

## Basic R - List, matrix and array

# Creating a list

- Lists are the R objects which contain elements of different types, like numbers, strings, vectors, matrix, function and even some lists inside it.
- We can create a list using *list()* function

```
x<-1
y<-c("A","B")
z<-matrix(5,2,2)
f<-function(x){f<-x^2}
t<-list(x,y,z)
mylist<-list(x,y,z,f,t)
```

# Creating a list

```
str(mylist)
```

```
## List of 5
## $ : num 1
## $ : chr [1:2] "A" "B"
## $ : num [1:2, 1:2] 5 5 5 5
## $ :function (x)
## ..- attr(*, "srcref")= 'srcref' int [1:8] 4 4 4 22 4 22 4
## ..- attr(*, "srcfile")=Classes 'srcfilecopy', 'srcfile'
## $ :List of 3
## ..$ : num 1
## ..$ : chr [1:2] "A" "B"
## ..$ : num [1:2, 1:2] 5 5 5 5
```

# Subsetting a list

- "[.]" extracts a sub-list, the result will be a list.

```
str(mylist[3])
```

```
## List of 1  
## $ : num [1:2, 1:2] 5 5 5 5
```

```
str(mylist[5])
```

```
## List of 1  
## $ :List of 3  
## ..$ : num 1  
## ..$ : chr [1:2] "A" "B"  
## ..$ : num [1:2, 1:2] 5 5 5 5
```

# Subsetting a list

- "[[.]]" extracts a single component from a list; it removes a level of hierarchy from the list.

```
str(mylist[[3]])
```

```
##  num [1:2, 1:2] 5 5 5 5
```

```
str(mylist[[5]])
```

```
## List of 3
```

```
## $ : num 1
```

```
## $ : chr [1:2] "A" "B"
```

```
## $ : num [1:2, 1:2] 5 5 5 5
```

# Naming elements of a list

```
names(mylist)<-c("x","y","z","f","t")  
mylist$f # using "$" accessor ~ [[.]]
```

```
## function(x){f<-x^2}  
print(mylist$f(mylist$z))
```

```
##      [,1] [,2]  
## [1,]   25   25  
## [2,]   25   25
```

# Manipulating List Elements

- Adding an element

```
mylist[6]<-list(c(1,2,3))  
mylist<-c(mylist,list(c(4,5,6)))
```

- Remove an element of a list

```
mylist[6]<-NULL  
str(mylist)
```

```
## List of 6  
## $ x: num 1  
## $ y: chr [1:2] "A" "B"  
## $ z: num [1:2, 1:2] 5 5 5 5  
## $ f:function (x)  
## ..- attr(*, "srcref")= 'srcref' int [1:8] 4 4 4 22 4 22 4  
## .. ..- attr(*, "srcfile")=Classes 'srcfilecopy', 'srcfile'  
## $ t:List of 3
```

# Practice

Write a function of a numeric vector  $\mathbf{x}$ , named *mysummary*, with output is a list:

- The first element, named "mean", is the average value of  $\mathbf{x}$
- The second element, named "sd", is the standard deviation of  $\mathbf{x}$
- The third element, named "quantile", is a numeric vector of 5 elements:  $\min(x)$ , 1<sup>st</sup> quartile, the median, 3<sup>rd</sup> quartile and  $\max(x)$
- The fourth element, names "plot", is the density plot of  $\mathbf{x}$



# Practice

```
mysummary<-function(x){  
  m<-mean(x,na.rm=TRUE)  
  sd<-sd(x,na.rm = TRUE)  
  quan<-c(min(x,na.rm=TRUE),  
           quantile(x,0.25,na.rm=TRUE),  
           quantile(x,0.5,na.rm=TRUE),  
           quantile(x,0.75,na.rm=TRUE),  
           max(x))  
  pl<-plot(density(x))  
  mylist<-list(m,sd,quan,pl)  
  names(mylist)<-c("mean","sd","quantile","plot")  
  mysummary<-mylist  
}
```

