp, 20, prob=T)
> lines(density(samp))
```

$$f(x; k, \theta) = x^{k-1} \frac{e^{-x/\theta}}{\theta^k \Gamma(k)} \text{ for } x > 0$$

$$\Gamma(z) = \int_0^{\infty} t^{z-1} e^{-t} dt$$





# R Graphics

- Save graph object
  - ✓ Use file formats that support help(Device) in Windows

```
# 히스토그램 그리기
samp <- rgamma(500,2,2)
hist(samp, 20, prob=T)
lines(density(samp))

# 그림을 파일로 저장하기: png 형식
png("Hist_dist.png")
hist(samp, 20, prob=T)
lines(density(samp))
dev.off()

# 그림을 파일로 저장하기: pdf 형식
pdf("Hist_dist.pdf")
hist(samp, 20, prob=T)
lines(density(samp))
dev.off()
```

 Hist_dist	2016-01-14 오후...	Adobe Acrobat D...	8KB
 Hist_dist	2016-01-14 오후...	PNG 파일	4KB



# R Graphics

- ggplot2: Make R graph more beautiful and informative!

✓ <http://ggplot2.org/>

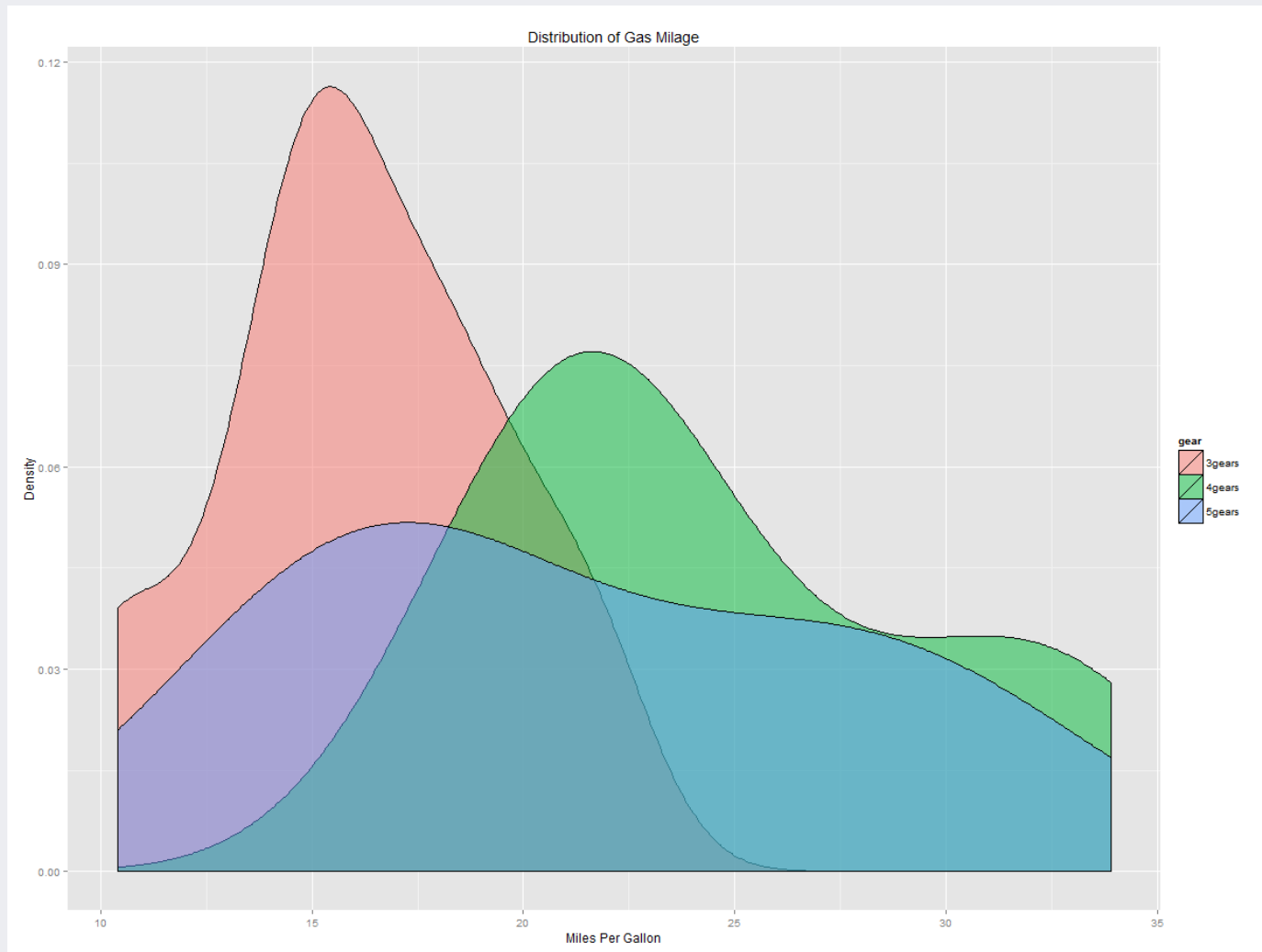
```
# 보다 다양한 그래프 생성을 위해 ggplot2 패키지 이용
install.packages("ggplot2")
library(ggplot2)
data(mtcars)

