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Console ~/     
> source('~/.active-rstudio-document', echo=TRUE)
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```
> #创建向量  
> a <- c(1, 2, 5, 3, 6, -2, 4)  
  
> b <- c("one", "two", "three")  
  
> c <- c(TRUE, TRUE, TRUE, FALSE, TRUE, FALSE)  
  
> #使用向量  
> a <- c(1, 2, 5, 3, 6, -2, 4)  
  
> a[3]  
[1] 5  
  
> a[c(1, 3, 5)]  
[1] 1 5 6  
  
> a[2:6]  
[1] 2 5 3 6 -2  
  
> a <- c("k", "j", "h", "a", "c", "m")  
  
> a[3]  
[1] "h"  
  
> a[c(1, 3, 5)]  
[1] "k" "h" "c"  
  
> a[2:6]  
[1] "j" "h" "a" "c" "m"  
  
> #创建矩阵  
> y <- matrix(1:20,nrow=5,ncol=4)  
  
> y  
      [,1] [,2] [,3] [,4]  
[1,]    1    6   11   16  
[2,]    2    7   12   17  
[3,]    3    8   13   18  
[4,]    4    9   14   19  
[5,]    5   10   15   20  
  
> cells <- c(1,26,24,68)  
  
> rnames <- c("R1", "R2")
```

Console ~/ ↗



```
> cnames <- c("C1", "C2")

> mymatrix <- matrix(cells, nrow=2, ncol=2, byrow=TRUE, dimnames=list(rnames, cnames))

> mymatrix
  C1 C2
R1 1 26
R2 24 68

> #byrow 默认为FALSE, R如何将传入的向量填充入矩阵, byrow时优先填满行
> mymatrix <- matrix(cells, nrow = 2, ncol = 2, byrow = FALSE, dimnames = list(rnames, cnames))

> mymatrix
  C1 C2
R1 1 24
R2 26 68

> #使用矩阵
> x <- matrix(1:10, nrow=2)

> x
      [,1] [,2] [,3] [,4] [,5]
[1,]    1    3    5    7    9
[2,]    2    4    6    8   10

> x[2,]
[1] 2 4 6 8 10

> #访问矩阵的方式matrix[row_index,col_index]row_index或col_index缺省则代表访问全部的列或者行
> x[,2]
[1] 3 4

> x[1,4]
[1] 7




> x[1, c(4,5)]
[1] 7 9




> #创建数组
> dim1 <- c("A1", "A2")

> dim2 <- c("B1", "B2", "B3")

> dim3 <- c("C1", "C2", "C3", "C4")

> z <- array(1:24, c(2,3,4), dimnames = list(dim1, dim2, dim3))
```

```
Console ~/     
> z  
, , C1  
  
      B1 B2 B3  
A1  1  3  5  
A2  2  4  6  
  
, , C2  
  
      B1 B2 B3  
A1  7  9 11  
A2  8 10 12  
  
, , C3  
  
      B1 B2 B3  
A1 13 15 17  
A2 14 16 18  
  
, , C4  
  
      B1 B2 B3  
A1 19 21 23  
A2 20 22 24  
  
> #创建数据框  
> patientID <- c(1,2,3,4)  
  
> age <- c(25, 34, 28, 52)  
  
> diabetes <- c("Type1", "Type2", "Type1", "Type1")  
  
> status <- c("Poor", "Improved", "Excellent", "Poor")  
  
> patientdata <- data.frame(patientID, age, diabetes, status)  
  
> patientdata  
  patientID age diabetes  status  
1         1  25   Type1    Poor  
2         2  34   Type2 Improved  
3         3  28   Type1 Excellent  
4         4  52   Type1    Poor  
  
> #指定数据框中的数据  
> patientdata[1:2]  
  patientID age
```

```
Console ~/   
1      1 25
2      2 34
3      3 28
4      4 52

> patientdata[c("diabetes","status")]
  diabetes status
1   Type1   Poor
2   Type2 Improved
3   Type1 Excellent
4   Type1   Poor

> patientdata$age
[1] 25 34 28 52

> #使用系数
> patientID <- c(1, 2, 3, 4)

> age <- c(25, 34, 28, 52)

> diabetes <- c("Type1", "Type2", "Type1", "Type1")

> status <- c("Poor", "Improved", "Excellent", "Poor")

> diabetes <- factor(diabetes)

> status <- factor(status, order=TRUE)

> patientdata <- data.frame(patientID, age, diabetes, status)

> str(patientdata)
'data.frame':  4 obs. of  4 variables:
 $ patientID: num  1 2 3 4
 $ age      : num  25 34 28 52
 $ diabetes : Factor w/ 2 levels "Type1","Type2": 1 2 1 1
 $ status   : Ord.factor w/ 3 levels "Excellent"<"Improved"<..: 3 2 1 3

> summary(patientdata)
  patientID      age
Min.   :1.00  Min.   :25.00
1st Qu.:1.75  1st Qu.:27.25
Median :2.50  Median :31.00
Mean   :2.50  Mean   :34.75
3rd Qu.:3.25  3rd Qu.:38.50
Max.   :4.00  Max.   :52.00
 diabetes      status
Type1:3  Excellent:1
```

```
> summary(patientdata)
  patientID      age
Min.   :1.00   Min.   :25.00
1st Qu.:1.75   1st Qu.:27.25
Median :2.50   Median :31.00
Mean   :2.50   Mean    :34.75
3rd Qu.:3.25   3rd Qu.:38.50
Max.   :4.00   Max.    :52.00
diabetes      status
Type1:3   Excellent:1
Type2:1   Improved :1
          Poor      :2

> #创建list
> g <- "My First List"

> h <- c(25, 26, 18, 39)

> j <- matrix(1:10, nrow=5)

> k <- c("one", "two", "three")

> mylist <- list(title=g, ages=h, j, k)

> mylist
$title
[1] "My First List"

$ages
[1] 25 26 18 39

[[3]]
      [,1] [,2]
[1,]    1    6
[2,]    2    7
[3,]    3    8
[4,]    4    9
[5,]    5   10

[[4]]
[1] "one"  "two"  "three"
```

> |