



Functions



Functions are created using the function () directive and are stored as R objects just like anything else. In particular, they are R objects of class of function".

```
f <- function(<arguments>) {
     ## Do something interesting
}
```



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Functions



Functions in R are "first class objects", which means that they can be treated much like any other R object. Importantly,

- Functions can be passed as arguments to other functions
- Functions can be nested, so that you can define a function inside of another function
- The return value of a function is the last expression in the function body to be evaluated.



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Function Arguments



Functions have *named arguments* which potentially have *default values*.

- The formal arguments are the arguments included in the function definition
- The formals function returns a list of all the formal arguments of a function
- Not every function call in R makes use of all the formal arguments
- Function arguments can be missing or might have default values



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Argument Matching



R functions arguments can be matched positionally or by name. So the following calls to ${\tt sd}$ are all equivalent

```
> mydata <- rnorm(100)
> sd(mydata)
> sd(x = mydata)
> sd(x = mydata, na.rm = FALSE)
> sd(na.rm = FALSE, x = mydata)
> sd(na.rm = FALSE, mydata)
```

Even though it's legal, it is not recommend to mess around with the order of the arguments too much, since it can lead to some confusion.

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Argument Matching



You can mix positional matching with matching by name. When an argument is matched by name, it is "taken out" of the argument list and the remaining unnamed arguments are matched in the order that they are listed in the function definition.

```
> args(lm)
function (formula, data, subset, weights, na.action, method = "qr",
    model = TRUE, x = FALSE, y = FALSE, qr = TRUE, singular.ok = TRUE,
    contrasts = NULL, offset, ...)
NULL
```

The following two calls are equivalent.

```
> lm(data = mydata, y~x, model = FALSE, 1:100)
> lm(y~x, mydata, 1:100, model = FALSE)
```

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Argument Matching



- Most of the time, named arguments are useful on the command line when you have a long argument list and you want to use the defaults for everything except for an argument near the end of the list
- Named arguments also help if you can remember the name of the argument and not its position on the argument list (plotting is a good example).

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Argument Matching



Function arguments can also be *partially* matched, which is useful for interactive work. The order of operations when given an argument is

- 1. Check for exact match for a named argument
- 2. Check for a partial match
- 3. Check for a positional match



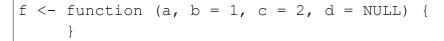
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Defining a Function





In addition to not specifying a default value, you can also set an argument value to ${\tt NULL}$.



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Lazy Evaluation



Arguments to functions are evaluated *lazily*, so they are evaluated only as needed.

```
f <- function(a,b){
    return(a^2)
}
> f(2)
[1] 4
```

This function never actually uses the argument b, so calling f(2) will not produce an error because the 2 gets positionally matched to a.

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Lazy Evaluation



```
f <- function (a, b) {
  print(a)
  print(b)
}</pre>
```

```
> f(45)
[1] 45
Error in print(b) : argument "b" is missing, with no default
```

Notice that "45" got printed first before the error was triggered. This is because b did not have to be evaluated until after print(a). Once the function tried to evaluate print(b) it had to throw an error.

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The "..." Argument



The ... argument indicate a variable number of arguments that are usually passed on to other functions.

• ... is often used when extending another function and you don't want to copy the entire argument list of the original function

```
myplot <- function(x, y, type = "]", ...){
  plot(x,y, type = type, ...)
}</pre>
```

• Generic functions use ... so that extra arguments can be passed to methods (more on this later)

```
> mean
function (x, ...)
UseMethod("mean")
<bytecode: 0x00000000391e720>
<environment: namespace:base>
```

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The "..." Argument



The ... argument is also necessary when the number of arguments passed to the function cannot be known in advance.

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Arguments Coming After the "..." Argument



One catch with ... is that any arguments that appear *after* ... on the argument list must be named explicitly and cannot be partially matched.

```
> args(paste)
function (..., sep = " ", collapse = NULL)
NULL
> paste("a", "b", sep = ":")
[1] "a:b"
> paste("a", "b", se = ":")
[1] "a b :"
```





Important Notes in Creating Functions



- Make sure all arguments are properly assigned
- The body should contain all necessary steps to be done by R
- return ([object]) function is used to provide a returning value for the function



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Try these!



- 1. Create a function named "square" that produces the squared value of a numeric argument.
- 2. Create a function named "sqroot" that produces the square root of a valid numerical argument but produces the message "ERROR: Invalid argument" otherwise.
- 3. Create a function named "sq" that prints every perfect squares until the specified number of observations (n), e.g. 1,2,9,..., n²



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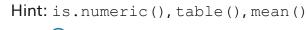
Try these!



