

# *The R Book*

## **Chapter 2: Essentials of the R Language**

### **Session 12**

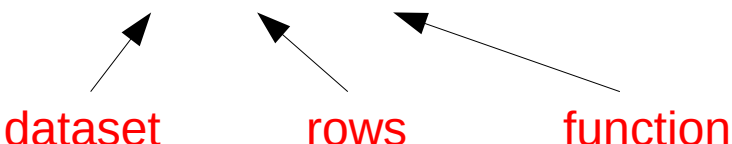
# Evaluating Functions with apply, sapply and lapply

## apply

- to apply functions to rows or columns of matrices or dataframes

```
> (X<-matrix(1:24,nrow=4))  
      [,1] [,2] [,3] [,4] [,5] [,6]  
[1,]    1    5    9   13   17   21  
[2,]    2    6   10   14   18   22  
[3,]    3    7   11   15   19   23  
[4,]    4    8   12   16   20   24
```

```
> apply(X, 1, sum)
```



dataset      rows      function

The diagram shows three red labels: 'dataset', 'rows', and 'function'. Three black arrows point from these labels to the arguments in the function call 'apply(X, 1, sum)'. The arrow from 'dataset' points to 'X', the arrow from 'rows' points to '1', and the arrow from 'function' points to 'sum'.

```
[1] 66 72 78 84
```

# Evaluating Functions with apply, sapply and lapply


## apply

- to apply functions to rows or columns of matrices or dataframes

```
> apply(X, 1, function(x) x^2+x)
```

	[, 1]	[, 2]	[, 3]	[, 4]
[1, ]	2	6	12	20
[2, ]	30	42	56	72
[3, ]	90	110	132	156
[4, ]	182	210	240	272
[5, ]	306	342	380	420
[6, ]	462	506	552	600

"anonymous function",  
not named



# Evaluating Functions with `apply`, `sapply` and `lapply`

## `sapply`

- to apply functions to vectors
- useful with complex iterative calculations

```
> sapply(3:7, seq)
```

```
[[1]]
```

```
[1] 1 2 3
```

```
[[2]]
```

```
[1] 1 2 3 4
```

```
[[3]]
```

```
[1] 1 2 3 4 5
```

```
[[4]]
```

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

```
[[5]]
```

```
[1] 1 2 3 4 5 6 7
```

# Evaluating Functions with `apply`, `sapply` and `lapply`

## `sapply`

Example : Decay of radioactive emissions, over a 50-day period

- use non-linear least squares to **estimate the decay constant a** in

$$y = \exp(-ax)$$

### a) upload a dataset

```
> sapdecay<-read.table("/home/.../sapdecay.txt", header=T)
> attach(sapdecay)
> names(sapdecay)
[1] "ex" "wy"
```

### b) calculate the sum of the squares $wy \sim$ predicted ( $yf$ ) values

```
> sumsq <- function(a,xv=ex,yv=wy)
+ { yf <- exp(-a*xv)
+ sum((yv-yf)^2) }
```

# Evaluating Functions with apply, sapply and lapply

## sapply

- use non-linear least squares to **estimate the decay constant a**

$$y = \exp(-ax)$$

c) get a rough idea of the decay constant a,

```
> lm(log(wy) ~ ex)
```

Call:

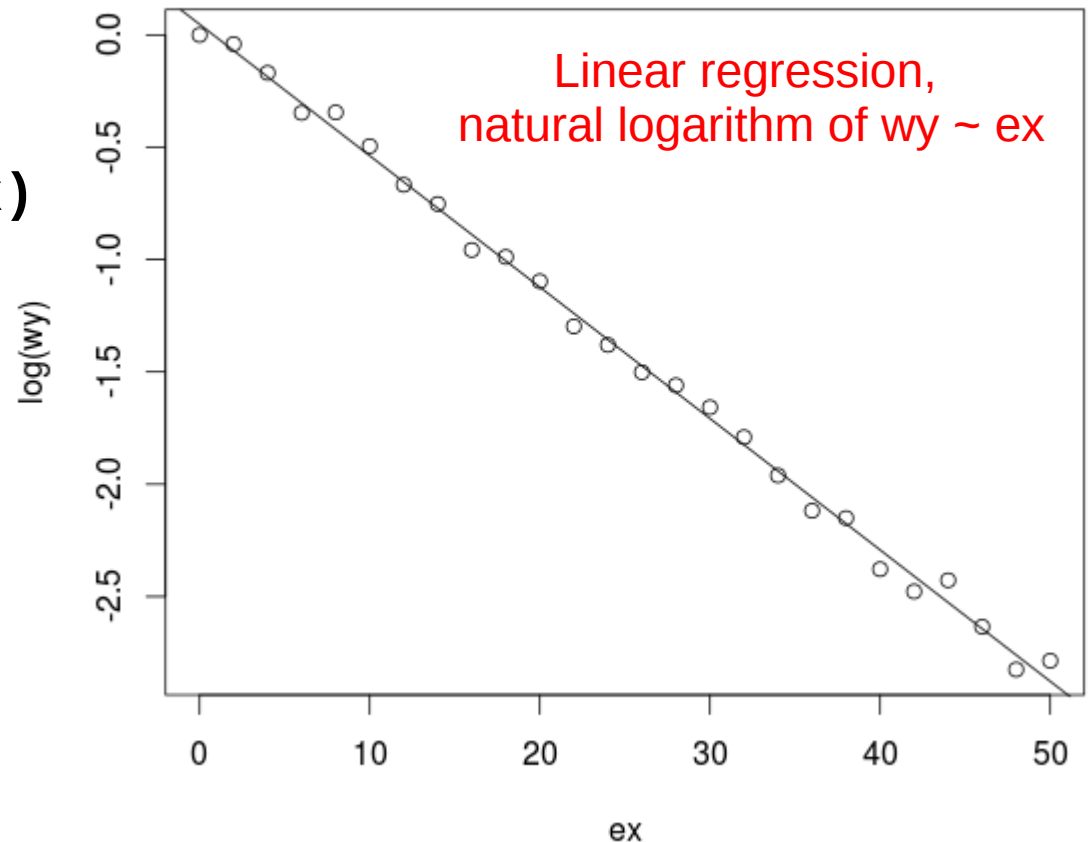
```
lm(formula = log(wy) ~ ex)
```

Coefficients:

(Intercept)	ex
0.04688	-0.05849

```
> plot(log(wy) ~ ex)
```

```
> abline(lm(log(wy) ~ ex))
```



# Evaluating Functions with apply, sapply and lapply

## sapply

- use non-linear least squares to **estimate the decay constant a**

d) generate a range of values for a on either side of 0.058

```
> a<-seq(0.01,0.2,.005)
```

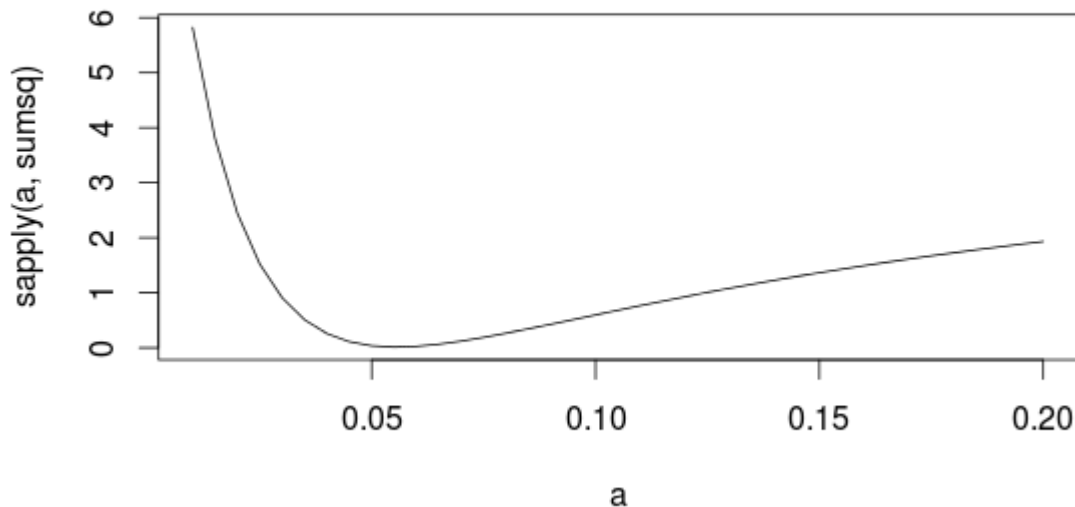
```
> a
```

```
[1] 0.010 0.015 0.020 0.025 0.030 0.035 0.040 0.045 0.050 0.055 0.060 0.065 0.070 0.075 0.080  
[16] 0.085 0.090 0.095 0.100 0.105 0.110 0.115 0.120 0.125 0.130 0.135 0.140 0.145 0.150 0.155  
[31] 0.160 0.165 0.170 0.175 0.180 0.185 0.190 0.195 0.200
```

e) use sapply to apply the sum of squares for each value «a»

```
> plot(a, sapply(a, sumsq), type="l")
```

