统计分析与建模

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R基础语法

Objects

- Vectors
- Arrays
- Matrices
- Lists
- data frames
- Data type and operation
 - Data type
 - Special Constants
 - Operation
- Control flow
 - Conditions
 - Loops
- Function
- I/O

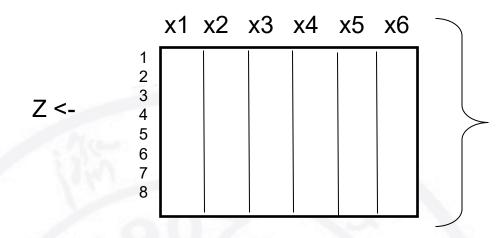


Objects

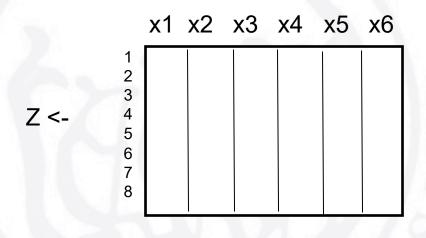
- Almost all things in R functions, datasets, results, etc. are Objects.
 - (graphics are written out and are not stored as objects)

- Objects are classified by two criteria:
 - MODE(...): how objects are stored in R character, numeric, logical, factor, list, & function
 - CLASS(...): how objects are treated by functions (important to know!) [vector], matrix, array, data.frame, & hundreds of special classes created by specific functions

Objects



The MODE of Z is determined **automatically** by the types of things stored in Z – numbers, characters, etc. If it is a mix, mode = list.

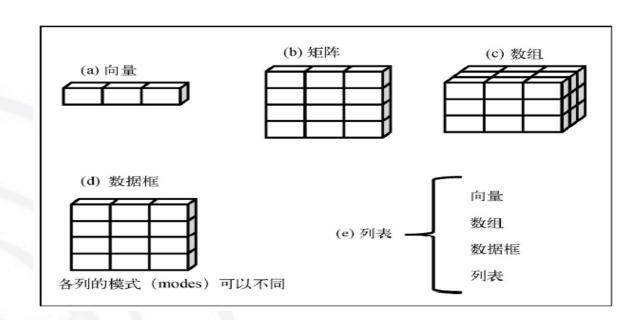


The CLASS of Z is either set by default depending, on how it was created, or is **explicitly set by user**. You can check the objects' class and change it. It determines how functions deal with Z.

```
DEMO
       a=c(1,2,3)
class(a)='small_num_list'
       (mode(a))
        (class(a))
```

Objects in R

- Vectors
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Vectors

- A vector is a single entity consisting of an ordered collection of elements of the same type
- To define a vector use the function c() (concatenate)
- You can include inputs to a vector of different data type however the resulting vector will make them all the same type
- You can transform the vector to any data type as well
 - as.numeric(), as.character()

DEMO

Vector生成

- ①函数c(...)为自定义量
- ②from:to产生一个序列
- ③seq()产生一个等差向量序列:
- seq(from = n, to = m, by = k, len = w)
- ④rep() 重复一个对象: rep(x, times)
- ⑤rnorm()随机产生正态分布向量: rnorm(number, mean, variance)

● Vector访问

- You can access specific elements of a vector
 - (1) 如何排除下标对应的元素?
 - (2) 满足条件的元素? which(...), which.max(...)

```
      > x[3]
      #下标为正数,取出下标对应的元素

      > x[-3]
      #下标为负数,排除下标对应的元素

      > x[3:5]
      #取出连续的元素

      > x[c(3,5,8)]
      #如果一次取出多个元素,需要用向量做下标,注意负号

      > x[which(x>6)]
      #取出满足条件的元素,要使用which()函数

      > x[which.max(x)]
      #取出最大元素,最小元素小标为which.min
```

● Vector更新

• You can change, add, or delete parts to a vector

```
13  x <- seq(1, 3, by=0.2)

14  x

15  x[8] <- 34

16  x

17  x[12] <- 3.2

18  x

19  x <- x[c(-8, -12)]

20  x
```