4. Create a function named "gendata" that produces a data frame with predetermined number of observations (n) that contains the variable "prev" where "prev" has a predermined starting value (s) with the next value as half of the previous value, i.e.,

new value = 0.5 * previous value

5. Create a function named "summ" that prints the mean of a vector argument but gives number of observation per level when a categorical vector argument is used.





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A Diversion on Binding Values to Symbol



How does R know which value to assign to which symbol? When I type,

```
> lm <- function(x){x*x}
> lm
function(x){x*x}
```

how does R know what value to assign to the symbol lm? Why doesn't it give it the value of lm that is in the **stats** package?



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A Diversion on Binding Values to **Symbol**



When R tries to bind a value to a symbol, it searches through a series of environments to find the appropriate value. When you are working on the command line and need to retrieve the value of an R object, the order is roughly

- 1. Search the global environment for a symbol name matching the one requested.
- 2. Search the namespaces of each of the packages on the search list

The search list can be found by using the search function.

```
search()
   ".GlobalEnv"
```

"package:graphics" "package:datasets"

"package:base"

"tools:rstudio" "package:grDevices" "package:utils"

"package:methods"

"Autoloads"

"package:stats"

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Binding Values to Symbol



- The global environment or the user's workspace is always the first element of the search list and the base package is always the last.
- The order of the packages on the search list matters!
- User's can configure which packages get loaded on startup so you cannot assume that there will be a set list of packages available.
- When a user loads a package with library the namespace of that package gets put in position 2 of the search list (by default) and everything else gets shifted down the list.
- Note that R has separate namespaces for functions and nonfunctions so it's possible to have an object named c and a function named c.





Scoping Rules



The scoping rules for R are the main feature that make it different from the original S language.

- The scoping rules determine how a value is associated with a free variable in a function
- R uses lexical scoping or static scoping. A common alternative is dynamic scoping.
- Related to the scoping rules is how R uses the search list to bind a value to a symbol
- Lexical scoping turns out to be particularly useful for simplifying statistical computations

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Lexical Scoping



Consider the following function.

```
f <- function(x,y) {
   x^2 + y / z
}</pre>
```

This function has 2 formal arguments x and y. In the body of the function there is another symbol z. In this case z is called a **free variable**. The scoping rules of a language determine how values are assigned to free variables. Free variables are not formal arguments and are not local variables (assigned inside the function body).

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Lexical Scoping



Lexical Scoping in R means that

the values of free variables are searched for in the environment in which the function was defined.

What is an environment?

- An environment is a collection of (symbol, value) pairs, i.e. x is a symbol and 3.14 might be its value.
- Every environment has a parent environment; it is possible for an environment to have multiple "children"
- the only environment without a parent is the empty environment
- A function + an environment = a closure or function closure.



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Lexical Scoping



Searching for the value for a free variable:

- If the value of a symbol is not found in the environment in which a function was defined, then the search is continued in the parent environment.
- The search continues down the sequence of parent environments until we hit the top-level environment; this usually the global environment (workspace) or the namespace of a package.
- After the top-level environment, the search continues down the search list until we hit the empty environment. If a value for a given symbol cannot be found once the empty environment is arrived at, then an error is thrown.







Lexical Scoping



Why does all this matter?

- Typically, a function is defined in the global environment, so that the values of free variables are just found in the user's workspace
- This behavior is logical for most people and is usually the "right thing" to do
- However, in R you can have functions defined inside other functions
 Languages like C don't let you do this
- Now things get interesting In this case the environment in which a function is defined is the body of another function!



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Lexical Scoping

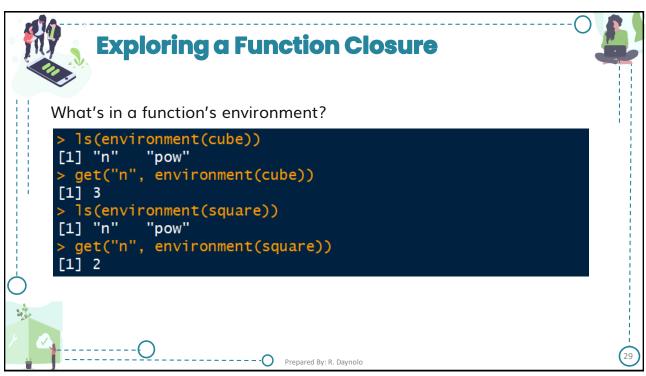


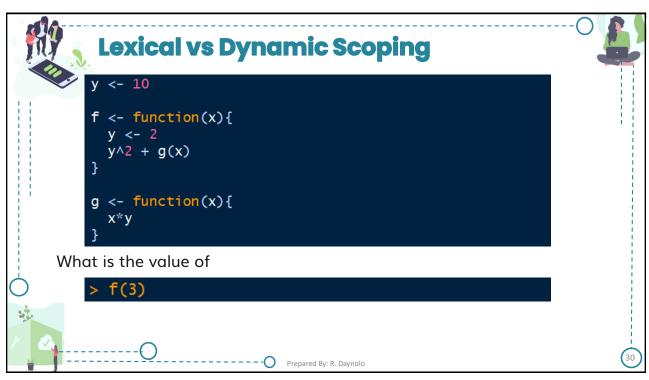
```
make.power <- function(n) {
   pow <- function(x) {
      x^n
   }
   pow
}</pre>
```