# 그래프 도서를 위한 팩터 생성
mtcars$gear <- factor(mtcars$gear, levels=c(3,4,5), labels=c("3gears", "4gears", "5gears"))
mtcars$am <- factor(mtcars$am, levels=c(0,1), labels=c("Automatic", "Manual"))
mtcars$cyl <- factor(mtcars$cyl, levels=c(4,6,8), labels=c("4cyl", "6cyl", "8cyl"))

# 연비(mpg)에 대한 커널 밀도 함수 추정
# 기어의 숫자에 따른 그룹(색상)별로 도시
qplot(mpg, data=mtcars, geom="density", fill=gear,
      alpha=I(.5), main="Distribution of Gas Milage",
      xlab="Miles Per Gallon", ylab="Density")
```

# R Graphics

- ggplot2

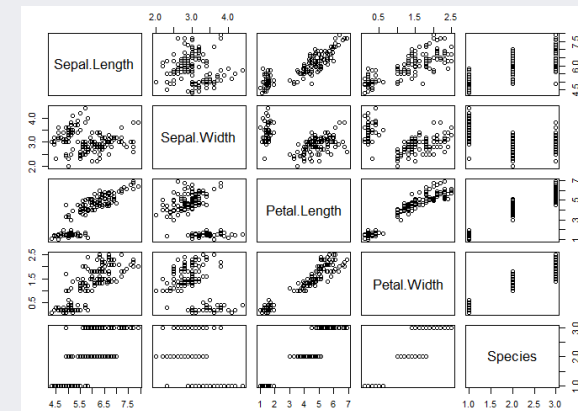
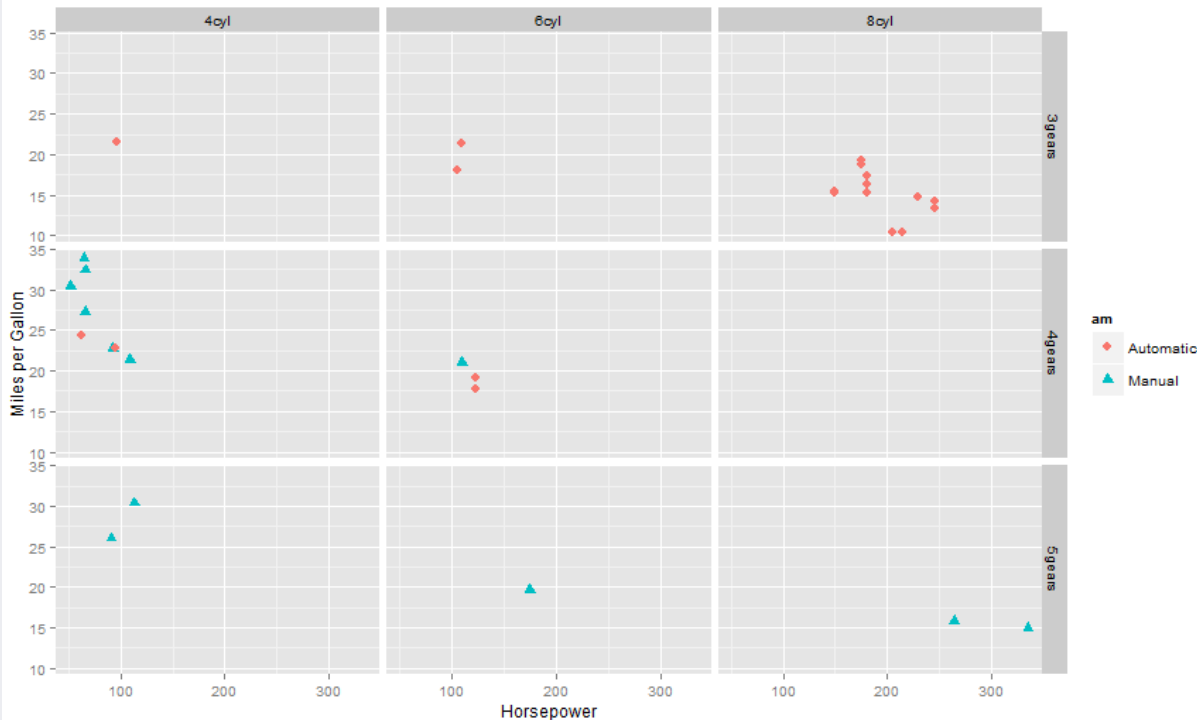


# R Graphics

- ggplot2

- ✓ Scatter plot with ggplot2 vs graphics

