



# Lecture 1: Introduction to the Course Basics of Data

S. Morteza Najibi

Shiraz University

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A decorative background pattern consisting of a grid of squares, each divided into four triangles by diagonal lines. The pattern is rendered in a light gray color on a dark gray background.

# Introduction

# Agenda



- ▶ Course overview and mechanics
- ▶ Built-in data types
- ▶ Built-in functions and operators
- ▶ First data structures: Vectors and arrays

# Why good statisticians learn to program



- ▶ *Independence*: Otherwise, you rely on someone else having given you exactly the right tool
- ▶ *Honesty*: Otherwise, you end up distorting your problem to match the tools you have
- ▶ *Clarity*: Making your method something a machine can do disciplines your thinking and makes it public; that's science

# How this class will work



- ▶ No programming knowledge presumed
- ▶ Some stats. knowledge presumed
- ▶ General programming mixed with data-manipulation and statistical inference
- ▶ Class will be *very* cumulative
- ▶ Keep up with the readings and assignments!

# assignment



Assignments, class notes, grading policies, useful links on R:

▶ <http://cw.sdat.ir>



# R Basics

# The class in a nutshell: Functional programming



2 sorts of things (**objects**): **data** and **functions**

▶ **Data:** things like 7, “seven”, 7.000, the matrix  $\begin{bmatrix} 7 & 7 & 7 \\ 7 & 7 & 7 \end{bmatrix}$

▶ **Functions:** things like `log`, `+` (two arguments), `<` (two), `mod` (two), `mean` (one)

*A function is a machine which turns input objects (**arguments**) into an output object (**return value**), possibly with **side effects**, according to a definite rule*



Programming is writing functions to transform inputs into outputs

Good programming ensures the transformation is done easily and correctly

Machines are made out of machines; functions are made out of functions, like

$$f(a, b) = a^2 + b^2$$

*The route to good programming is to take the big transformation and break it down into smaller ones, and then break those down, until you come to tasks which the built-in functions can do*

# Before functions, data

Different kinds of data object

All data is represented in binary format, by **bits** (TRUE/FALSE, YES/NO, 1/0)

- ▶ **Booleans** Direct binary values: **TRUE** or **FALSE** in R
- ▶ **Integers**: whole numbers (positive, negative or zero), represented by a fixed-length block of bits
- ▶ **Characters** fixed-length blocks of bits, with special coding; **strings** = sequences of characters
- ▶ **Floating point numbers**: a fraction (with a finite number of bits) times an exponent, like  $1.87 \times 10^6$ , but in binary form
- ▶ **Missing or ill-defined values**: **NA**, **NaN**, etc.

# Operators



- ▶ **Unary** - for arithmetic negation, ! for Boolean
- ▶ **Binary** usual arithmetic operators, plus ones for modulo and integer division; take two numbers and give a number

## some basics



```
7+5
```

```
## [1] 12
```

```
7-5
```

```
## [1] 2
```

```
7*5
```

```
## [1] 35
```

```
7^5
```

```
## [1] 16807
```

```
7/5
```

```
## [1] 1.4
```

```
7 %% 5
```

```
## [1] 2
```

```
7 %/% 5
```

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

# The R Console



Basic interaction with R is by typing in the **console**, a.k.a. **terminal** or **command-line**

You type in commands, R gives back answers (or errors)

Menus and other graphical interfaces are extras built on top of the console

# Operators cont'd.

**Comparisons** are also binary operators; they take two objects, like numbers, and give a Boolean

```
7 > 5
```

```
## [1] TRUE
```

```
7 < 5
```

```
## [1] FALSE
```

```
7 >= 7
```

```
## [1] TRUE
```

```
7 <= 5
```

```
## [1] FALSE
```

```
7 == 5
```

```
## [1] FALSE
```

```
7 != 5
```

```
## [1] TRUE
```



# Boolean operators

Basically “and” and “or”:

```
(5 > 7) & (6*7 == 42)
```

```
## [1] FALSE
```

```
(5 > 7) | (6*7 == 42)
```

```
## [1] TRUE
```

(will see special doubled forms, `&&` and `||`, later)