This function returns another function as its value

```
> cube <- make.power(3)
> square <- make.power(2)
> cube(3)
[1] 27
> square(3)
[1] 9
```

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Lexical vs Dynamic Scoping



- With lexical scoping the value of y in the function g is looked up in the environment in which the function was defined, in this case the global environment, so the value of y is 10.
- With dynamic scoping, the value of y is looked up in the environment from which the function was called (sometimes referred to as the *calling environment*).
 - In R the calling environment is known as the *parent frame*
- So the value of y would be 2.



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Lexical vs Dynamic Scoping



When a function is *defined* in the global environment and is subsequently *called* from the global environment, then the defining environment and the calling environment are the same. This can sometimes give the appearance of dynamic scoping.

```
> g <- function(x) {
+ a <- 3
+ x + a + y
+ }
> g(2)
Error in g(2) : object 'y' not found
> y <- 3
> g(2)
[1] 8
```



Other Languages



Other languages that support lexical scoping

- Scheme
- Perl
- Python
- Common Lisp (all languages converge to Lisp)



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Consequences of Lexical Scoping



- In R, all objects must be stored in memory
- All functions must carry a pointer to their respective defining environments, which could be anywhere
- In S-PLUS, free variables are always looked up in the global workspace, so everything can be stored on the disk because the "defining environment" of all functions is the same.



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Application: Optimization



Why is any of this information useful?

- Optimization routines in R like optim, nlm, and optimize require you to pass a function whose argument is a vector of parameters (e.g. a log-likelihood)
- However, an object function might depend on a host of other things besides its parameters (like data)
- When writing software which does optimization, it may be desirable to allow the user to hold certain parameters fixed



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Maximizing a Normal Likelihood



Write a "constructor" function

```
make.NegLogLik <- function(data, fixed=c(FALSE,FALSE)) {
   params <- fixed
   function(p) {
     params[!fixed] <- p
     mu <- params[1]
     sigma <- params[2]
     a <- -0.5*length(data)*log(2*pi*sigma^2)
     b <- -0.5*sum((data-mu)^2) / (sigma^2)
     -(a + b)
   }
}</pre>
```

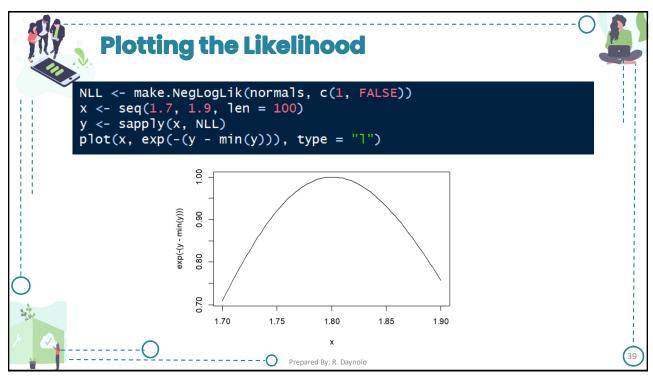
Note: Optimization functions in R minimize functions, so you need to use the negative log-likelihood.

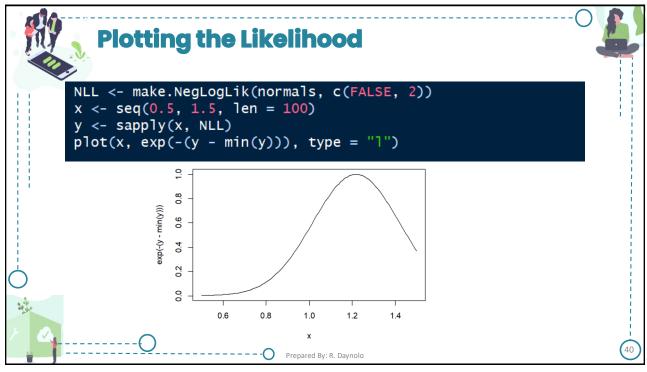


```
Maximizing a Normal Likelihood

> set.seed(1); normals <- rnorm(100, 1, 2)
> NLL <- make.NegLogLik(normals)
> NLL
function(p) {
    params[!fixed] <- p
    mu <- params[1]
    sigma <- params[2]
    a <- -0.5*length(data)*log(2*pi*sigma^2)
    b <- -0.5*sum((data-mu)^2) / (sigma^2)
    -(a + b)
}

<pre>
    bytecode: 0x0000000005ffd430>
    cenvironment: 0x0000000005flc8b8>
> ls(environment(NLL))
[1] "data" "fixed" "params"
```







Lexical Scoping Summary



- Objective functions can be "built" which contain all of the necessary data for evaluating the function
- No need to carry around long argument lists useful for interactive and exploratory work.
- · Code can be simplified and cleaned up
- Reference: Robert Gentleman and Ross Ihaka (2000). "Lexical Scope and Statistical Computing," JCGS, 9, 491–508.