Vector运算

```
27 a <- c(1, 2, 3, 4, 5)

28 a + 1

29 a^2

30 sqrt(a)

31 sum(a)
```

- (1) 如果向量长度不同? ——v1+v2
- (2) 如何获得向量长度? ——length(v1)
- (3) 如何连接向量? ——c(v1,v2)

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Array生成

Arrays are the R data objects which can store data in two or more dimensions

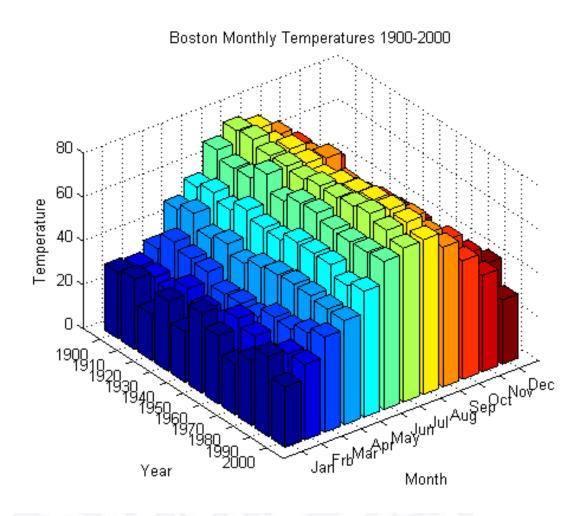
```
> my_array <- array(1:24, dim=c(4,6))
> my_array
    [,1] [,2] [,3] [,4] [,5] [,6]
[1,]
[2,] 2 6 10 14
[3,] 3 7 11 15 19 23
[4,] 4 8 12 16
                       20 24
> my_array <- array(c(1, 4, 8, 10), dim=c(2,2))
> my_array
    [,1] [,2]
[1,]
[2,]
        10
```

Array排列更新

You can always change the dimensions of an existing array

```
[,1] [,2] [,3] [,4] [,5] [,6]
[1,]
                          21
[2,] 2 6 10
                          22
                 14
[3,] 3 7 11 15 19
                         23
[4,] 4 8
             12
                 16
                          24
> dim(my_array) <- c(6,4)</pre>
> my_array
    [,1] [,2] [,3] [,4]
[1,]
[2,] 2 8 14
                20
[3,] 3 9 15 21
[4,] 4
         10 16 22
[5,]
         11
             17
                 23
[6,]
         12
                  24
             18
```

Example of Array Use



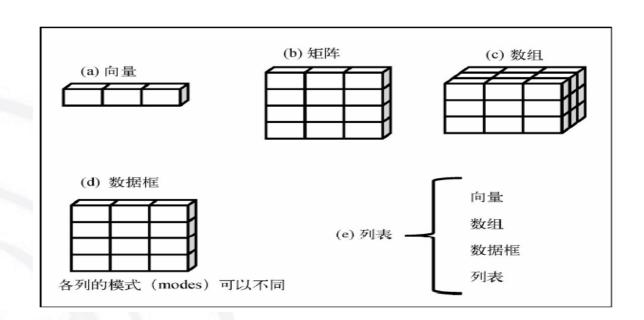
- Array访问
- Array更新
- Array运算

单元素访问 a[i, j, k]

多元素访问 a[c(i1, i2), c(j1, j2), c(k1, k2)]

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Matrice生成

- An array with two dimensions
- You can change the ordering of the matrix to order by row

matrix(data,nrow=,ncol=,byrow=T)

```
# 创建一个 3x3 的矩阵
A <- matrix(1:9, nrow = 3)
B <- matrix(9:1, nrow = 3)

# 矩阵乘法
C <- A %*% B
print(C)
```

Matrice访问

You can select specific elements and subsets of a matrix

方法1: 使用下标和方括号来选择矩阵中的行、列或元素

- y[i,]: 返回矩阵y中的第i行
- y[,j]: 返回第j列
- y[i,j]: 返回第i行第j列元素
- y[i,-j]: 返回第i行,但排除第j列元素
- y[-i, j]: 返回第j行,但排除第i行元素