# More types

`typeof()` function returns the type

`is.foo()` functions return Booleans for whether the argument is of type *foo*

`as.foo()` (tries to) “cast” its argument to type *foo* — to translate it sensibly into a *foo*-type value

```
typeof(7)
```

```
## [1] "double"
```

```
is.numeric(7)
```

```
## [1] TRUE
```

```
is.na(7)
```

```
## [1] FALSE
```

```
is.na(7/0)
```

```
## [1] FALSE
```

```
is.na(0/0)
```

```
## [1] TRUE
```

Why is  $7/0$  not NA, but  $0/0$  is?

```
is.character(7)
```

```
## [1] FALSE
```

```
is.character("7")
```

```
## [1] TRUE
```

```
is.character("seven")
```

```
## [1] TRUE
```

```
is.na("seven")
```

```
## [1] FALSE
```

```
as.character(5/6)
```

```
## [1] "0.833333333333333"
```

```
as.numeric(as.character(5/6))
```

```
## [1] 0.8333333
```

```
6*as.numeric(as.character(5/6))
```

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

```
5/6 == as.numeric(as.character(5/6))
```

```
## [1] FALSE
```

(why is that last FALSE?)

# Data can have names

We can give names to data objects; these give us **variables**

A few variables are built in:

```
pi
```

```
## [1] 3.141593
```

Variables can be arguments to functions or operators, just like constants:

```
pi*10
```

```
## [1] 31.41593
```

```
cos(pi)
```

```
## [1] -1
```

Most variables are created with the **assignment operator**, `<-` or `=`

```
approx.pi <- 22/7  
approx.pi
```

```
## [1] 3.142857
```

```
diameter.in.cubits = 10  
approx.pi*diameter.in.cubits
```

```
## [1] 31.42857
```



# The assignment operator also changes values:

```
circumference.in.cubits <- approx.pi*diameter.in.cubits  
circumference.in.cubits
```

```
## [1] 31.42857
```

```
circumference.in.cubits <- 30  
circumference.in.cubits
```

```
## [1] 30
```



# Workspace

# Using Names



Using names and variables makes code: easier to design, easier to debug, less prone to bugs, easier to improve, and easier for others to read

Avoid “magic constants”; use named variables you will be graded on this!

Named variables are a first step towards **abstraction**

# The workspace



What names have you defined values for?

```
ls()
```

```
## [1] "approx.pi"                "circumference.in.cubits"  
## [3] "diameter.in.cubits"
```

```
objects()
```

```
## [1] "approx.pi"                "circumference.in.cubits"  
## [3] "diameter.in.cubits"
```

Getting of variables:

```
rm("circumference.in.cubits")  
ls()
```

```
## [1] "approx.pi"                "diameter.in.cubits"
```

# First data structure: vectors



Group related data values into one object, a **data structure**

A **vector** is a sequence of values, all of the same type

```
x <- c(7, 8, 10, 45)
x
```

```
## [1] 7 8 10 45
```

```
is.vector(x)
```

```
## [1] TRUE
```

`c()` function returns a vector containing all its arguments in order `x[1]` is the first element, `x[4]` is the 4th element

`x[-4]` is a vector containing all but the fourth element

`vector(length=6)` returns an empty vector of length 6; helpful for filling things up later

```
weekly.hours <- vector(length=5)  
weekly.hours[5] <- 8
```

# Vector arithmetic



Operators apply to vectors “pairwise” or “elementwise”:

```
y <- c(-7, -8, -10, -45)
x+y
```

```
## [1] 0 0 0 0
```

```
x*y
```

```
## [1] -49 -64 -100 -2025
```

# Recycling



**Recycling** repeat elements in shorter vector when combined with longer

```
x + c(-7,-8)
```

```
## [1] 0 0 3 37
```

```
x^c(1,0,-1,0.5)
```

```
## [1] 7.000000 1.000000 0.100000 6.708204
```

Single numbers are vectors of length 1 for purposes of recycling:

```
2*x
```

```
## [1] 14 16 20 90
```



Can also do pairwise comparisons:

```
x > 9
```

```
## [1] FALSE FALSE TRUE TRUE
```

Note: returns Boolean vector

Boolean operators work elementwise:

```
(x > 9) & (x < 20)
```

```
## [1] FALSE FALSE TRUE FALSE
```