```
# 각 기어(gear)-실린더 조합에 따른 연비(mpg)와 마력(hp)의 산점도  
# 각 산점도에서 변속기(am)는 색상과 모양으로 구분됨  
qplot(hp, mpg, data=mtcars, shape=am, color=am,  
       facets=gear~cyl, size=I(3),  
       xlab="Horsepower", ylab="Miles per Gallon")
```

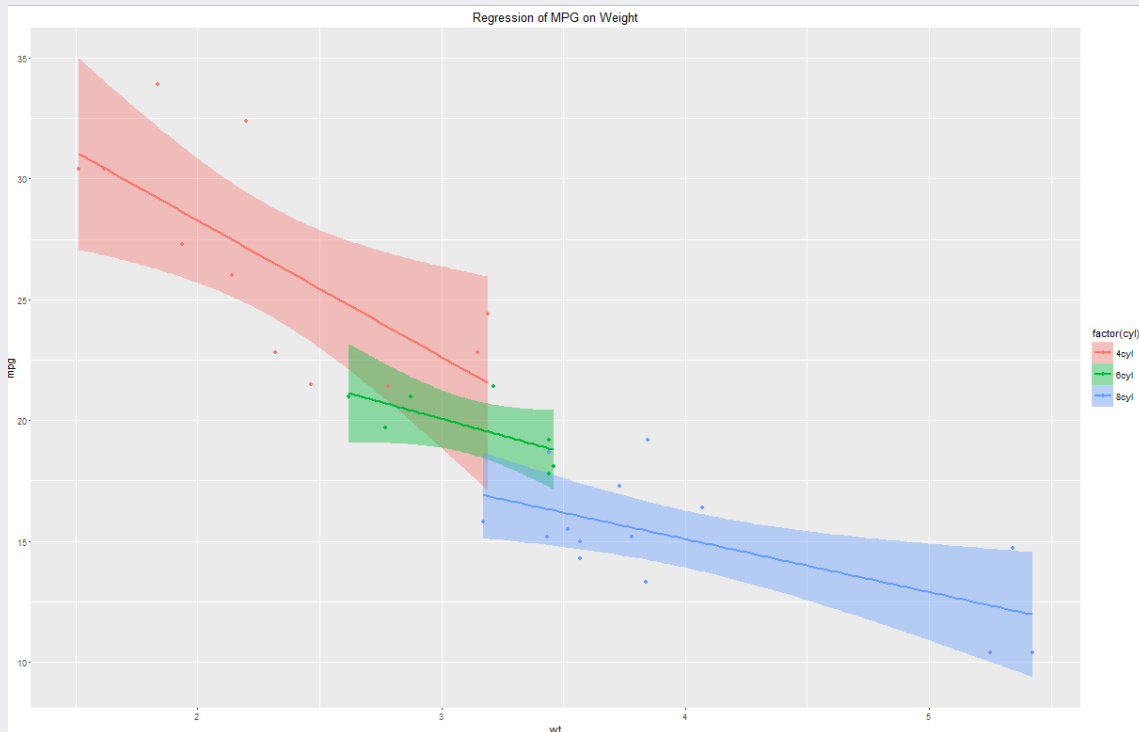


# R Graphics

- ggplot2

✓ Can use various conditions and options

```
# 실린더 갯수에 따라 공차중량(wt)과 연비(mpg)를 회귀선으로 표현  
p <- ggplot(mtcars, aes(y=mpg, x=wt, colour=factor(cyl)))  
p <- p + ggtitle("Regression of MPG on Weight")  
p <- p + stat_smooth(method=lm, aes(fill = factor(cyl))) + geom_point()  
p
```

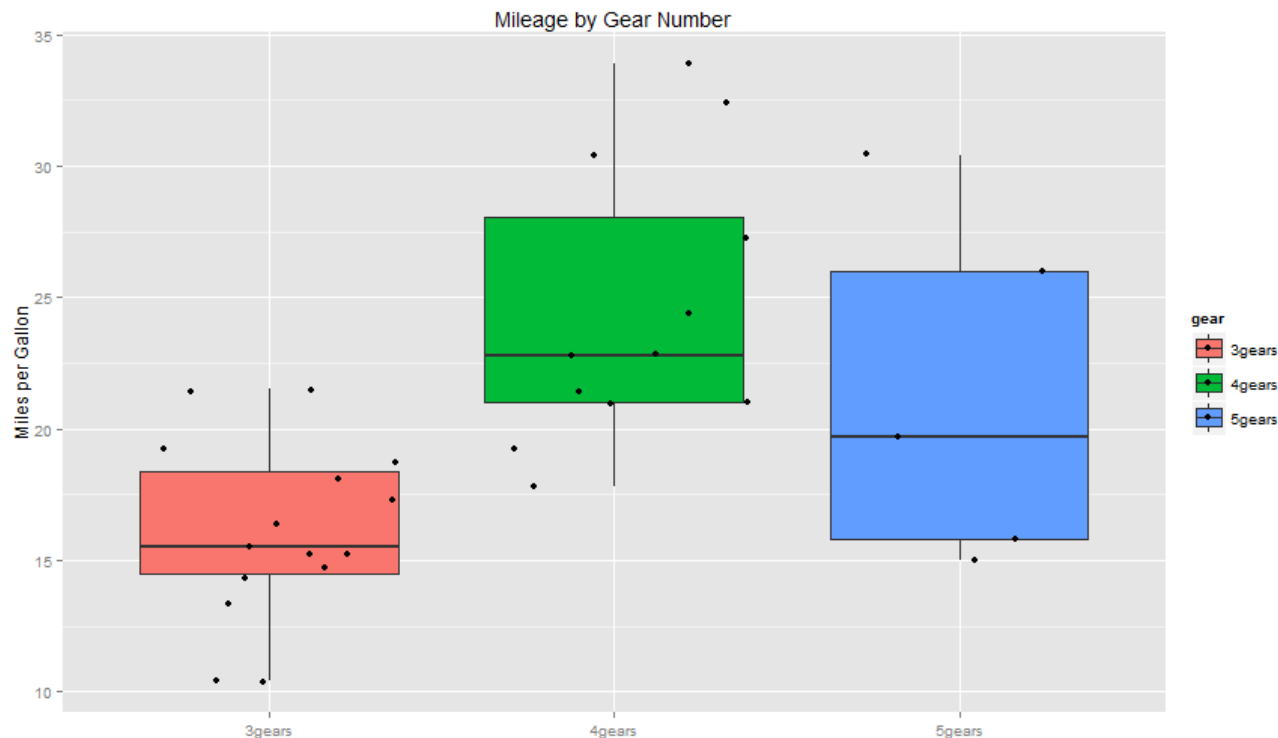


# R Graphics

- ggplot2

✓ Can use various conditions and options

