

<https://intro2r.com/> Chapter 2

CSCI 297b, Spring 2023

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R basics

- R is case sensitive. `anova` is not the same as `Anova`
- Anything following `#` is a comment and is ignored by R
- Comments should be used liberally
- Commands are separated by a newline or a semicolon ;
- A continuation prompt, `+`, means the previous line is not finished
- If execution hangs and does not stop, try the escape key or the stop button

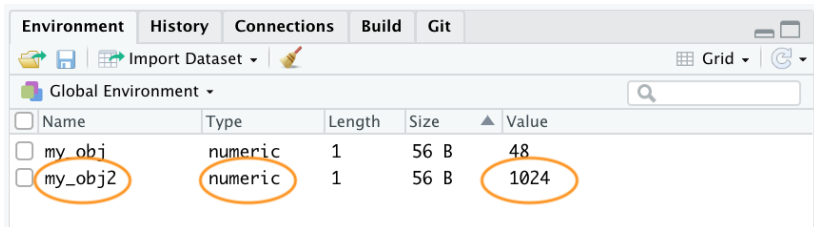
Some builtin R functions

```
log(1)           # logarithm to base e
## [1] 0
log10(1)         # logarithm to base 10
## [1] 0
exp(1)           # natural antilog
## [1] 2.718282
sqrt(4)          # square root
## [1] 2
4^2              # 4 to the power of 2
## [1] 16
pi               # not a function but useful
## [1] 3.141593
```

Objects and assignment

```
> my_obj <- 1729
> my_obj2 <- "R is cool"
> my_obj
[1] 1729
> my_obj3 <- my_obj / 2
> my_obj3
[1] 864.5
> my_obj4 <- my_obj + my_obj3
> my_obj4
[1] 2593.5
> my_obj5 <- my_obj + my_obj2
Error in my_obj + my_obj2 : non-numeric argument to binary operator
>
```

The Environment Tab



The screenshot shows the RStudio Environment Tab with the following table of objects:

<input type="checkbox"/>	Name	Type	Length	Size	▲	Value
<input type="checkbox"/>	my_obj	numeric	1	56 B		48
<input type="checkbox"/>	my_obj2	numeric	1	56 B		1024

The values 1024 and the word 'numeric' are circled in orange in the original image.

Naming Objects

There are two hard problems in computer science: cache invalidation, naming things, and off-by-1 errors.

— Leon Bambrick

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Two often conflicting goals:

- Short
- Meaningful

Name conventions

```
output_summary <- "my analysis"      # snake case
output.summary <- "my analysis"      # dot case
outputSummary <- "my analysis"       # camel case
OutputSummary <- "my analysis"       # Pascal case
output-summary <- "my analysis"      # kebab case
```

- Snake case used by textbook
- Google style recommends Pascal for function names
- Kebab case is illegal in R
- Dots illegal in many other languages
- Camel case is my favorite

Don't use existing names

```
data <- read.table("mydatafile", header = TRUE)  
#data is a function!
```

Do exercise 2, part 1

The `c()` function

```
my_vec <- c(2,3,1,6,4,3,3,7)
mean(my_vec)      # returns the mean of my_vec
## [1] 3.625
var(my_vec)       # returns the variance of my_vec
## [1] 3.982143
sd(my_vec)        # returns the standard deviation of my_vec
## [1] 1.995531
length(my_vec)    # returns the number of elements in my_vec
## [1] 8
```

Sequences

```
my_seq <- 1:10      # create regular sequence
my_seq
## [1] 1 2 3 4 5 6 7 8 9 10
my_seq2 <- 10:1     # in decending order
my_seq2
## [1] 10 9 8 7 6 5 4 3 2 1
my_seq2 <- seq(from = 1, to = 5, by = 0.5)
my_seq2
## [1] 1.0 1.5 2.0 2.5 3.0 3.5 4.0 4.5 5.0
my_seq3 <- rep(2, times = 10)  # repeats 2, 10 times
my_seq3
## [1] 2 2 2 2 2 2 2 2 2 2
my_seq4 <- rep("abc", times = 3)  # repeats 'abc' 3 times
my_seq4
## [1] "abc" "abc" "abc"
```

Sequences

```
my_seq5 <- rep(1:5, times = 3)  # repeats the series 1 to  
                                # 5, 3 times
```

```
my_seq5
```