# Matrix

- Matrix is the R objects in which the elements are arranged in a two-dimensional.
- Matrices contains elements of the same type i.e. we can only create a matrix containing only numbers or only logical values.
- Matrices containing numeric elements are very useful in data analytics.

```
M<-matrix(1:6,2,3) # 2*3 matrix  
M
```

```
##      [,1] [,2] [,3]  
## [1,]    1    3    5  
## [2,]    2    4    6
```

# Accessing elements of a matrix

```
# Access the element at 2nd column and 1st row.  
print(M[1,2])
```

```
## [1] 3
```

```
# Access the element at 2nd column and 3th row.  
print(M[2,3])
```

```
## [1] 6
```

```
# Access only the 2nd row.  
print(M[2,])
```

```
## [1] 2 4 6
```

```
# Access only the 3rd column.  
print(M[,3])
```

```
## [1] 5 6
```

# Matrix computations

```
print(M*2)
```

```
##           [,1] [,2] [,3]
## [1,]         2    6   10
## [2,]         4    8   12
```

```
M1<-matrix(rep(c(1,2),3),2,3)
```

```
M*M1
```

```
##           [,1] [,2] [,3]
## [1,]         1    3    5
## [2,]         4    8   12
```

```
M+M1
```

```
##           [,1] [,2] [,3]
## [1,]         2    4    6
## [2,]         4    6    8
```

# Matrix computations

Operation/formula	Note
$M_1 \% * \% M_2$	Matrix multiplication, where $M_1$ is a $n * m$ and $M_2$ is a $m * k$ matrix
$t(M)$	The transpose matrix of matrix $M$
$det(M)$	The determination of squared matrix $M$
$solve(M)$	The inversed matrix of squared matrix $M$
$chol(M)$	The Cholesky decomposition
$eigen(M)\$values$	The eigenvalues of the matrix $M$
$eigen(M)\$vectors$	The eigenvectors of the matrix $M$

# Matrix application

Matrix representation of an image

```
dat <- read_mnist() # size 60000 * 784

# matrix 28*28 of an image
MM<-matrix(mnist$train$images[8,],28,28)
MM

image(1:28,1:28,MM,
      col = gray.colors(10,start=0,end=1)) #black-white

MM<-ifelse(MM>0,255,0)
image(1:28,1:28,MM,
      col = gray.colors(10,start=0,end=1))
```

# Matrix application

Solve the system equation

$$9x + 8y + 9z + 2t = 42$$

$$5x + 2y + 7z + 3t = 45$$

$$6x + 4y + 3z + 6t = 53$$

$$8x + 2y + 5z + 6t = 63$$

# Matrix application

```
M<-matrix(c(c(9,5,6,8),  
             c(8,2,4,2),  
             c(9,7,3,5),  
             c(2,3,6,6)),4,4)  
v<-matrix(c(42,45,53,63),4,1)
```

*# SOLUTION (x,y,z,t)*

```
solve(M)%*%v
```

```
##      [,1]  
## [1,]    1  
## [2,]   -1  
## [3,]    3  
## [4,]    7
```



# Matrix application

Calculation of least squares estimates by matrix manipulation: You are fitting a multiple linear regression model

$$Y_i = \beta_0 + \beta_1 X_1 + \cdots + \beta_p X_p$$

- a. You are using Boston data set in package MASS, the dependent variable ( $Y$ ) is *medv* and there are 3 independent variables: *lstat*, *crim*, *rm*.
- b. Set up the vector  $Y$  and matrix  $\mathbf{X}$
- c. Using the following formula to calculate the least squares estimate:  
$$\hat{\beta} = (X'X)^{-1}X'Y$$
- d. Compare result in (c.) to the result from *lm* function