```
# 기어의 숫자에 따른 연비의 상자그림  
# 실제 관측치들을 점으로 표현  
qplot(gear, mpg, data=mtcars, geom=c("boxplot", "jitter"),  
      fill=gear, main="Mileage by Gear Number",  
      xlab="", ylab="Miles per Gallon")
```



# AGENDA

**01** Conditions and Loops

---

**02** Functions

---

**03** R Graphics

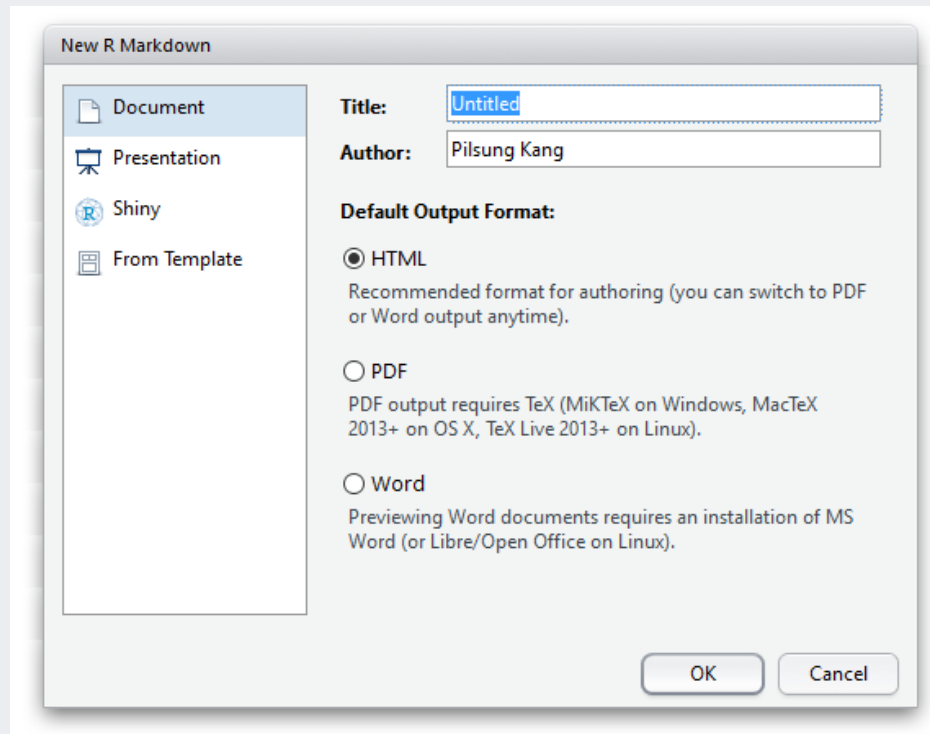
---

**04** Documentation

---

# Documentation

- Documentation using R Markdown
  - ✓ install “knitr” package
- Create an RMD file
  - ✓ File → New File → R Markdown



# Documentation

- Meta information of the file

- ✓ Title, author, date, output file format, etc.

```
1 ▸ ---  
2 title: "ggplot2_examples"  
3 author: "Pilsung Kang"  
4 date: "2016년 1월 14일"  
5 output: html_document  
6 ▸ ---
```

- ✓ How to embed R script in rmd file

- ✓ ```{r, eval = T, echo = T}

- eval = T: run R script, F: do not run R script
- echo = T: print the script in the output file, F: do not print the script

```
```{r, eval = T, echo = T}  
library(ggplot2)  
data(mtcars)  
head(mtcars)  
```
```



# Documentation

- Text in rmd file

- ✓ Parts without ``` are printed as they are

- ✓ Use markdown syntax

- Ex: `**pilsung kang**` becomes **pilsung kang** in the output file