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

```
my_seq6 <- rep(1:5, each = 3)   # repeats each element of  
                                # the series 3 times
```

```
my_seq6
```

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

```
my_seq7 <- rep(c(3, 1, 10, 7), each = 3) # repeats each  
                                           # element of the  
                                           # series 3 times
```

```
my_seq7
```

```
## [1] 3 3 3 1 1 1 10 10 10 7 7 7
```

```
## Alternative approach:
```

```
in_vec <- c(3, 1, 10, 7)
```

```
my_seq7 <- rep(in_vec, each = 3)
```

```
my_seq7
```

```
## [1] 3 3 3 1 1 1 10 10 10 7 7 7
```

Positional indexing

```
my_vec          # remind ourselves what my_vec looks like
## [1] 2 3 1 6 4 3 3 7
my_vec[3]       # extract the 3rd value
## [1] 1

# if you want to store this value in another object
val_3 <- my_vec[3]
val_3
## [1] 1
my_vec[c(1, 5, 6, 8)]
## [1] 2 4 3 7
my_vec[3:8]
## [1] 1 6 4 3 3 7
```

Logical indexing

```
my_vec          # remind ourselves what my_vec looks like
## [1] 2 3 1 6 4 3 3 7
my_vec[my_vec > 4]
## [1] 6 7
my_vec > 4
## [1] FALSE FALSE FALSE  TRUE FALSE FALSE FALSE  TRUE
my_vec[c(FALSE, FALSE, FALSE, TRUE, FALSE, FALSE, FALSE, TRUE)]
## [1] 6 7
```

Logical indexing

```
my_vec          # remind ourselves what my_vec looks like
## [1] 2 3 1 6 4 3 3 7
my_vec[my_vec >= 4]      # values greater or equal to 4
## [1] 6 4 7
my_vec[my_vec < 4]       # values less than 4
## [1] 2 3 1 3 3
my_vec[my_vec <= 4]      # values less than or equal to 4
## [1] 2 3 1 4 3 3
my_vec[my_vec == 4]      # values equal to 4
## [1] 4
my_vec[my_vec != 4]      # values not equal to 4
## [1] 2 3 1 6 3 3 7
```


Boolean expressions

```
my_vec          # remind ourselves what my_vec looks like
## [1] 2 3 1 6 4 3 3 7
val26 <- my_vec[my_vec < 6 & my_vec > 2]
val26
## [1] 3 4 3 3
val63 <- my_vec[my_vec > 6 | my_vec < 3]
val63
## [1] 2 1 7
```

Replacing elements

```
my_vec          # remind ourselves what my_vec looks like
## [1] 2 3 1 6 4 3 3 7
```

```
## replace the 4th element with 500
```

```
my_vec[4] <- 500
```

```
my_vec
```

```
## [1] 2 3 1 500 4 3 3 7
```

```
# replace the 6th and 7th element with 100
```

```
my_vec[c(6, 7)] <- 100
```

```
my_vec
```

```
## [1] 2 3 1 500 4 100 100 7
```

```
# replace element that are less than or equal to 4 with 1000
```

```
my_vec[my_vec <= 4] <- 1000
```

```
my_vec
```

```
## [1] 1000 1000 1000 500 1000 100 100 7
```

Do exercise 2, part 2

Sorting elements

```
my_vec
## [1] 1000 1000 1000 500 1000 100 100 7

vec_sort <- sort(my_vec)
vec_sort
## [1] 7 100 100 500 1000 1000 1000 1000

vec_sort2 <- sort(my_vec, decreasing = TRUE)
vec_sort2
## [1] 1000 1000 1000 1000 500 100 100 7

vec_sort3 <- rev(sort(my_vec))
vec_sort3
## [1] 1000 1000 1000 1000 500 100 100 7
```

Ordering elements

```
height <- c(180, 155, 160, 167, 181)
```

```
height
```

```
## [1] 180 155 160 167 181
```

```
p.names <- c("Joanna", "Charlotte", "Helen", "Karen", "Amy")
```

```
p.names
```

```
## [1] "Joanna"      "Charlotte" "Helen"      "Karen"      "Amy"
```

```
height_ord <- order(height)
```

```
height_ord
```

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

```
height[height_ord]
```

```
## [1] 155 160 167 180 181
```

```
names_ord <- p.names[height_ord]
```

```
names_ord
```

```
## [1] "Charlotte" "Helen"      "Karen"      "Joanna"     "Amy"
```