← Type "l" for lines



```
> sumsq<-function(a,xv=ex,yv=wy)  
+ { yf <- exp(-a*xv)  
+ sum((yv-yf)^2) }
```

# Evaluating Functions with apply, sapply and lapply

## sapply

- use non-linear least squares to **estimate the decay constant a**

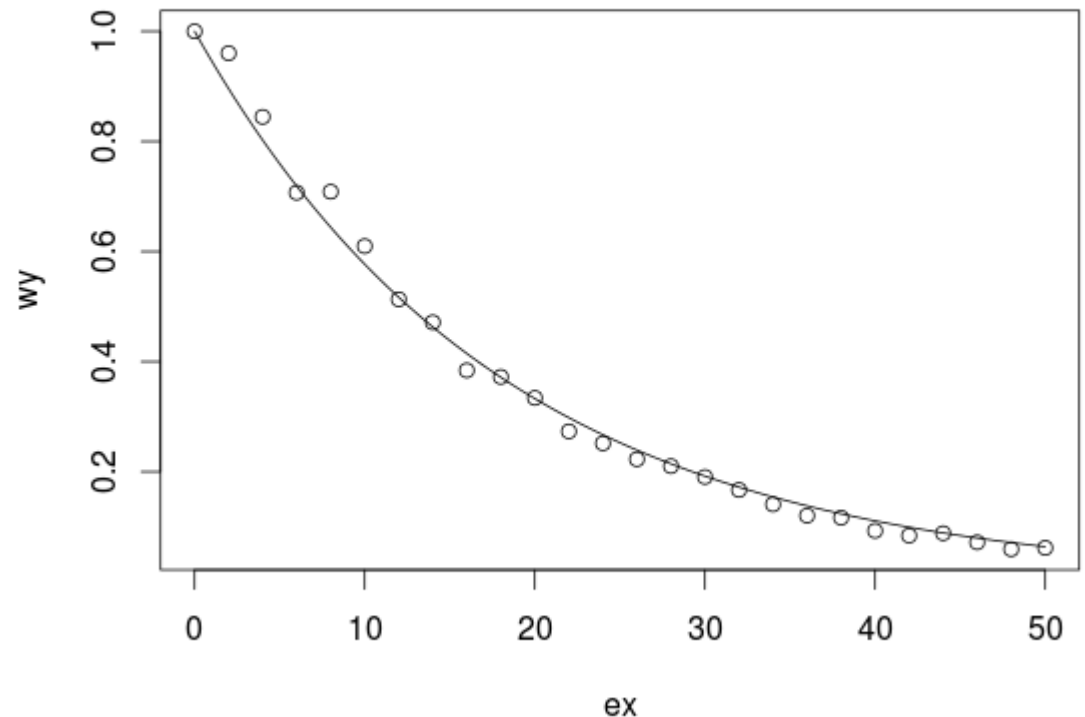
f) determine « a » with the smallest sum of squares

```
> a[min(sapply(a, sumsq)) == sapply(a, sumsq)]  
[1] 0.055
```

g) plot decay values with regression function and fitted « a »

```
> plot(ex, wy)  
> xv <- seq(0, 50, 0.1)  
> lines(xv, exp(-0.055 * xv))
```