方法2: 使用向量和方括号来选择矩阵中的行、列或元素

- y[c(1,3),c(2,4)]: 返回第1,3行,第2,4列元素
- y[c(1,3),-c(2,4)]:返回第1,3行,但排除第2,4列元素

Matrice运算

- 横向合并矩阵:cbind();
- 纵向合并矩阵:rbind();
- 转置:t(y);
- · 将矩阵转化为向量: as.vector();
- 返回矩阵维度: dim()、nrow()和ncol();
- 对矩阵各列求和:colSums();
- 求矩阵各列的均值:colMeans();
- 对矩阵各行求和:rowSums();
- 求矩阵各列的均值:rowMeans();
- 计算行列式:det();

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● Data Frame生成

```
Pitchers_Speeds <- data.frame(
Pitchers_Name = c("Nolan Ryan","Bob Feller","Aroldis Chapman","Aroldis Chapman","Joel Zumaya"),

MPH = c(108.1, 107.6, 106, 105.1, 104.8),

stringsAsFactors = FALSE

Pitchers_Speeds
```

```
> Pitchers_Speeds
    Pitchers_Name MPH
1    Nolan Ryan 108.1
2    Bob Feller 107.6
3 Aroldis Chapman 106.0
4 Aroldis Chapman 105.1
5    Joel Zumaya 104.8
```

● Data Frame访问

- d[r,]: rth row of object d
- d[,c]: cth column of object d
- d[r,c]: entry in row r and column c of object d
- d["age"]: extract column "age" from object d
- d\$age: extract column "age" from object d

Data Frame函数

- colnames()
- rownames()
- nrow()
- ncol()

● Data Frame函数

• merge() Two data frames can be merged into one data frame using the function merge.

```
test1 <- read.delim("test1.txt", sep =" ")</pre>
test1
   name year BA
1 Dick 1963 0.12 0.27
  Gose 1970 0.53 0.74
   Rolf 1971 0.53 0.28
4 Heleen 1974 0.81 0.29
test2 <- read.delim("test2.txt", sep =" ")</pre>
   name year
1 Dick 1963 0.42 0.12
2 Gose 1970 0.26 0.57
   Rolf 1971 0.87 0.37
4 Heleen 1974 0.86 0.15
test.merge <- merge(test1,test2)</pre>
test.merge
            BA HR
name year
1 Dick 1963 0.12 0.27 0.42 0.12
  Gose 1970 0.53 0.74 0.26 0.57
3 Heleen 1974 0.81 0.29 0.86 0.15
   Rolf 1971 0.53 0.28 0.87 0.37
```

● Data Frame函数



aggregate()

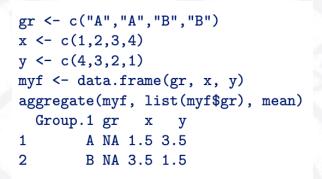
The function aggregate is used to aggregate data frames. It splits the data frame into groups and applies a function on each group.

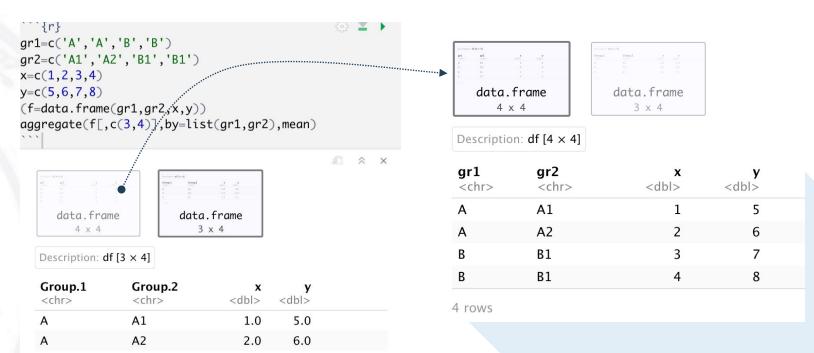
B1

3 rows

3.5

7.5





● Data Frame函数



 The function stack can be used to stack columns of a data frame into one column and one grouping column

```
group1 <- rnorm(3)
group2 <- rnorm(3)
group3 <- rnorm(3)
df <- data.frame(group1,group2, group3)</pre>
```

group1 <dbl></dbl>	group2 <dbl></dbl>	group3 <dbl></dbl>
-0.8935259	-1.05613011	0.7447249
-0.6174068	1.92203525	-0.7599139
0.6729324	-0.08939907	-0.4169499