To compare whole vectors, best to use `identical()` or `all.equal()`:

```
x == -y
```

```
## [1] TRUE TRUE TRUE TRUE
```

```
identical(x,-y)
```

```
## [1] TRUE
```

```
identical(c(0.5-0.3,0.3-0.1),c(0.3-0.1,0.5-0.3))
```

```
## [1] FALSE
```

```
all.equal(c(0.5-0.3,0.3-0.1),c(0.3-0.1,0.5-0.3))
```

```
## [1] TRUE
```

The background of the slide is a dark blue color with a subtle, repeating geometric pattern of light blue triangles. The triangles are arranged in a way that creates a sense of depth and structure, with some pointing upwards and others downwards, forming a larger, more complex pattern.

# Simple Functions

# Functions on vectors



Lots of functions take vectors as arguments: - `mean()`, `median()`, `sd()`, `var()`, `max()`, `min()`, `length()`, `sum()`: return single numbers - `sort()` returns a new vector - `hist()` takes a vector of numbers and produces a histogram, a highly structured object, with the side-effect of making a plot - Similarly `ecdf()` produces a cumulative-density-function object - `summary()` gives a five-number summary of numerical vectors - `any()` and `all()` are useful on Boolean vectors

# Addressing vectors



Vector of indices:

```
x[c(2,4)]
```

```
## [1] 8 45
```

Vector of negative indices

```
x[c(-1,-3)]
```

```
## [1] 8 45
```

(why that, and not 8 10?)

# Boolean vector

```
x[x>9]
```

```
## [1] 10 45
```

```
y[x>9]
```

```
## [1] -10 -45
```

`which()` turns a Boolean vector in vector of TRUE indices:

```
places <- which(x > 9)  
places
```

```
## [1] 3 4
```

```
y[places]
```

```
## [1] -10 -45
```

# Named components

You can give names to elements or components of vectors

```
names(x) <- c("v1", "v2", "v3", "fred")  
names(x)
```

```
## [1] "v1"    "v2"    "v3"    "fred"
```

```
x[c("fred", "v1")]
```

```
## fred    v1  
##    45     7
```

note the labels in what R prints; not actually part of the value

`names(x)` is just another vector (of characters):

```
names(y) <- names(x)
sort(names(x))
```

```
## [1] "fred" "v1"    "v2"    "v3"
```

```
which(names(x)=="fred")
```

```
## [1] 4
```



# Take-Aways



- ▶ We write programs by composing functions to manipulate data
- ▶ The basic data types let us represent Booleans, numbers, and characters
- ▶ Data structure let us group related values together
- ▶ Vectors let us group values of the same type
- ▶ Use variables rather a profusion of magic constants
- ▶ Name components of structures to make data more meaningful

# Peculiarities of floating-point numbers



The more bits in the fraction part, the more precision

The R floating-point data type is a **double**, a.k.a. **numeric** back when memory was expensive, the now-standard number of bits was twice the default

Finite precision  $\Rightarrow$  arithmetic on **doubles**  $\neq$  arithmetic on  $\mathbb{R}$ .

```
0.45 == 3*0.15
```

```
## [1] FALSE
```

```
0.45 - 3*0.15
```

```
## [1] 5.551115e-17
```

Often ignorable, but not always - Rounding errors tend to accumulate in long calculations - When results should be  $\approx 0$ , errors can flip signs - Usually better to use `all.equal()` than exact comparison

```
(0.5 - 0.3) == (0.3 - 0.1)
```

```
## [1] FALSE
```

```
all.equal(0.5-0.3, 0.3-0.1)
```

```
## [1] TRUE
```



# Peculiarities of Integers

Typing a whole number in the terminal doesn't make an integer; it makes a double, whose fractional part is 0

```
is.integer(7)
```

```
## [1] FALSE
```

This looks like an integer

```
as.integer(7)
```

```
## [1] 7
```

To test for being a whole number, use `round()`:

```
round(7) == 7
```

```
## [1] TRUE
```

A decorative background pattern consisting of a grid of squares, each divided into four triangles by diagonal lines. The pattern is rendered in a light gray color on a dark gray background.

**Thank You**