## Vectors and Matrices in R

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# Creating Vectors in R

Using the "combine" operator c:

```
vd <- c(1,2,3,4)
vd
```

[1] 1 2 3 4

[1] 1 2 3 4

This is not quite the same:

```
class(vd)
```

[1] "numeric"

class(vi)

[1] "integer"

```
Creating vectors of a given size and type:
```

```
vd <- numeric(4)
vd
[1] 0 0 0 0
vi <- integer(4)</pre>
vi
[1] 0 0 0 0
vl <- logical(4)
vl
```

#### [1] FALSE FALSE FALSE

```
vc <- character(4)
vc
```

[1] "" "" "" ""

The: symbol generates a vector of a range of values:

We can also reverse:

or use variables to define the range:

### Ambiguous cases:

Often better to use seq:

$$seq(1.5, 3.5, by = 0.5)$$

$$seq(3.5, 1.5, by = -0.5)$$

$$seq(1.5, 3.5, length.out = 6)$$

[1] 1.5 1.9 2.3 2.7 3.1 3.5

## **Vector Operations**

You can find reference to most of these from ART 2.4

Vector addition adds each dimension of vectors.

```
a \leftarrow c(1,2,3,4)

b \leftarrow c(1,2,3,4)

( c \leftarrow a + b ) # Same as c \leftarrow a + b; c
```

[1] 2 4 6 8

Subtraction -, Multiplication \*, Division / work similarly.

**Do not** confuse \* with %\*%: \* is element-wise multiplication.

Logical operations performed on each element of the vector:

$$a \leftarrow c(1,2,3,4)$$
  
 $a > 2$ 

#### [1] FALSE FALSE TRUE TRUE

Many math functions operate on each element of the input vector (sin,cos,exp,log):

```
\exp(c(1,2,3,4))
```

[1] 2.718282 7.389056 20.085537 54.598150

This type of vector operations are called **vector-to-vector**.

An example of a **vector-to-scalar** computation is:

```
a \leftarrow c(1,2,3,4)
sum(a)
```

[1] 10

and an example of vector-to-matrix computation is:

```
b <- c(4,3,2,1)
tcrossprod(a, b)
```

```
[,1] [,2] [,3] [,4]
[1,] 4 3 2 1
[2,] 8 6 4 2
[3,] 12 9 6 3
[4,] 16 12 8 4
```

which is computing  $ab^T$ .

# Indexing Element(s) in a Vector

Use [] to index a single element.

### Indexing start from 1 not 0!

Indexing an element beyond the range of the vector returns NA.

[1] NA

and doing

[1] 1 2 3 4 10

works! Quite different from C.

### Generally better to allocate memory in advance:

```
tic <- Sys.time()
x <- c(1)
for(ii in 2:1e6){
   x[ii] <- ii
}
Sys.time() - tic</pre>
```

#### Time difference of 0.615715 secs

```
tic <- Sys.time()
x <- numeric(1e6)
for(ii in 2:1e6){
   x[ii] <- ii
}
Sys.time() - tic</pre>
```

Time difference of 0.1520305 secs

You can index more than one elements at a time:

Note: strictly speaking b is a vector of doubles!

Let's try:

Potentially leading to unexpected results. This is because:

```
as.integer(c(1.6, 3.6))
```

[1] 1 3

See also as.double, as.logical, as.character, ...

Use: to access multiple contiguous elements in a vector.

$$a[1:3]$$
 # same as  $a[c(1, 2, 3)]$ 

You can also use conditional expression to index a vector:

[1] 7 5

Let's break it down step by step:

$$(1 < -a > 2)$$

[1] TRUE FALSE FALSE TRUE

a[1]

[1] 7 5

```
Note: common source of errors:

1

[1] TRUE FALSE FALSE TRUE

class(1)

[1] "logical"

Suppose that at some point in your code:
```

```
1 <- as.integer(1)
1</pre>
```

[1] 1 0 0 1

Then this might not be what you expected:

```
a[1]
```

[1] 7 7

You are selecting a[1] twice, not a[c(1, 4)].

# Exploring R's documentation

How to get documentation on \*, +, : etc?

For functions we do:

```
?mean
```

This won't work:

```
?*
# Error: unexpected '*' in "?*"
```

### We need to type

```
?"*"
?":"
?"?"
?"%*%"
```

#### Matrix Construction

You can create a matrix from a vector using matrix function:

```
a <- c(1,2,3,4)
A <- matrix(a, nrow = 2)
A
```

nrow parameter specifies the number of rows in the matrix.

```
a <- c(1,2,3,4)
A <- matrix(a, nrow = 1)
A
```

By default, matrix treats your input vector 1:4 as a **column-major order**.

If you want to fill the matrix in **rows-major order**:

You can get dimension of a matrix using dim function.

[1] 2 2

Otherwise you can use ncol(A) and nrow(A).

## Indexing Elements in a Matrix

```
(A \leftarrow matrix(1:4, nrow = 2))
     [,1] [,2]
[1,] 1 3
[2,] 2 4
A[1,1]
[1] 1
A[1:2,1] # Not a matrix anymore!
[1] 1 2
You can access i-th row of a matrix by using [i,].
A[1,] # Do A[ , 1] for first column
[1] 1 3
```

```
As for vectors you can index using logical vectors:
( A \leftarrow matrix(1:9, nrow = 3) )
     [,1] [,2] [,3]
[1,] 1 4 7
[2,] 2 5 8
[3,] 3 6 9
lr <- c(TRUE, FALSE, TRUE)</pre>
A[lr, ]
     [,1] [,2] [,3]
[1,] \quad 1 \quad 4 \quad 7
[2,] 3 6 9
lc <- c(TRUE, FALSE, TRUE)</pre>
A[lr, lc]
     [,1] [,2]
[1,] 1 7
[2,] 3
```

```
To access the diagonal do:

( A <- matrix(1:9, nrow = 3) )

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

diag(A)

To over-write it:

## Matrix Operators

The +, -, \*, / symbols work in an element-wise fashion.