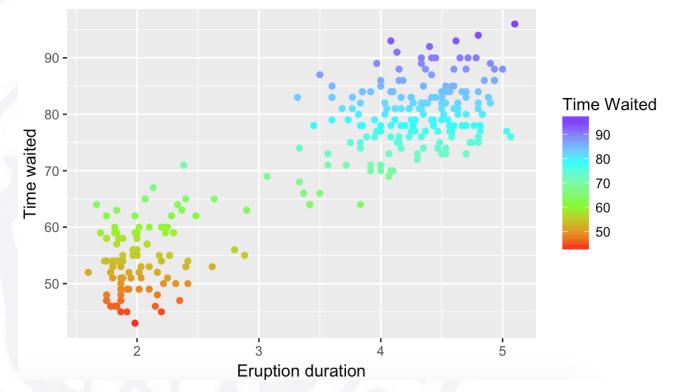
stack(df)

values <dbl></dbl>	ind <fctr></fctr>
-0.89352595	group1
-0.61740677	group1
0.67293242	group1
-1.05613011	group2
1.92203525	group2
-0.08939907	group2
0.74472487	group3
-0.75991386	group3
-0.41694994	group3

● Data Frame应用

```
ggplot(faithful, aes(x = faithful$eruptions, y = faithful$waiting, colour = faithful$waiting)) +
geom_point(aes()) +
scale_colour_gradientn(name = "Time Waited", colours=rainbow(4)) +
labs(x = "Eruption duration", y = "Time waited") +
ggtitle("Plot of Old Faithful Waiting and Eruption Times")
```

Plot of Old Faithful Waiting and Eruption Times



> faithful

	eruptions	waiting
1	3.600	79
2	1.800	54
3	3.333	74
4	2.283	62
5	4.533	85
6	2.883	55
7	4.700	88
8	3.600	85
9	1.950	51
10	4.350	85
11	1.833	54
12	3.917	84
13	4.200	78
14	1.750	47

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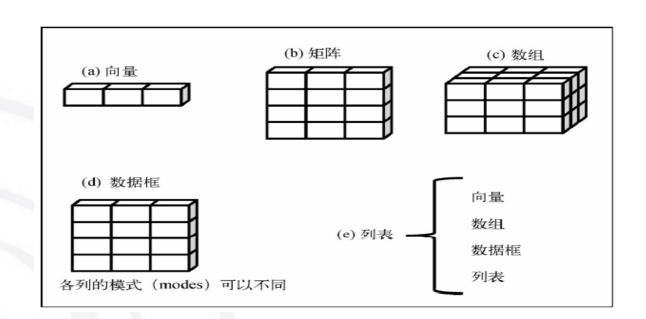
● List 生成&访问

 A list is an object containing elements of any type, including other objects

```
mylist <- list(a = 1:3, b = "hello", c = matrix(1:4, 2, 2))
mylist[1] # 返回包含元素a的列表
mylist["a"] # 返回包含元素a的列表
mylist$a # 访问元素 a
mylist[[1]] # 访问元素 a
```

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扩展阅读: ts

```
myts1 <- ts(data = rnorm(100), start=c(1987), freq = 12)
tsp(myts1)
myts2 <- ts(data = matrix(rnorm(100),ncol=2), start=c(1987,4), freq=12)
tsp(myts2)

[1] 1987.00 1995.25 12.00
[1] 1987.250 1991.333 12.000</pre>
```