# Matrix application

```
library(MASS)
n<-nrow(Boston)
X<-matrix(c(rep(1,n),Boston$lstat,Boston$crim,Boston$rm),n,4)
Y<-matrix(Boston$medv,n,1)
#c. b0,b1,b2,b3
t(solve(t(X)%*%X) %*% t(X) %*% Y)
```

```
##           [,1]      [,2]      [,3]      [,4]
## [1,] -2.562251 -0.5784858 -0.1029409  5.216955
```

```
#d. check with lm
lm(medv~lstat+crim+rm,data=Boston)
```

```
##
## Call:
## lm(formula = medv ~ lstat + crim + rm, data = Boston)
##
## Coefficients:
```

# Array

- Arrays are objects which can store data in more than 2 dimensions.
- If we create an array of dimension  $(n, m, k)$ , it creates  $k$  rectangular matrices each with  $n$  rows and  $m$  columns.
- Arrays can store only data type.

```
# 2 matrix 3*4
```

```
M<-array(1:24,dim=c(3,4,2))
```

```
M
```

```
## , , 1
```

```
##
```

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

```
## [1,]    1    4    7   10
```

```
## [2,]    2    5    8   11
```

```
## [3,]    3    6    9   12
```

```
##
```

# Accessing array elements

```
M[, ,1] # 3*4 matrix
```

```
##      [,1] [,2] [,3] [,4]  
## [1,]    1    4    7   10  
## [2,]    2    5    8   11  
## [3,]    3    6    9   12
```

```
M[,4,1] # vector length = 3
```

```
## [1] 10 11 12
```

```
M[1,2,2] # a number
```

```
## [1] 16
```

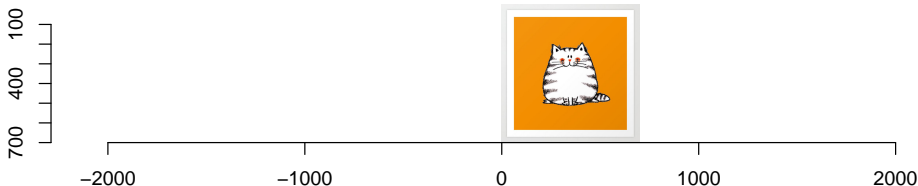
# Application of array

Images and videos are 4-dimension array

```
library(imager)
setwd("/Users/yenvv/Documents/Stochastic_simulation_with_R/Images")
img<-load.image("cat.jpg")
dim(img) # width, height, depth, spectrum
```

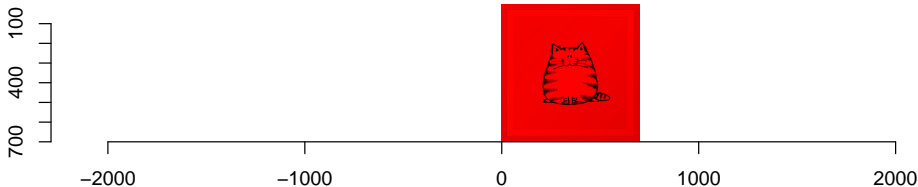
```
## [1] 700 700 1 3
```

```
plot(img) # plot a 4-dimension array
```



# Application of array

```
img1<-img  
img1[,,,2:3]<-0 # red only  
plot(img1) # red only
```

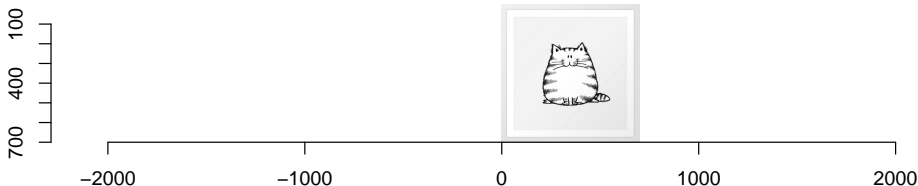


```
img1<-img  
img1[,,,1:2]<-0 # blue only  
plot(img1) # blue only
```