```
30 기어 단계에 따라서 **연비의 분포(miles per gallon)**를 도시합니다.  
31  
32 ~~~{r, eval = T, echo = T}  
33 # Kernel density plots for mpg  
34 # grouped by number of gears (indicated by color)  
35 qplot(mpg, data=mtcars, geom="density", fill=gear,  
36       alpha=I(.5), main="Distribution of Gas Milage",  
37       xlab="Miles Per Gallon", ylab="Density")  
38 ~~~
```

- For more information about markdown syntax

- <https://gist.github.com/ihoneymon/652be052a0727ad59601>
      - <http://blog.kalkin7.com/2014/02/05/wordpress-markdown-quick-reference-for-koreans/>

# Documentation

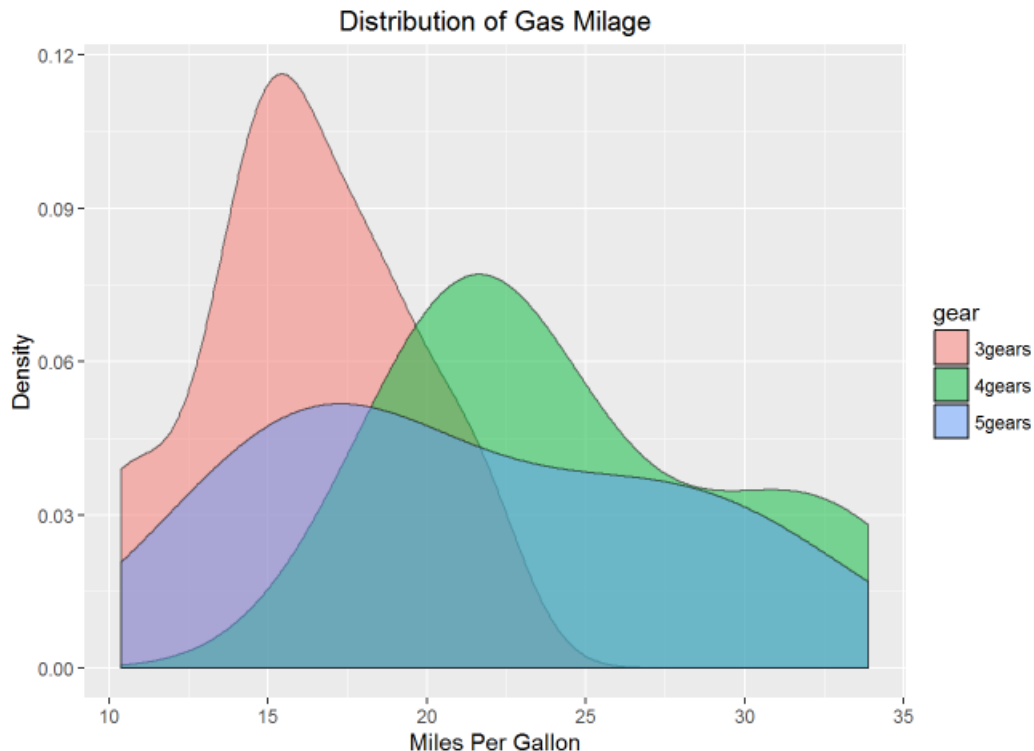
- Output style

- ✓ Differences with eval and echo options

기어 단계에 따라서 연비의 분포(miles per gallon)를 도시합니다.

```
# Kernel density plots for mpg
# grouped by number of gears (indicated by color)
qplot(mpg, data=mtcars, geom="density", fill=gear,
      alpha=I(.5), main="Distribution of Gas Milage",
      xlab="Miles Per Gallon", ylab="Density")
```

**echo = T**  
Print R script



**eval = T**  
Print the result of running R script

# Documentation

- Output style

- ✓ Differences with eval and echo options

기어 단계에 따라서 연비의 분포(miles per gallon)를 도시합니다.