Vectorization

```
# create a vector
my_vec2 <- c(3, 5, 7, 1, 9, 20)

# multiply each element by 5
my_vec2 * 5
## [1] 15 25 35 5 45 100

# create a second vector
my_vec3 <- c(17, 15, 13, 19, 11, 0)

# add both vectors
my_vec2 + my_vec3
## [1] 20 20 20 20 20 20

# multiply both vectors
my_vec2 * my_vec3
## [1] 51 75 91 19 99 0
```

Vectorization recycling

```
my_vec2 <- c(3, 5, 7, 1, 9, 20)
my_vec4 <- c(1, 2)

# add both vectors - quiet recycling!
my_vec2 + my_vec4
## [1] 4 7 8 3 10 22
```

Missing data

```
temp <- c(7.2, NA, 7.1, 6.9, 6.5, 5.8, 5.8, 5.5, NA, 5.5)
temp
## [1] 7.2 NA 7.1 6.9 6.5 5.8 5.8 5.5 NA 5.5
```

```
mean_temp <- mean(temp)
mean_temp
## [1] NA
```

```
mean_temp <- mean(temp, na.rm = TRUE)
mean_temp
## [1] 6.2875
```

- Some functions may deal with NAs differently.
- Always consult the documentation.

R help

```
## show the help page:  
help("mean")  
?mean
```

```
## search all help pages:  
help.search("mean")  
??mean
```

- Also use the Help tab in RStudio
- Usually the most helpful part of a help page is the examples.

R help

```
apropos("mean")  
## [1] ".colMeans"      ".rowMeans"      "colMeans"      "kmeans"  
## [5] "mean"           "mean_temp"     "mean.Date"     "mean.de  
## [9] "mean.difftime"  "mean.POSIXct"  "mean.POSIXlt"  "rowMean  
## [13] "vec_mean"       "weighted.mean"
```

- Find all functions with "mean" in their name.

```
RSiteSearch("regression")
```

- Search for keywords and phrases in function help pages and vignettes for all CRAN packages, and in CRAN task views.

General R resources

- <https://cran.r-project.org/other-docs.html>
R-Project: User contributed documentation
- <https://journal.r-project.org/>
The R Journal: Journal of the R project for statistical computing
- <http://swirlstats.com/>
Swirl: An R package that teaches you R from within R
- <https://www.rstudio.com/resources/cheatsheets/>
RStudio's printable cheatsheets
- <http://rseek.org/>
Rseek A custom Google search for R-related sites

Getting help

- <https://www.google.com/>

Google it!: Try Googling any error messages you get. It's not cheating and everyone does it! You'll be surprised how many other people have probably had the same problem and solved it.

- <http://stackoverflow.com/questions/tagged/r>

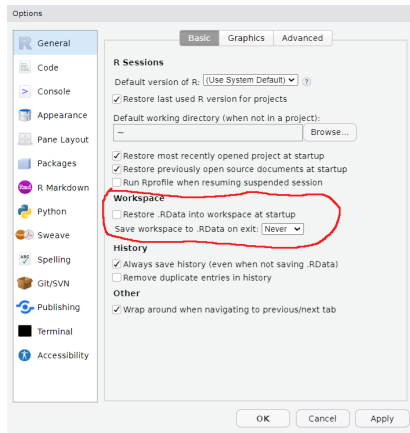
Stack Overflow: There are many thousands of questions relevant to R on Stack Overflow. Here are the most popular ones, ranked by vote. Make sure you search for similar questions before asking your own, and make sure you include a reproducible example to get the most useful advice. A reproducible example is a minimal example that lets others who are trying to help you to see the error themselves.

Saving stuff

- Normally the only thing you need to save is your R script.
- Put everything you do in the script.
- Save it, and the next time you start just source it.

Global options

- Tools → Global Options ...
- Turn off .RData save and restore
- This prevents new R sessions from being influenced by previous R sessions.



Saving objects

- Occasionally you want to save an object rather than recompute it.
- Some objects may take minutes or hours to compute.

```
## save a single object to a file  
save(nameOfObject, file = "name_of_file.RData")
```

```
## save all the objects in your workspace in a single file  
save.image(file = "name_of_file.RData")
```

```
## reload whatever you saved  
load(file = "name_of_file.RData")
```

Do exercise 2, part 3