

Utility Functions	
<code>getwd()</code>	Find the working dir.
<code>setwd("C:/file/path")</code>	Set the working dir.
<code>ls()</code>	List variables
<code>rm(object)</code>	Delete object
<code>str(object)</code>	Displays internal structure of R object
<code>help.start()</code>	Launch help console
<code>install.packages("package_name")</code>	Install package
<code>library(package_name)</code>	Load package
<code>detach(package:-package_name)</code>	Remove package
<code>scan()</code>	Read data values

Lists	
<code>list(x=1:5, y=c('a', 'b'))</code>	Create list
<code>is.list()</code>	<code>as.list()</code>
Check if the arg is a list	Force the arg to list
<code>lapply(list_name, function)</code>	
Apply function over a list and return as list	
<code>sapply(list_name, function)</code>	
Return as suitable data structure(vector)	

Strings	
<code>c("String1","String2")</code>	Create a string vector
<code>toString(x)</code>	Convert to string
<code>noquote(string)</code>	Print string w/o quote
<code>sprintf()</code>	Print text & var values
<code>cat()</code>	Concatenate & print
<code>toupper(string)</code>	Convert to uppercase
<code>tolower(string)</code>	Convert to lowercase
<code>substr(string,n,m)</code>	Extract substrings in a string from n to m
<code>strsplit(string," ")</code>	Split elements of string
<code>paste(c("a","b"),"c")</code>	<code>paste0(c("a","b"),"c")</code>
Concatenate vectors	Concat w/o separator

Probability Distributions	
<code>rbinom(n, size, prob)</code>	Binomial
<code>rpois(n, lambda)</code>	Poisson
<code>runif(n, min=0, max=1)</code>	Uniform
<code>rnorm(n, mean=0, sd=1)</code>	Normal
<code>rexp(n, rate=1)</code>	Exponential

Vectors	
<code>c(2, 4, 6)</code>	Numeric vector
<code>c("one","two","thr")</code>	Character vector
<code>c(TRUE,FALSE)</code>	Logical vector
<code>rep(1:2,times=3)</code>	Repeat a vector
<code>rep(1:2,each=3)</code>	Repeat the elements
<code>which.min()</code>	<code>which.max()</code>
Index of the min	Index of the max

Data Frames	
<code>data.frame(x=1:2, y=c('a', 'b'))</code>	Create data frame
<code>View(df_name)</code>	See full dataframe
<code>head(df_name)</code>	See first 6 rows
<code>tail(df_name)</code>	See last 6 rows
<code>df_name[cond,]</code>	Row filter
<code>df_name[c("column")]</code>	Column filter
<code>df_name[cond1,][,cond2]</code>	Row and Column filter

Functions	
<pre>function_name <- function(var){ Do something return (new_variable)}</pre>	
<code>args(function_name)</code>	Arguments of func
<code>body(function_name)</code>	Body of func



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Flow Control

If Statement -

```
if (condition){
  Do something
} else {
  Do something different}
-----
- -----
```

Ifelse Statement -

```
ifelse (condition, Do
something, Do something
different)
-----
- -----
```

Switch Statement -

```
switch ("beta", " alpha= 1,b -
eta =2, gamma=3,4)
```

Loops

For Loop -

```
for (var in sequence){Do something
}
```

Visualizations

barplot()	plot()	qqnorm()
pie()	plot(density())	qplot()
mosaicplot()	pairs()	boxplot()
hist()	matplot()	ggplot()

Arrays

array(1:24, dim=c(4,3,2), dimnames=.....)
Create array with 4 rows, 3 cols and 2 groups

Matrices

m1 <- matrix(1:12, now=4, ncol=3, dimnames=....)
Create a matrix with 4 rows and 3 columns

t(m) Transpose of matrix

rbind(m1,m2) **cbind(m1,m2)**
Combine by row Combine by column

The following applies to arrays also:

dimnames(m) **dim(m)** **colnames(m)**
rownames(m) **nrow(m)** **ncol(m)**

Descriptive Statistics

summary(object) Summary of object

class(object) Find class of an R object
while (condition){ Do something }

length(object) Get length of an object

quantile(x) Find quantiles

rowMeans(x)/ **rowSums(x)/**
colMeans(x) **colSums(x)**

table(x) Build a contingency table

describe(object) Description of object

subset(x,cond) Create subsets

Hypothesis Testing

t.test(data, mu=3)
One sample two-sided t-test

t.test(data, mu=3, alternative='greater')
One sample one-sided t-test

t.test(data1, data2, mu=0.5)
Two sample two-sided t-test

t.test(data1, data2, mu=0.5, alternative='less')
Two sample one-sided t-test

t.test(post_data, pre_data, paired=TRUE)
Paired test

wilcox.test(data, mu=8, alt='less')
Wilcoxon test

cor.test(data1, data2) **chisq.test(data)**
Correlation test Chi-square test

ks.test(data1, data2) **shapiro.test(data)**
If both are from same Normality test
distn

aov(data1~ data2) **lm(data1~ data2)**
ANOVA Regression



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