Attributes

```
\``{r ts-2}
  attributes(myts1)

$tsp
[1] 1987.00 1995.25 12.00

$class
[1] "ts"
```

```
Jan
                     Feb
                               Mar
    0.046128499 0.729604206 1.571093199 0.793378194 -2.124475262
1989 1.136379750 0.354477352 -0.020180716 0.502346831 -2.049895972
1990 -1.605261218 -1.097419428 0.056647473 0.904257794 -1.492322196
1991 -0.997676795 0.732180442 -0.644546109 0.403500684 -0.056973110
1992 -0.713443569 1.015353141 -1.619333396 0.222546252 0.318044949
1993 -0.812307978 -0.776311435 0.116474834 0.634127039 -0.019213579
1994 0.129156883 0.975929852 1.528174631 0.054049106 1.329992724
1995 0.347079293 -1.662569406 1.134133996 -0.708107736
                               Aug
           Jun
                     Jul
                                          Sep
                                                    0ct
1987 -1.723999355 -1.711261593 0.308002424 -1.811040362 0.529584847
1989 -1.399155638 -0.537394067 -3.368797340 1.591608556 1.128888373
1991 -0.751655757 -0.637122966 -0.278422051 1.645725778 1.280306695
1992 -1.072479106 1.272251578 -0.276779046 1.189197255 0.229086277
1993 0.091579973 -0.206504772 -0.006067391 1.696747114 -0.892035033
1995
           Nov
                     Dec
    0.752834818 -1.912643758
1988 1.785845775 1.455185240
1989 0.201966987 -0.444604725
1990 0.565319694 0.044472613
1991 -0.139837665 0.361251897
1992 0.007191677 0.751474723
1993 -1.244280470 0.578293815
1994 1.141872976 1.196038968
1995
```

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Data Types

- Logical True/False
- Integer 4, 6, 2
- Numeric 3.4, 566, 2.34
- Character "a", "hello"
- Factor "Female", "Male"
- Date & Times "2021-09-15", " 2021-09-15 13:30:00"

Assigned Variables

Make sure to take into account data type when doing this

```
> a <- 1
> b <- 2
> a+b
[1] 3
> r <- "Hello"
> t <- "World"
> paste(r,t, sep = " ")
[1] "Hello World"
```

Character

nchar()

```
x <- c("a","b","c")
mychar1 <- "This is a test"
mychar2 <- "This is another test"
charvector <- c("a", "b", "c", "test")
nchar(mychar1)
[1] 15
nchar(charvector)
[1] 1 1 1 4</pre>
```

substring()

```
x <- c("Gose", "Longhow", "David")
substring(x,first=2,last=4)
[1] "ose" "ong" "avi"</pre>
```

paste()

```
paste("number",1:10, sep=".")
[1] "number.1" "number.2" "number.3" "number.4"
[5] "number.5" "number.6" "number.7" "number.8"
[9] "number.9" "number.10"
```

grep () Finding patterns in character objects

```
car.names <- row.names(cars)
grep("Volvo", car.names)
[1] 37</pre>
```

• gsub()

String Split

strsplit()

```
strsplit(x = c("Some text", "another string", split = NULL)
[[1]]
[1] "S" "o" "m" "e" " "t" "e" "x" "t"
[[2]]
 [1] "a" "n" "o" "t" "h" "e" "r" " "s" "t" "r" "i" "n" "g"
strsplit(
  x = c("Some~text" , "another-string", "Amsterdam is a nice city"),
  split = "[~-]"
[[1]]
[1] "Some" "text"
[[2]]
[1] "another" "string"
[[3]]
[1] "Amsterdam is a nice city"
```

Factor(因子)

- **名义型**变量是没有顺序之分的类别变量。下表中,糖尿病类型Diabetes(Type1、Type2)是名义型变量的一例。即使在数据中Type1编码为1而Type2编码为2,这也并不意味着二者是有序的。
- **有序型**变量表示一种顺序关系,而非数量关系。病情Status (poor, improved, excellent) 是顺序型变量的典型示例。我们知道,病情为poor (较差) 病人的状态不如improved (病情好转)的病人,但并不知道相差多少。