```
# Kernel density plots for mpg
# grouped by number of gears (indicated by color)
qplot(mpg, data=mtcars, geom="density", fill=gear,
      alpha=I(.5), main="Distribution of Gas Milage",
      xlab="Miles Per Gallon", ylab="Density")
```

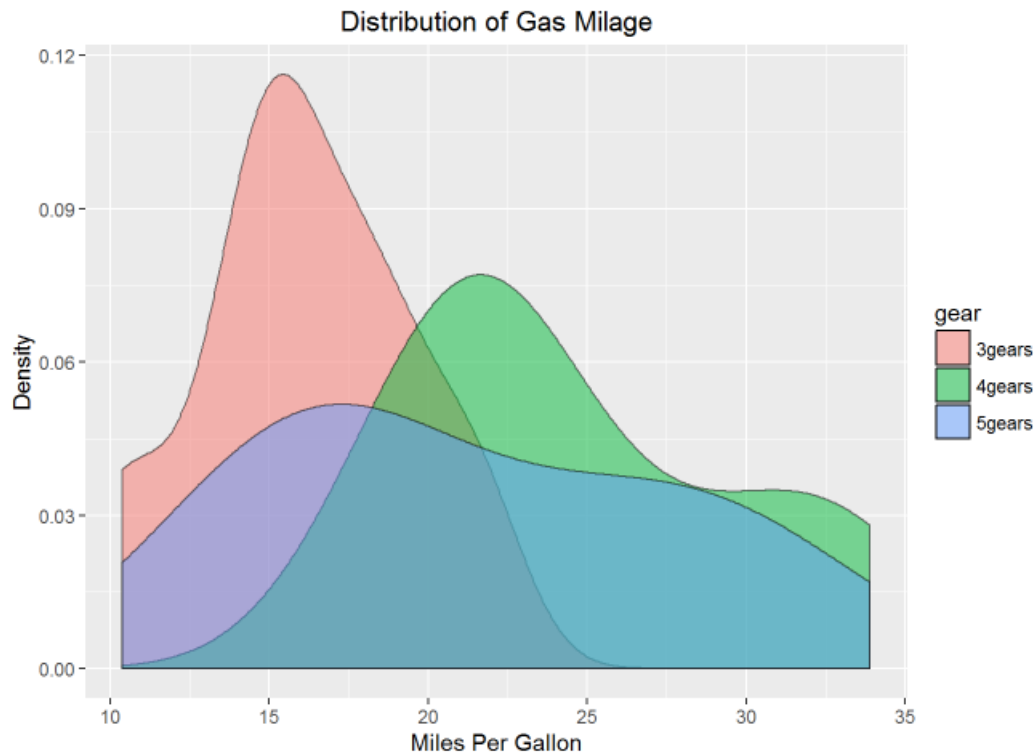
**echo = T**  
Print R script

**eval = F**  
Nothing appears because R script is not run

# Documentation

- Output style
  - ✓ Differences with eval and echo options

기어 단계에 따라서 연비의 분포(miles per gallon)를 도시합니다.



**echo = F**  
Do not print R script

**eval = T**  
Print the result of running R script

# Documentation

- Output style

✓ save html file format

## ggplot2\_examples

Pilsung Kang

2016년 1월 14 일

R 마크다운을 이용하여 R 스크립트와 결과물을 HTML 파일로 불러오는 예제.

ggplot2 패키지를 호출하고 R에 내장된 mtcars 데이터를 불러옵니다.

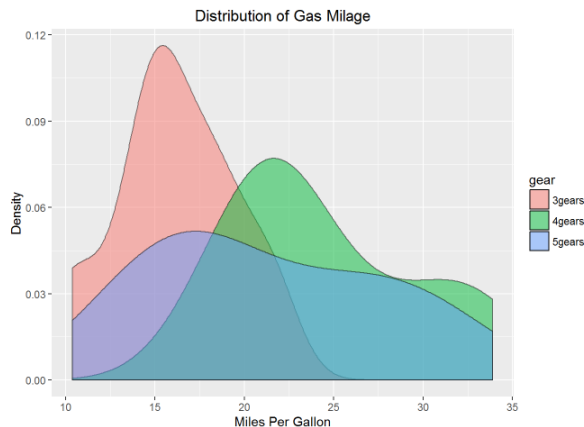
```
library(ggplot2)
data(mtcars)
head(mtcars)
```

```
##           mpg cyl  disp  hp  drat   wt  qsec vs am gear carb
## Mazda RX4      21.0   6  160 110 3.90 2.620 16.46 0 1   4   4
## Mazda RX4 Wag  21.0   6  160 110 3.90 2.875 17.02 0 1   4   4
## Datsun 710      22.8   4  108  93 3.85 2.320 18.61 1 1   4   1
## Hornet 4 Drive  21.4   6  258 110 3.08 3.215 19.44 1 0   3   1
## Hornet Sportabout 18.7   8  360 175 3.15 3.440 17.02 0 0   3   2
## Valiant         18.1   6  225 105 2.76 3.460 20.22 1 0   3   1
```

mtcars의 세 가지 변수에 대해서 팩터 형태로 변환을 하고 레이블을 부여합니다. 1. 기어 단계: 3단계, 4단계, 5단계 2. 변속기 종류: 자동, 수동 3. 엔진 실린더 수: 4개, 5개, 6개

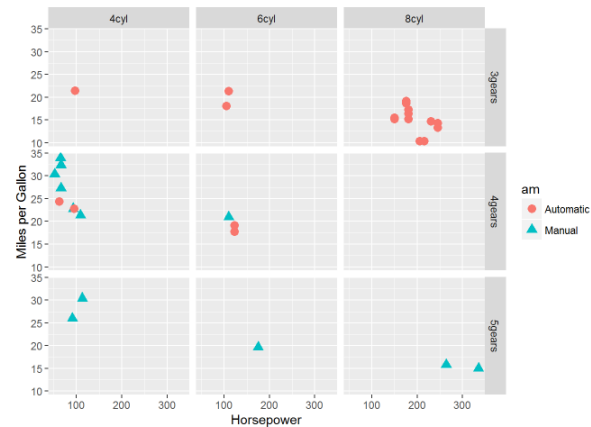
```
# create factors with value labels
mtcars$gear <- factor(mtcars$gear, levels=c(3,4,5), labels=c("3gears", "4gears", "5gears"))
mtcars$am <- factor(mtcars$am, levels=c(0,1), labels=c("Automatic", "Manual"))
mtcars$cyl <- factor(mtcars$cyl, levels=c(4,6,8), labels=c("4cyl", "6cyl", "8cyl"))
```

기어 단계에 따라서 연비의 분포(miles per gallon)를 도시합니다.



변속기 종류(자동, 수동) 및 엔진 실린더 수(4개, 5개, 6개)에 따라 마력(Horsepower)과 연비(Miles per Gallon)를 산점도로 나타냅니다.

