# Data Classes

#### Module 3

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June 15, 2105

#### Functions - Intro

- R revolves around functions: denoted by [function name]()
- Every function takes an input, defined by arguments, often provided by the user
- Many functions have default settings for these arguments
- example([function name]) shows you how it is used

# R Help

• If you know the name of a function, ?[function name] or help([function name]) will pop up the help menu

```
> ## ?str
> ## help("str")
```

#### **Functions - Intro**

For example, length is a function we briefly covered last module. You can try typing ?length in the console and reading the help file.

You can also see examples of running a function using example() [which is another function!]

#### Data Classes:

- One dimensional classes ('vectors'):
  - Character: strings or individual characters, quoted
  - Numeric: any real number(s)
  - Integer: any integer(s)/whole numbers
  - Factor: categorical/qualitative variables
  - Logical: variables composed of TRUE or FALSE

## **Data Classes:**

- Two dimensional classes:
  - data.frame: traditional 'Excel' spreadsheets
    - \* Each column can have a different class, from above
  - Matrix: two-dimensional data, composed of rows and columns. Unlike data frames, the entire matrix is composed of one R class, e.g. all numeric or all characters.

## Character and numeric

We have already covered character and numeric

```
> class(c("Andrew","Jaffe"))

[1] "character"
> class(c(1,4,7))

[1] "numeric"
```

Recall that c() and class() are both functions!

# Integer

Integer is a special subset of numeric that contains only whole numbers

A sequence of numbers is an example of the integer class

```
> x = seq(from=1,to=5) # seq() is a function
> x

[1] 1 2 3 4 5
> class(x)
```

[1] "integer"

# Integer

The colon: is a shortcut for making sequences of numbers. It makes consecutive integer sequence from [num1] to [num2] by 1

```
> 1:5
```

[1] 1 2 3 4 5

## **Factor**

factor are special character vectors where the elements have pre-defined groups or 'levels'. You can think of these as qualitative or categorical variables:

```
> x = factor(c("boy","girl","girl","boy","girl"))
> x

[1] boy girl girl boy girl
Levels: boy girl
```

#### > class(x)

#### [1] "factor"

Note that levels are, by default, alphabetical or alphanumerical order.

# Logical

logical is a class that only has two possible elements: TRUE and FALSE

```
> x = c(TRUE, FALSE, TRUE, TRUE, FALSE)
> class(x)
```

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

sum() and mean() work on logical vectors - they return the total and proportion of TRUE elements, respectively.

# Logical

Note that logical elements are NOT in quotes.

```
> z = c("TRUE", "FALSE", "TRUE", "FALSE")
> class(z)
```

[1] "character"

## **Vector functions**

Useful functions for exploring vectors (and other data types):

- length()
- head() and tail()
- table()
- subset() and brackets ([])
- unique()
- sum(), mean(), median(), min(), max()

## Head and Tail

- head() shows the first 6 (default) elements of an R object
- tail() shows the last 6 (default) elements of an R object
- str() shows the structure of an R object

## Head and Tail

```
> z = 1:100 # recall a sequence from 1 to 100
> head(z)

[1] 1 2 3 4 5 6
> tail(z)

[1] 95 96 97 98 99 100
> str(z)

int [1:100] 1 2 3 4 5 6 7 8 9 10 ...
```

These functions show a brief snapshot of the data which is useful for exploratory data analysis.

# **Table**

table() is the basic tabulation function, which is often more useful for character and factor vectors From the manual: "table uses the cross-classifying factors to build a contingency table of the counts at each combination of factor level"

# **Data Subsetting**

Brackets are used to select/subset/extract data in R

```
> x1 = 10:20
> x1
```

```
[1] 10 11 12 13 14 15 16 17 18 19 20
```

```
> length(x1)
```

[1] 11

# **Data Subsetting**

```
> x1[1] # selecting first element

[1] 10
> x1[3:4] # selecting third and fourth elements

[1] 12 13
> x1[c(1,5,7)] # selecting first, fifth, and seventh elements

[1] 10 14 16
```

This is probably the most powerful and useful function in R. If you master this, you can literally do anything with R. Everything in the 'data analysis pipeline' revolves around subsetting (as you will soon see)