病人编号 (PatientID)	入院时间 (AdmDate)	年龄 (Age)	糖尿病类型 (Diabetes)	病情 (Status)
1	10/15/2009	25	Type1	Poor
2	11/01/2009	34	Type2	Improved
3	10/21/2009	28	Type1	Excellent
4	10/28/2009	52	Type1	Poor

Factor

- 名义型
 - diabetes<-c("type1","type2", "type1", "type1")
- 有序型
 - status<-c("Poor", "Improved", "Excellent", "Poor")
 - status <- factor(status, ordered=TRUE)
 - status<-factor(status, levels=c("Poor","Improved", "Excellent"))

Creating factors from continuous data

```
Example 1
x < -1:15
breaks <-c(0,5,10,15,20)
cut(x,breaks)
[1] (0,5] (0,5]
                         (0,5] (0,5] (5,10] (5,10] (5,10]
                   (0,5]
[10] (5,10] (10,15] (10,15] (10,15] (10,15]
Levels: (0,5] (5,10] (10,15] (15,20]
Example 2
cut( x, breaks=5)
 [1] (0.986,3.79] (0.986,3.79] (0.986,3.79] (3.79,6.6]
                                                     (3.79,6.6]
 [6] (3.79,6.6] (6.6,9.4] (6.6,9.4] (6.6,9.4] (9.4,12.2]
[11] (9.4,12.2] (9.4,12.2] (12.2,15] (12.2,15]
Levels: (0.986,3.79] (3.79,6.6] (6.6,9.4] (9.4,12.2] (12.2,15]
Example 3
x \leftarrow rnorm(15)
cut(x, breaks=3, labels=c("low", "medium", "high"))
           medium medium medium high
 [1] high
                                            low
                                                  high
                                                         low
                                                               low
[11] high
                        medium high
           low
                  low
Levels: low medium high
```

Date & Time

```
dz = z[2] -z[1]
dz
data.class(dz)
Time difference of 351 days
[1] "difftime"
```

```
t1 <- as.POSIXct("2003-01-23")
t2 <- as.POSIXct("2003-04-23 15:34")
t1
t2
[1] "2003-01-23 W. Europe Standard Time"
[1] "2003-04-23 15:34:00 W. Europe Daylight Time"
# pasting 4 character dates and 4 character times together
dates <-c("02/27/92", "02/27/92", "01/14/92", "02/28/92")
times <- c("23:03:20", "22:29:56", "01:03:30", "18:21:03")
x <- paste(dates, times)</pre>
z <- strptime(x, "%m/%d/%y %H:%M:%S")</pre>
zt <- as.POSIXct(z)</pre>
zt
[1] "1992-02-27 23:03:20 W. Europe Standard Time"
[2] "1992-02-27 22:29:56 W. Europe Standard Time"
[3] "1992-01-14 01:03:30 W. Europe Standard Time"
 [4] "1992-02-28 18:21:03 W. Europe Standard Time"
zt + 13 Unit: second
 [1] "1992-02-27 23:03:33 W. Europe Standard Time"
 [2] "1992-02-27 22:30:09 W. Europe Standard Time"
 [3] "1992-01-14 01:03:43 W. Europe Standard Time"
 [4] "1992-02-28 18:21:16 W. Europe Standard Time"
 t2 <- as.POSIXct("2004-01-23 14:33")
t1 <- as.POSIXct("2003-04-23")
 d < -t2-t1
 Time difference of 275.6479 days
```

Date & Time

- Functions
 - weekdays ()
 - months ()
 - quarters ()
 - Sys.time ()
 - •
- Packages
 - ZOO
 - chron
 - tseries
 - Rmetrics
 - •

```
weekdays(zt)
[1] "Thursday" "Thursday" "Tuesday" "Friday"
```

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Special Constants

- NA
- Inf
- -Inf
- TRUE
- FALSE

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Math Operators

Operator	Description
+	addition
-	subtraction
*	multiplication
/	division
^ or **	exponentiation
x %% y	modulus (x mod y) 5%%2 is 1
x %/% y	integer division 5%/%2 is 2