- use :  
**xv <- seq(0, 50, 10)**





# Evaluating Functions with apply, sapply and lapply

## sapply

- use non-linear least squares to **estimate the decay constant a**

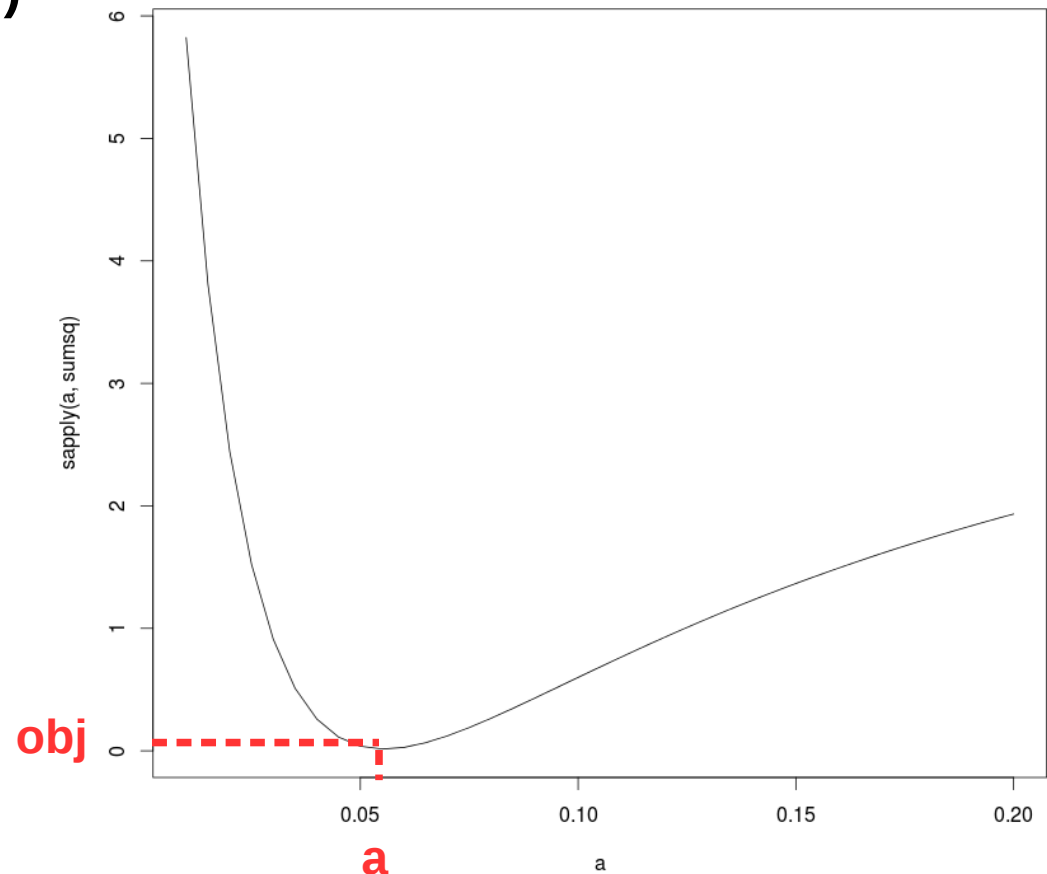
### h) streamlined version using the optimize function

```
> fa<-function(a) sum((wy-exp(-a*ex))^2)
> optimize(fa,c(0.01,0.1))
```

```
$minimum
[1] 0.05538411
```

```
$objective
[1] 0.01473559
```

```
> sumsq(0.05538411)
[1] 0.01473559
```



# Evaluating Functions with apply, sapply and lapply

## Lists and lapply

```
> a<-c("a", "b", "c", "d")
> b<-c(1, 2, 3, 4, 4, 3, 2, 1)
> c<-c(T, T, F)
```

character information  
numeric information  
logical information

```
> list.object<-list(a, b, c)
> class(list.object)
[1] "list"
```

combine into a list

---

```
> list.object
[[1]]
[1] "a" "b" "c" "d"

[[2]]
[1] 1 2 3 4 4 3 2 1

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

```
> lapply(list.object, length)
[[1]]
[1] 4

[[2]]
[1] 8

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

# Evaluating Functions with apply, sapply and lapply

## Lists and lapply

```
> a<-c("a", "b", "c", "d")
> b<-c(1, 2, 3, 4, 4, 3, 2, 1)
> c<-c(T, T, F)
```

character information  
numeric information  
logical information

```
> list.object<-list(a, b, c)
> class(list.object)
[1] "list"
```

combine into a list

---

```
> list.object
[[1]]
[1] "a" "b" "c" "d"

[[2]]
[1] 1 2 3 4 4 3 2 1

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

```
> lapply(list.object, class)
[[1]]
[1] "character"

[[2]]
[1] "numeric"

[[3]]
[1] "logical"
```

# Looking for runs of numbers within vectors

Function `rle` ('run length encoding')

```
> (poisson<-rpois(150,0.7))
```

```
 [1] 1 0 0 0 0 1 0 0 1 1 0 1 1 0 0 0 1 2 1 0 0 1 1 0 0 1 2 0 1 2 0 1 1 0
0 0 3 1 0 0 1 1 0 1 0 1 1 0 1 0 0 0 1
 [54] 0 0 0 2 1 1 0 0 1 1 0 1 0 0 0 2 1 0 2 2 1 1 0 1 2 1 0 0 0 0 0 1 1 2
0 2 0 0 2 0 0 0 3 0 2 1 0 3 1 0 1 1 1
[107] 0 2 1 0 0 1 0 1 1 1 1 1 0 0 1 0 0 0 0 0 2 1 1 2 0 1 0 0 1 0 1 0 0
0 0 1 1 1 0 0 0 1 1
```