# Application of array

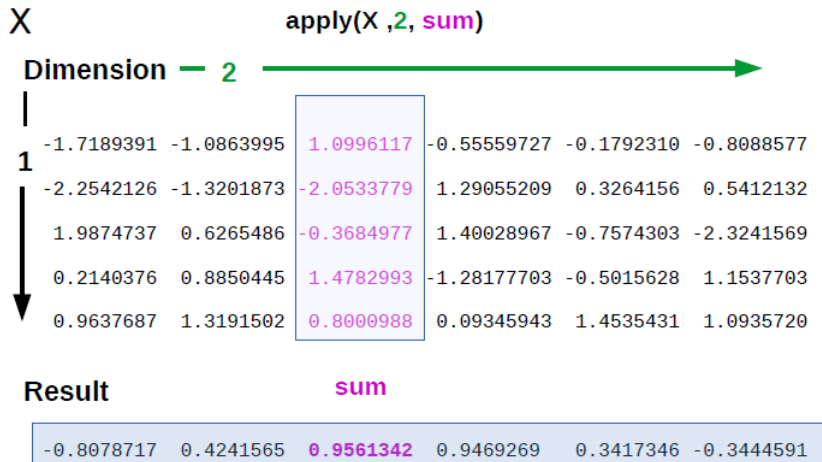
```
arr1<-img[, ,1,1] # 2-dimension  
img1<-array(arr1,dim=c(700,700,1,1))  
plot(as.cimg(img1))
```



```
# add random (noise) into  
arr2<-arr1+array(runif(700*700,0,2),dim=c(700,700))  
img2<-array(arr2,dim=c(700,700,1,1)) # re dim  
plot(as.cimg(img2))
```



# apply() function





# *apply()* function

- This function allow crossing the data in a number of ways and avoid use of loop constructs.
- *apply()* function can manipulate data from matrices, arrays, lists and dataframe.
- *apply()* function form the basis of more complex combinations and helps to perform operations with very few lines of code including: *lapply()* , *sapply()*, *vapply()*, *mapply()*, *rapply()*, and *tapply()* functions.

# *apply()* function

Using `apply()` function to normalize (to  $[0,1]$ ) variables in Boston data set.

```
normalize<-function(x){x<-(x-min(x))/(max(x)-min(x))}
```

```
# normalize all variables (columns) of Boston
```

```
dat<-apply(Boston,2,normalize)
```

```
class(dat)
```

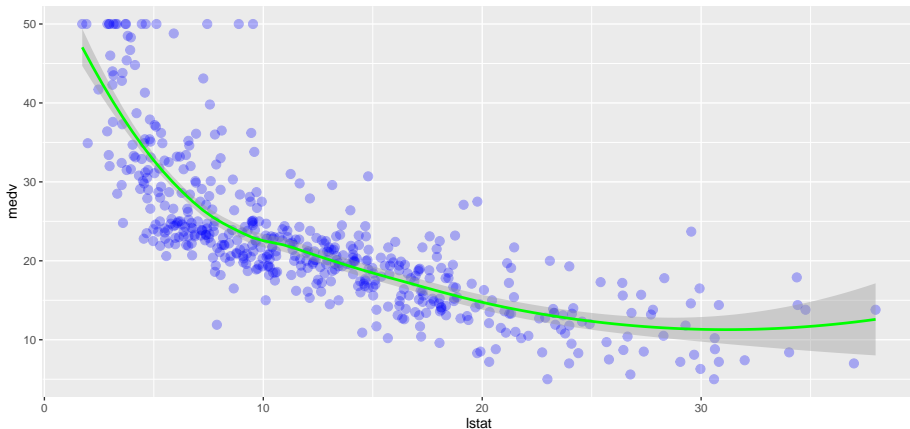
```
## [1] "matrix" "array"
```

```
dat<-as.data.frame(dat,colnames=names(Boston))
```

# *apply()* function

Using `apply()` function to normalize (to  $[0,1]$ ) variables in Boston data set.

Boston data set before normalizing



# *apply()* function

Using `apply()` function to normalize (to  $[0,1]$ ) variables in Boston data set.

Boston data set after normalizing