Relational Operators

Sign Meaning

- == Equals
- != Does not equal
- > Greater than
- >= Greater than or equal
 - < Less than
- <= Less than or equal to
- %in% Included in
- is.na() Is a missing value

```
```{r}
a=c(1,NA,3,3)
is.na(a)
a[is.na(a)]
a[!is.na(a)]
```

```
[1] FALSE TRUE FALSE FALSE
[1] NA
[1] 1 3 3
```

# String operation

- nchar()
- paste(str1,str2,sep)
- strsplit(string,sep)
- substr(string,start,stop)
- chartr(old,new,string)

# Object operation

函数	用途
length(object)	显示对象中元素/成分的数量
dim(object)	显示对象的维度
str(object)	显示对象的结构
class(object)	显示对象的类别
mode(object)	显示对象的模式
names(object)	显示象中各成分的名称
c(object,object,)	将对象合并入一个向量
cbind(object,object,)	按列合并对象
rbind(object,object,)	按行合并对象

# Convert operation

From To	Vector	Matrix	Data frame
Vector	c(x,y)	cbind(x,y) rbind(x,y)	data.frame(x,y)
Matrix	as.vector(mymatrix)		as.data.frame(mymatrix)
Data frame		as.matrix(myframe)	

# Apply operation

```
fr for_loop}
set.seed(431)
 (mat43=replicate(4,sample(3,3)))
 (mat43l=rep(0,4))|
 for (j in 1:4) {mat43l[j]=max(mat43[,j])}
 mat43l
```

```
[,1] [,2] [,3] [,4]
[1,] 3 2 2 3
[2,] 2 1 3 2
[3,] 1 3 1 1
[1] 0 0 0 0
[1] 3 3 3 3
```

```
apply(mat43,2,max)
 ## [1] 3 3 3 3
 `{r apply}
apropos("apply")
[1] ".rs.api.applyTheme" ".rs.applyTheme"
[3] ".rs.applyTransform"
 "apply"
[5] "dendrapply"
 "eapply"
[7] "kernapply"
 "lapply"
 "rapply"
[9] "mapply"
 "tapply"
[11] "sapply"
[13] "vapply"
```

#### 扩展阅读:

《An introduction to R》 6 Efficient calculations

### Contents

- Objects
  - Vectors
  - Arrays
  - Matrices
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- I/O

### Conditions

if (cond) {expr1} else {expr2}

```
> w = 3
> if(w < 5)
 {
 d=2
 }else{
 d=10
 }
> d
```

#### switch

## Loops

# While-loop while (cond) {expr}

```
while(tmp < 100){
 tmp <- tmp + rbinom(1,10,0.5)
 n <- n +1
}</pre>
```

# For-loop for (var in vec) {expr}

```
for(i in 2:length(x))
{
 x[i] <- x[i] + phi*x[i-1]
}</pre>
```

# Repeat-loop

```
repeat
{
 some expressions
}
```

In the repeat loop some expressions are repeated 'infinitely', so repeat loops will have to contain a break statement to escape them.

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#### Build-in functions

#### Data creation

- read.table: read a table from file. Arguments: header=TRUE: read first line as titles of the columns; sep=",": numbers are separated by commas; skip=n: don't read the first n lines.
- write.table: write a table to file
- c: paste numbers together to create a vector
- array: create a vector, Arguments: dim: length
- matrix: create a matrix, Arguments: ncol and/or nrow: number of rows/columns
- data.frame: create a data frame
- list: create a list
- rbind and cbind: combine vectors into a matrix by row or column

#### Data processing

- seq: create a vector with equal steps between the numbers
- rnorm: create a vector with random numbers with normal distribution (other distributions are also available)
- sort: sort elements in increasing order
- t: transpose a matrix
- aggregate(x,by=ls(y),FUN="mean"): split data set x into subsets (defined by y) and computes means of the subsets. Result: a new list.
- na.approx: interpolate (in zoo package). Argument: vector with NAs. Result: vector without NAs.
- cumsum: cumulative sum. Result is a vector.
- rollmean: moving average (in the zoo package)
- paste: paste character strings together
- substr: extract part of a character string

#### Function

#### name=function(arglist){expr}