```
> rle(poisson)
```

Run Length Encoding

```
 lengths: int [1:91] 1 4 1 2 2 1 2 3 1 1 ... list [[1]], [[2]]
 values  : num [1:91] 1 0 1 0 1 0 1 0 1 2 ...
```

```
> max(rle(poisson)[[1]]) Max repeat length
[1] 6
```

```
> which(rle(poisson)[[1]]==6) Position in [[1]] ... and also in [[2]]
[1] 78
```

```
> rle(poisson)[[2]][78] Value [78] in [[2]]
[1] 0
```

# Looking for runs of numbers within vectors

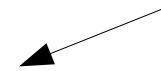
Function rle ('run length encoding')

```
> (poisson<-rpois(150,0.7))
  [1] 1 0 0 0 0 1 0 0 1 1 0 1 1 0 0 0 1 2 1 0 0 1 1 0 0 1 2 0 1 2 0 1 1 0
0 0 3 1 0 0 1 1 0 1 0 1 1 0 1 0 0 0 1
  [54] 0 0 0 2 1 1 0 0 1 1 0 1 0 0 0 2 1 0 2 2 1 1 0 1 2 1 0 0 0 0 0 1 1 2
0 2 0 0 2 0 0 0 3 0 2 1 0 3 1 0 1 1 1
 [107] 0 2 1 0 0 1 0 1 1 1 1 1 0 0 1 0 0 0 0 0 2 1 1 2 0 1 0 0 1 0 1 0 0
0 0 1 1 1 0 0 0 1 1
```

In a function:

```
> run.and.value<-function (x) {
+
+   a<- max(rle(poisson)[[1]])
+   b<-rle(poisson)[[2]][which(rle(poisson)[[1]] == a)]
+
+   cat("length = ",a," value = ",b, "\n")}
```

Value [a] in [[2]]



```
> poisson<-rpois(150,0.7)
> run.and.value(poisson)
length = 6 value = 0
```

# Saving Data Produced within R to Disc

Export a vector into a file (single column)

```
> poisson<-rpois(150,0.7)
```

```
> write(poisson, "/home/michael/ownCloud/BioInfo/poisson.txt",1)
```

# Saving Data Produced within R to Disc

Export a vector into a file (single column)

```
> poisson<-rpois(150,0.7)
```

```
> write(poisson, "/home/michael/ownCloud/BioInfo/poisson.txt",1)
```

Export a table or a matrix of numbers to file:

```
> xmat<-matrix(rpois(100000,0.75),nrow=1000)
```

```
> write.table(xmat, "/home/michael/ownCloud/BioInfo/table.txt",  
col.names=F,row.names=F)
```

# Saving Data Produced within R to Disc

Export a vector into a file (single column)

```
> poisson<-rpois(150,0.7)
> write(poisson,"/home/michael/ownCloud/BioInfo/poisson.txt",1)
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Export a table or a matrix of numbers to file:

```
> xmat<-matrix(rpois(100000,0.75),nrow=1000)
> write.table(xmat,"/home/michael/ownCloud/BioInfo/table.txt",
col.names=F,row.names=F)
```

```
> xmat.table<-table(xmat) ← builds a contingency table
```

```
> xmat.table
```

```
xmat
```

	0	1	2	3	4	5	6	7
	47289	35235	13464	3284	621	95	11	1

```
> write.table(xmat.table,"/home/michael/ownCloud/BioInfo/
xmattable.txt",col.names=F,row.names=F)
```



# Saving Data Produced within R to Disc

Export a vector into a file (single column)

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> poisson<-rpois(150,0.7)
> write(poisson,"/home/michael/ownCloud/BioInfo/poisson.txt",1)
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```
> xmat<-matrix(rpois(100000,0.75),nrow=1000)
> write.table(xmat,"/home/michael/ownCloud/BioInfo/table.txt",
col.names=F,row.names=F)
```

```
> xmat.table<-table(xmat) ← builds a contingency table
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```
> xmat.table
```

```
xmat
```

	0	1	2	3	4	5	6	7
	47289	35235	13464	3284	621	95	11	1

```
> write.table(xmat.table,"/home/michael/ownCloud/BioInfo/
xmattable.txt",col.names=F,row.names=F)
```

```
> write.table(unclass(xmat.table),"/home/michael/ownCloud/
BioInfo/xmattableu.txt",col.names=F,row.names=F)
```