```
# Scatterplot of mpg vs. hp for each combination of gears and cylinders
# in each facet, transmission type is represented by shape and color
ggplot(hp, mpg, data=mtcars, shape=am, color=am,
       faceta=gear~cyl, size=I(3),
       xlab="Horsepower", ylab="Miles per Gallon")
```



실린더의 수별로 공차중량(wt)과 연비(mpg)사이의 선형 식을 도출합니다. 추정된 연비의 95% 신뢰구간을 음영으로 표시합니다.

```
# Separate regressions of mpg on weight for each number of cylinders
p <- ggplot(mtcars, aes(y=mpg, x=wt, colour=factor(cyl)))
p <- p + ggtitle("Regression of MPG on Weight")
p <- p + stat_smooth(method=lm, aes(fill = factor(cyl))) + geom_point()
p
```



# R Markdown Cheat Sheet

## R Markdown Cheat Sheet

learn more at [rmarkdown.rstudio.com](http://rmarkdown.rstudio.com)



### .Rmd files

An R Markdown (.Rmd) file is a record of your research. It contains the code that a scientist needs to reproduce your work along with the narration that a reader needs to understand your work.



### Reproducible Research

At the click of a button, or the type of a command, you can rerun the code in an R Markdown file to reproduce your work and export the results as a finished report.



### Dynamic Documents

You can choose to export the finished report as a html, pdf, MS Word, ODT, RTF, or markdown document; or as a html or pdf based slide show.

### Workflow

**1 Open a new .Rmd file** at File ► New File ► R Markdown. Use the wizard that opens to pre-populate the file with a template

**2 Write document** by editing template

**3 Knit document to create report** Use knit button or `render()` to knit

**4 Preview Output** in IDE window

**5 Publish** (optional) to web or server

### Interactive Documents

Turn your report into an interactive Shiny document in 4 steps



- 1 Add runtime: shiny to the YAML header.
- 2 Call Shiny input functions to embed input objects.
- 3 Call Shiny render functions to embed reactive output.
- 4 Render with `rmarkdown::run` or click Run Document in RStudio IDE

### .Rmd structure

**YAML Header**  
Optional section of render (e.g. pandoc) options written as key:value pairs (YAML).

- At start of file
- Between lines of ---

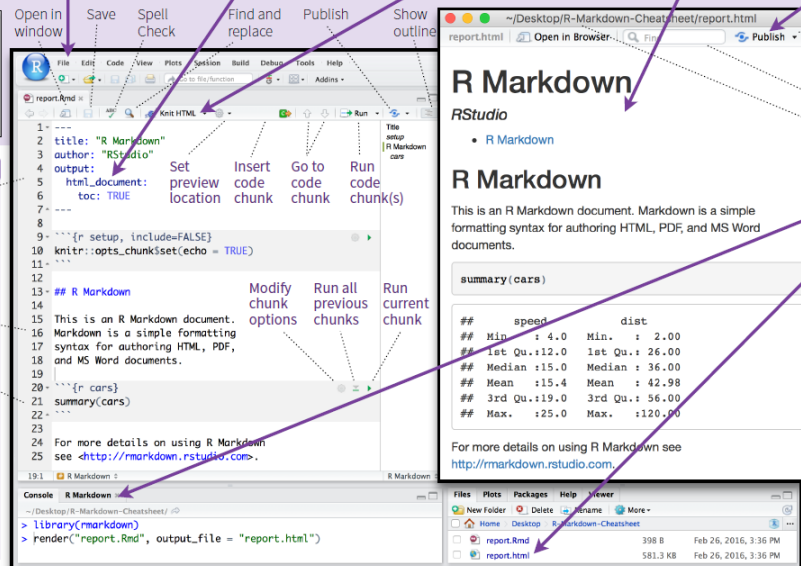
**Text**  
Narration formatted with markdown, mixed with:

**Code chunks**  
Chunks of embedded code. Each chunk:

- Begins with ````{r}`
- ends with `````

R Markdown will run the code and append the results to the doc.

It will use the location of the .Rmd file as the working directory



### render()

Use `rmarkdown::render()` to render/knit at cmd line. Important args:

- input** - file to render
- output\_format** - render output format
- output\_options** - List of render options (as in YAML)
- output\_file** - output file
- output\_dir** - output directory
- params** - list of params to use
- envir** - environment to evaluate code chunks in
- encoding** - of input file

```
---
output: html_document
runtime: shiny
---
```{r, echo = FALSE}
numericInput("n",
  "How many cars?", 5)
renderTable({
  head(cars, input$n)
})
```
```

| 5      |       |  |
|--------|-------|--|
| speed  | dist  |  |
| 1 4.00 | 2.00  |  |
| 2 4.00 | 10.00 |  |
| 3 7.00 | 4.00  |  |
| 4 7.00 | 22.00 |  |
| 5 8.00 | 16.00 |  |

Embed a complete app into your document with `shiny::shinyAppDir()`

\* Your report will be rendered as a Shiny app, which means you must choose an html output format, like `html_document`, and serve it with an active R Session.

### Embed code with knitr syntax

#### Inline code

Insert with ``r <code>``. Results appear as text without code.

Built with `r getRVersion()` → Built with 3.2.3

#### Code chunks

One or more lines surrounded with ````{r}` and `````. Place chunk options within curly braces, after `r`. Insert with ````{r options}`

```
```{r echo=TRUE}
getRVersion()
```
```

#### Global options

Set with `knitr::opts_chunk$set()`, e.g.