```
ExpAnd <- function(vec=seq(4,25,3),exponent=2,addto=3)
{
 # Function that takes argument vec to the power
 # exp, adds add and then outputs the result
 out<-vec^exponent+addto
 return(out)
}
ExpAnd()</pre>
```

```
> fun1 = function(arg1, arg2)
 {
 w = arg1 ^ 2
 return(arg2 + w)
 }
> fun1(arg1 = 3, arg2 = 5)
[1] 14
```

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- · 1/0

## I/O: txt file

```
file.show("matHap.txt")
```

```
"DYS19" "DXYS156Y" "DYS389m" "DYS389n" "DYS389p"
"H1" 14 12 4 12 3 10 8 10 1 4 15 13 0 1 1
"H3" 15 13 4 13 3 9 8 10 1 4 13 12 0 1 1
"H4" 15 11 5 11 3 10 8 10 1 4 11 14 0 1 1
```

```
haptable=read.table("matHap.txt")
haptable
```

```
##
 DYS19 DXYS156Y DYS389m DYS389p DYS389p
H1
 14
 12
 12
H3
 15
 13
 13
 15
 11
 13
H5
 11
 12
H7
 13
 12
H8
 16
 11
 12
```

```
> d = data.frame(a = c(3,4,5),
b = c(12,43,54)
> d
 b
 a
1 3 12
2 4 43
3 5 54
> write.table(d, file="tst0.txt",
row.names=FALSE)
> d2 = read.table(file="tst0.txt",
header=TRUE)
> d2
 b
 a
1 3 12
2 4 43
3 5 54
```

## I/O: csv file

file.show("Haplotype.csv")

```
> read.csv
function (file, header = TRUE, sep = ",", quote = "\"", dec = ".",
 fill = TRUE, comment.char = "", ...)
read.table(file = file, header = header, sep = sep, quote = quote,
 dec = dec, fill = fill, comment.char = comment.char, ...)
<bytecode: 0x7fbbca2577b0>
<environment: namespace:utils>
```

Hap=read.csv("/Users/susan/Dropbox/stat32/Slides\_ABCofR/Haplotype.csv")
head(Hap)

```
Hapscan<-scan("/Users/susan/Dropbox/stat32/Slides_ABCofR/Haplotype.csv",nlines=5,what="")
Hapscan</pre>
```

```
[1] "Individual,DYS19,DXYS156Y,DYS389m,DYS389n,
[2] "H1,14,12,4,12,3,10,8,10,1,4,15,13,0,1,1"
[3] "H3,15,13,4,13,3,9,8,10,1,4,13,12,0,1,1"
[4] "H4,15,11,5,11,3,10,8,10,1,4,11,14,0,1,1"
[5] "H5,17,13,4,11,3,10,7,10,1,4,14,12,0,1,1"
```

write.csv(data,file="data.csv",row.names = FALSE)

# I/O: excel file

```
install.packages("readxl")
```

```
library(readx1)
?read_excel
Hapall <- read_excel("/Users/susan/RWork/data/Haplotype.xlsx")</pre>
```

Find more at <a href="https://cran.r-project.org/doc/manuals/r-release/R-data.html">https://cran.r-project.org/doc/manuals/r-release/R-data.html</a>

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# Canvas 随堂测试

# 课后练习

- 分析faithful数据
  - 保存到csv文件(write.csv)
  - 从csv文件读入(read.csv)
  - · 查看数据一阶统计特征(summary)
  - 画数据散点图(plot)
  - · 画数据分布图(hist)
  - 根据以上分析,回答下面问题
    - 如果游客刚错过一次喷泉,您建议游客多久后回来看下一次喷泉?
    - 你能根据等待的时间来估计下一次喷泉能喷多久吗?



# Thanks

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