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Coding Standards for R



- 1. Always use text files / text editor
- 2. Indent your code
- 3. Limit the width of your code (more or less 80 columns)
- 4. Limit of length of individual functions



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Indenting



- Indenting improves readability
- Fixing line length (80 columns) prevents lots of nesting and very long functions
- Suggested: Indents of 4 spaces at minimum; 8 spaces ideal



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Dates and Times in R



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Dates and Times in R



R has developed a special representation of dates and times

- Dates are represented by the Date class
- Times are represented by the POSIXct or the POSIXlt class
- Dates are stored internally as the number of days since 1970-01-01
- Times are stored internally as the number of seconds since 1970-01-01



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Dates are represented by the Date class and can be coerced from a character string using the as.Date() function.

```
> x <- as.Date("1970-01-01")
> x
[1] "1970-01-01"
> unclass(x)
[1] 0
> unclass(as.Date("1970-01-02"))
[1] 1
```

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Times in R



Times are represented using the POSIXCt or the POSIXIt class

- POSIXct is just a very large integer under the hood; it use a useful class when you want to store times in something like a data frame
- POSIXIt is a list underneath and it stores other useful information like the day of the week, day of the year, month, day of the month

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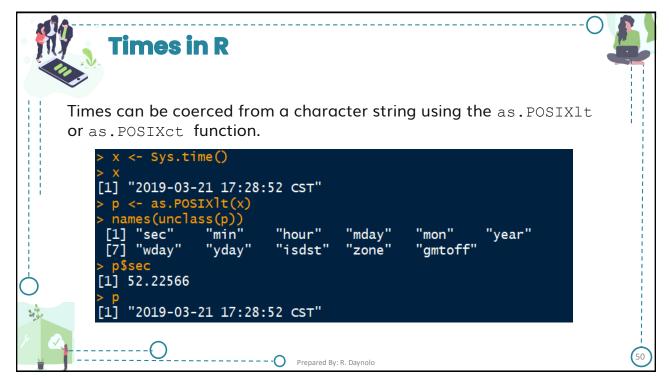
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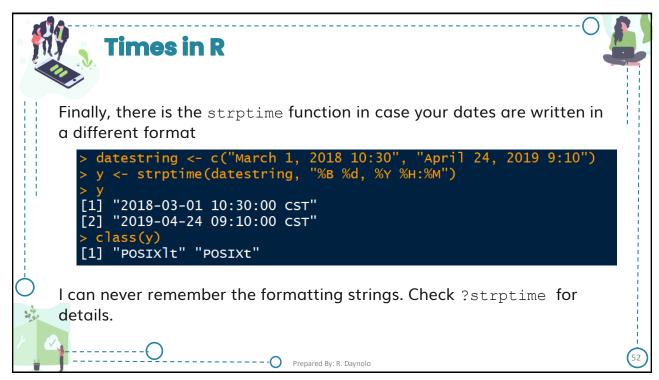
There are a number of generic functions that work on dates and times

- weekdays: give the day of the week
- months: give the month name
- quarters: give the quarter number ("Q1", "Q2", "Q3", or "Q4")

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Operations on Dates and Times



You can use mathematical operations on dates and times (well, just + and -). You can do comparisons too (==, <=)

```
> a <- as.Date("2012-01-01")
> b <- strptime("9 Jan 2011 11:34:21", "%d %b %Y %H:%M:%S")
> a-b
Error in a - b : non-numeric argument to binary operator
In addition: Warning message:
Incompatible methods ("-.Date", "-.POSIXT") for "-"
> a <- as.POSIXIt(a)
> a-b
Time difference of 356.8511 days
```



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Operations on Dates and Times



Even keeps track of leap years, leap seconds, daylight savings, and time zones

```
> q <- as.Date("2012-03-01")
> w <- as.Date("2012-02-28")
> q-w
Time difference of 2 days
> q <- as.POSIXct("2012-10-25 01:00:00")
> w <- as.POSIXct("2012-10-25 06:00:00", tz = "GMT")
> w-q
Time difference of 13 hours
```





Summary



- Dates and times have special classes in R that allow for numerical and statistical calculations
- Dates use the Date class
- Times use the POSIXct and POSIXlt class
- Character strings can be coerced to Date/Time classes using the strptime function or the
- as.Date, as.POSIXlt, or as.POSIXct



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