```
A <- matrix(1:4, nrow = 2, byrow = T)
B <- matrix(1:4, nrow = 2, byrow = T)
A + B
```

The **%\*%** symbol is used for matrix multiplication **AB**:

```
A %*% B

[,1] [,2]
[1,] 7 10
[2,] 15 22
```

### Some subtleties

An R vector is not a column vector (i.e., a matrix with 1 column):

```
A \leftarrow matrix(1:4, nrow = 2, byrow = T)
B \leftarrow matrix(1:2, nrow = 2, byrow = T)
( v \leftarrow A \% *\% B ) # NOT the same as c(5, 11)
      [,1]
[1,] 5
[2,] 11
dim(v)
[1] 2 1
(v \leftarrow as.vector(v)) # Same as c(5, 11)
[1] 5 11
dim(v) # instead, use length(v) to get number of elements
```

NULL

#### R "recycles" without telling you:

```
a \leftarrow c(1, 1, 1, 1)

b \leftarrow c(1, 2)

a * b # Same as a * c(b, b)
```

#### [1] 1 2 1 2

#### This issues a warning:

```
b2 <- c(1, 2, 3)
a * b2 # Same as a * c(b2, b2[1])
```

#### [1] 1 2 3 1

# Warning: longer object length is not a multiple of
# shorter object length

# Matrix Transposition

Matrix transpose is done by t function:

```
a \leftarrow matrix(1:6, nrow = 2)
а
    [,1] [,2] [,3]
[1,] 1 3 5
[2,] 2 4 6
t(a)
    [,1] [,2]
[1,] 1 2
[2,] 3 4
[3,] 5 6
```

Note this behaviour:

```
b \leftarrow c(1, 2, 3)
b
[1] 1 2 3
t(b)
     [,1] [,2] [,3]
[1,] 1 2
t(t(b))
     [,1]
[1,]
[2,] 2
[3,]
```

Again, in R a vector constructed with c(), a:b, numeric(n) is not the same as a matrix with one column. Big source of errors!

### Matrix Inversion

Recall given a square, full-rank matrix A,  $A^{-1}$  is such that

$$\mathbf{A}^{-1}\mathbf{A}=\mathbf{I}.$$

In R we use the solve function:

```
A <- matrix(1:4, nrow = 2)
solve(A)

[,1] [,2]
[1,] -2 1.5
[2,] 1-0.5

A %*% solve(A)
```

You should never invert a matrix unless you have to  $(\mathbf{A}\mathbf{x} = \mathbf{b})!$ 

### Conclusion

To create vectors you can use:

- ► c(x1, x2, ...)
- numeric(n), integer(n), ...
- ▶ a:b
- seq(a, b, by = eps) or seq(a, b, length.out = n)

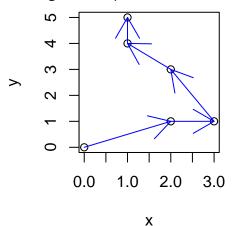
Vector and matrix indexing (or subsetting) via integer or logical vectors.

Distinction between elements-wise and vector or matrix operations.

#### Homework

Problem: given a sequence of locations in space:

- a. Find the length and direction of the segments separating each pair of locations.
- b. Find the total length of the path.



## Homework: part 1

Start by writing a function dist(a, b), where a and b are vectors represent two points in a space.

dist outputs the Euclidean distance between two points.

#### Requirements:

- ► Your function should work for input vectors in **any dimension**.
- dist(v,0) will return the length of vector v.
- No for-loops.
- Make test cases, and check your function.

## Homework: part 2

Write an R function angle, takes two vectors: a and b.

It outputs the angle between a and b in degrees.

- ▶ The result should be returned **in degrees**, not in radian.
- Use dist function you just wrote.
- Make test cases, and check your function.
- Hint: Type ?acos in command line and read the documentation.

# Homework: part 3 (submit)

Consider a sequence of n location in d-dimensional space. E.g.:

```
x y
loc1 0 0
loc2 2 1
loc3 3 1
loc4 2 3
loc5 1 4
loc6 1 5
```

Write a function that, given a matrix of locations in 2D, returns:

- 1. a vector containing the length (Euclidean norm) of each step
- 2. a vector containing the orientation of each step
  - ► for orientation you should use the angle between the x-axis and the step
- 3. a scalar representing the total length of the path

Hint: to write a function that returns several objects you can use:

```
my_function <- function(){
  a <- 1:2
  b <- "Hello!"
  return( list(a, b) )
}</pre>
```

```
x <- my_function()
x[[1]]</pre>
```

[1] 1 2

x[[2]]

[1] "Hello!"

That's all you need to know about lists for the moment!

Hint: if you want to check your code visually you can do:

```
locs \leftarrow matrix(c(0, 2, 3, 2, 1, 1,
                  0, 1, 1, 3, 4, 5), 6, 2
plot(locs)
# YOUR CODE
for(ii in list of segments){
  # YOUR CODE to compute start (x0, y0) and
  # end (x1, y1) of segments
  arrows(x0, y0, x1, y1)
```

Challenge (do not submit): Can you extend your function beyond 2D? Can you do this in C++?

Hint: in D > 2 you need more than one angle to determine the direction of a vector!