```
```{r include=FALSE}
knitr::opts_chunk$set(echo = TRUE)
```
```

#### Important chunk options

- cache** - cache results for future knits (default = FALSE)
- cache.path** - directory to save cached results in (default = "cache/")
- child** - file(s) to knit and then include (default = NULL)
- collapse** - collapse all output into single block (default = FALSE)
- comment** - prefix for each line of results (default = "##")

- dependson** - chunk dependencies for caching (default = NULL)
- echo** - Display code in output document (default = TRUE)
- engine** - code language used in chunk (default = 'R')
- error** - Display error messages in doc (TRUE) or stop render when errors occur (FALSE) (default = FALSE)
- eval** - Run code in chunk (default = TRUE)

- fig.align** - 'left', 'right', or 'center' (default = 'default')
- fig.cap** - figure caption as character string (default = NULL)
- fig.height**, **fig.width** - Dimensions of plots in inches
- highlight** - highlight source code (default = TRUE)
- include** - Include chunk in doc after running (default = TRUE)

- message** - display code messages in document (default = TRUE)
- results** (default = 'markup')  
'asis' - passthrough results  
'hide' - do not display results  
'hold' - put all results below all code
- tidy** - tidy code for display (default = FALSE)
- warning** - display code warnings in document (default = TRUE)

Options not listed above: R.options, aniopts, autodep, background, cache.comments, cache.lazy, cache.rebuild, cache.vars, dev, dev.args, dpi, engine.opts, engine.path, fig.asp, fig.env, fig.ext, fig.keep, fig.lp, fig.path, fig.pos, fig.process, fig.retina, fig.scap, fig.show, fig.showtext, fig.subcap, interval, out.extra, out.height, out.width, prompt, purr, ref.label, render, size, split, tidy.opts

### Parameters

Parameterize your documents to reuse with different inputs (e.g., data sets, values, etc.)

**1 Add parameters**

Create and set parameters in the header as sub-values of `params`

```
---
params:
  n: 100
  d: lr Sys.Date()
---
```

**2 Call parameters**

Call parameter values in code as `params$name`

```
Today's date
is `r params$d`
```

**3 Set parameters**

Set values with `Knit` with `parameters` or the `params` argument of `render()`

```
render("doc.Rmd",
  params = list(n = 1, d = as.Date("2015-01-01")))
```

# R Markdown Cheat Sheet

## Pandoc's Markdown

Write with syntax on the left to create effect on right (after render)

### Plain text

End a line with two spaces to start a new paragraph.

`*italics*` and `**bold**`

``verbatim code``

`sub/superscript^2^~2~`

`~~strikethrough~~`

escaped: `\* \_ \\`

endash: `--`, emdash: `---`

equation: `$A = \pi * r^2$`

equation block:

`$$E = mc^2$$`

`> block quote`

`# Header1 {#anchor}`

`## Header 2 {#css_id}`

`### Header 3 {#css_class}`

`#### Header 4`

`##### Header 5`

`##### Header 6`

`<!--Text comment-->`

`\textbf{Text ignored in HTML}`

`<em>HTML ignored in pdfs</em>`

`<http://www.rstudio.com>`

`[link](www.rstudio.com)`

`Jump to [Header 1](#anchor)`

`image:`

`![Caption](smallor.png)`

`* unordered list`

`+ sub-item 1`

`+ sub-item 2`

`- sub-sub-item 1`

`* item 2`

`Continued (indent 4 spaces)`

`1. ordered list`

`2. item 2`

`1) sub-item 1`

`A. sub-sub-item 1`

`(@) A list whose numbering`

`continues after`

`(@) an interruption`

`Term 1`

`Definition 1`

`: Definition 1`

`Right Left Default Center`

`12 12 12 12`

`123 123 123 123`

`1 1 1 1`

`12 12 12 12`

`123 123 123 123`

`1 1 1 1`

`12 12 12 12`

`123 123 123 123`

`1 1 1 1`

`12 12 12 12`

`123 123 123 123`

`1 1 1 1`

`12 12 12 12`

`123 123 123 123`

`1 1 1 1`

Plain text

End a line with two spaces to start a new paragraph.

`italics` and `bold`

`verbatim code`

`sub/superscript^2`

`strikethrough`

escaped: `\* \_ \\`

endash: `--`, emdash: `---`

equation: `A = \pi * r^2`

equation block:

`E = mc^2`

`> block quote`

`# Header1 {#anchor}`

`## Header 2 {#css_id}`

`### Header 3 {#css_class}`

`#### Header 4`

`##### Header 5`

`##### Header 6`

`<!--Text comment-->`

`\textbf{Text ignored in HTML}`

`<em>HTML ignored in pdfs</em>`

`<http://www.rstudio.com>`

`[link](www.rstudio.com)`

`Jump to [Header 1](#anchor)`

`image:`

`![Caption](smallor.png)`

`* unordered list`

`+ sub-item 1`

`+ sub-item 2`

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