#### **Matrices**

```
> n = 1:9 # sequence from first number to second number incrementing by 1
[1] 1 2 3 4 5 6 7 8 9
> mat = matrix(n, nrow = 3)
> mat
     [,1] [,2] [,3]
[1,]
                   7
        1
             4
[2,]
        2
             5
                   8
[3,]
        3
                   9
```

# Matrix (and Data frame) Functions

These are in addition to the previous useful vector functions: \* nrow() displays the number of rows of a matrix or data frame \* ncol() displays the number of columns \* dim() displays a vector of length 2: # rows, # columns \* colnames() displays the column names (if any) and rownames() displays the row names (if any)

# **Data Selection**

Matrices have two "slots" you can use to select data, which represent rows and columns, that are separated by a comma, so the syntax is matrix[row,column].

```
> mat[1,1] # individual entry: row 1, column 1

[1] 1
> mat[1,] # first row

[1] 1 4 7
> mat[,1] # first columns

[1] 1 2 3
```

#### **Data Selection**

Note that the class of the returned object is no longer a matrix

```
> class(mat[1,])

[1] "integer"
> class(mat[,1])

[1] "integer"
```

# **Data Frames**

The data.frame is the other two dimensional variable class.

Again, data frames are like matrices, but each column is a vector that can have its own class. So some columns might be character and others might be numeric, while others maybe a factor.

We can look at some of the example data frames that come with R.

## **Data Frames**

```
> data(iris) ## just use some data in R already
> names(iris) ## get the column names

[1] "Sepal.Length" "Sepal.Width" "Petal.Length" "Petal.Width"
[5] "Species"
```

```
> str(iris) # easy snapshot of data, like `describe` in Stata
               150 obs. of 5 variables:
'data.frame':
$ Sepal.Length: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
$ Sepal.Width: num 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
$ Petal.Length: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
 $ Petal.Width : num 0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
              : Factor w/ 3 levels "setosa", "versicolor", ...: 1 1 1 1 1 1 1 1 1 1 ...
 $ Species
> head(iris, 3) # get top 3 rows
  Sepal.Length Sepal.Width Petal.Length Petal.Width Species
          5.1
                       3.5
                                    1.4
1
                                                0.2 setosa
2
           4.9
                       3.0
                                    1.4
                                                0.2 setosa
3
          4.7
                       3.2
                                    1.3
                                                0.2 setosa
```

## **Data Selection**

Data frames have special ways to select data, specifically by a \$ and the column name.

```
> head(iris$Petal.Length)

[1] 1.4 1.4 1.3 1.5 1.4 1.7

> class(iris$Petal.Width)

[1] "numeric"
```

## **Data Selection**

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[1] 3.5 3.0 3.2 3.1 3.6 3.9

3.2 setosa

You can also subset data frames like matrices, using row and column indices, but using column names is generally safer and more reproducible.

```
> head(iris[,2])
```

You can also use the bracket notation, but specify the name(s) in quotes if you want more than 1 column. This allows you to subset rows and columns at the same time

```
> iris[1:3,c("Sepal.Width","Species")]

Sepal.Width Species
1    3.5 setosa
2    3.0 setosa
```

# **Data Frames**

You can make your own data frames from "scratch" too, either from a matrix or using the data.frame function:

## **Data Frames**

> names(iris2)

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You can add variables to a data.frame using \$ as well:

```
> iris2 = iris # copy `iris` to a new df
> iris2$Index=1:nrow(iris2)
> head(iris2)
  Sepal.Length Sepal.Width Petal.Length Petal.Width Species Index
1
           5.1
                       3.5
                                    1.4
                                                0.2 setosa
           4.9
                       3.0
                                                0.2 setosa
                                                                2
2
                                    1.4
3
           4.7
                       3.2
                                    1.3
                                                0.2 setosa
                                                                3
           4.6
                                                0.2 setosa
4
                       3.1
                                    1.5
                                                                4
5
           5.0
                       3.6
                                    1.4
                                                0.2 setosa
                                                                5
6
           5.4
                       3.9
                                                0.4 setosa
                                                                6
                                    1.7
```

```
[1] "Sepal.Length" "Sepal.Width" "Petal.Length" "Petal.Width" [5] "Species" "Index"
```

# Data Classes: Extended

Extensions of "normal" data classes:

- N-dimensional classes:
  - Arrays: any extension of matrices with more than 2 dimensions, e.g. 3x3x3 cube
  - Lists: more flexible container for R objects.