R Reference

Basic Operations

- Use variable <- value to assign a value to a variable.
- # starts a comment.
- Statements in a block must be surrounded by curly braces { }.
- ?thing displays the help for thing.
- length(thing) produces the length of a collection.
- c(value1, value2, value3, ...) creates a vector.
- vector_name[i] selects the i'th value from a vector.
- mean, max, and min() calculate simple statistics.
- plot creates simple visualizations.
- list.files(pattern = "txt") returns the names of all files that contain "txt" in their name.

Data frames

- dim(dat) gives the dimensions of a data frame.
- dat[x, y] selects a single element from a data frame.
- dat[i,] selects the i'th row; dat[, i] selects the i'th column.
- low:high specifies a slice including elements from low to high.
- apply(dat, 1, mean) calculates the mean of each row.
- apply(dat, 2, mean) calculates the mean of each column.

Functions

- name <- function(...args...) defines a new function.
- name <- function(arg = default) specifies a default value for a parameter.
- Call a function using name(...values...).

Control Flow

• Create a for loop to process elements in a collection one at a time:

```
for (variable in collection) {
    ...body...
}
```

• Create a conditional using if and else:

```
if (condition_1) {
    ...body...
} else if (condition_2) {
    ...body...
} else {
    ...body...
}
```

• Use == to test for equality.

- X & Y is only true if both X and Y are true.
- X | Y is true if either X or Y, or both, are true.
- Use stopifnot(condition) to check that something is true when the program is running.

Using R from the command-line

- commandArgs(trailingOnly = TRUE) returns the command-line arguments.
- file("stdin") reads from standard input.
- cat(vec, sep = "\n") writes to standard ouput each element of vec on its own line.