

# 13

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## *Non-standard evaluation*

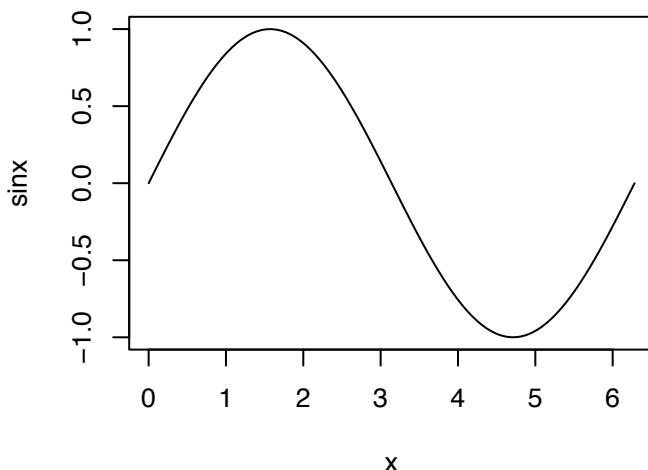
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“Flexibility in syntax, if it does not lead to ambiguity, would seem a reasonable thing to ask of an interactive programming language.”

— Kent Pitman

R has powerful tools for computing not only on values, but also on the actions that lead to those values. If you’re coming from another programming language, they are one of the most surprising features of R. Consider the following simple snippet of code that plots a sine curve:

```
x <- seq(0, 2 * pi, length = 100)
sinx <- sin(x)
plot(x, sinx, type = "l")
```



Look at the labels on the axes. How did R know that the variable on the x axis is called `x` and the variable on the y axis is called `sinx`? In most programming languages, you can only access the values of a function’s arguments. In R, you can also access the code used to compute them.

This makes it possible to evaluate code in non-standard ways: to use what is known as **non-standard evaluation**, or NSE for short. NSE is particularly useful for functions when doing interactive data analysis because it can dramatically reduce the amount of typing.

### *Outline*

- Section 13.1 teaches you how to capture unevaluated expressions using `substitute()`.
- [Section 13.2](#) shows you `subset()` works with combining `substitute()` with `eval()` to allow you to succinctly select rows from a data frame.
- [Section 13.3](#) discusses scoping issues specific to NSE, and will show you how to resolve them.
- [Section 13.4](#) shows why every function that uses NSE should have an escape hatch, a version that uses regular evaluation.
- [Section 13.5](#) teaches you how to use `substitute()` to work with functions that don't have an escape hatch.
- [Section 13.6](#) finishes off the chapter with a discussion of the downsides of NSE.

### *Prerequisites*

Before reading this chapter, make sure you're familiar with environments (Chapter 8) and lexical scoping (Section 6.2). You'll also need to install the `pryr` package with `install.packages("pryr")`. Some exercises require the `plyr` package, which you can install from CRAN with `install.packages("plyr")`.

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## 13.1 Capturing expressions

`substitute()` makes non-standard evaluation possible. It looks at a function argument and instead of seeing the value, it sees the code used to compute the value:

```
f <- function(x) {  
  substitute(x)  
}  
f(1:10)  
#> 1:10  
  
x <- 10  
f(x)  
#> x  
  
y <- 13  
f(x + y^2)  
#> x + y^2
```

For now, we won't worry about exactly what `substitute()` returns (that's the topic of Chapter 14), but we'll call it an expression.

`substitute()` works because function arguments are represented by a special type of object called a **promise**. A promise captures the expression needed to compute the value and the environment in which to compute it. You're not normally aware of promises because the first time you access a promise its code is evaluated in its environment, yielding a value.

`substitute()` is often paired with `deparse()`. That function takes the result of `substitute()`, an expression, and turns it into a character vector.

```
g <- function(x) deparse(substitute(x))  
g(1:10)  
#> [1] "1:10"  
g(x)  
#> [1] "x"  
g(x + y^2)  
#> [1] "x + y^2"
```

There are a lot of functions in Base R that use these ideas. Some use them to avoid quotes:

```
library(ggplot2)  
# the same as  
library("ggplot2")
```

Other functions, like `plot.default()`, use them to provide default labels. `data.frame()` labels variables with the expression used to compute them:

```
x <- 1:4
y <- letters[1:4]
names(data.frame(x, y))
#> [1] "x" "y"
```

We'll learn about the ideas that underlie all these examples by looking at one particularly useful application of NSE: `subset()`.

### 13.1.1 Exercises

1. One important feature of `deparse()` to be aware of when programming is that it can return multiple strings if the input is too long. For example, the following call produces a vector of length two:

```
g(a + b + c + d + e + f + g + h + i + j + k + l + m +
  n + o + p + q + r + s + t + u + v + w + x + y + z)
```

Why does this happen? Carefully read the documentation. Can you write a wrapper around `deparse()` so that it always returns a single string?

2. Why does `as.Date.default()` use `substitute()` and `deparse()`? Why does `pairwise.t.test()` use them? Read the source code.
3. `pairwise.t.test()` assumes that `deparse()` always returns a length one character vector. Can you construct an input that violates this expectation? What happens?
4. `f()`, defined above, just calls `substitute()`. Why can't we use it to define `g()`? In other words, what will the following code return? First make a prediction. Then run the code and think about the results.

```
f <- function(x) substitute(x)
g <- function(x) deparse(f(x))
g(1:10)
g(x)
g(x + y ^ 2 / z + exp(a * sin(b)))
```

---

## 13.2 Non-standard evaluation in subset

While printing out the code supplied to an argument value can be useful, we can actually do more with the unevaluated code. Take `subset()`, for example. It's a useful interactive shortcut for subsetting data frames: instead of repeating the name of data frame many times, you can save some typing:

```
sample_df <- data.frame(a = 1:5, b = 5:1, c = c(5, 3, 1, 4, 1))
```

```
subset(sample_df, a >= 4)
```

```
#>   a b c
```

```
#> 4 4 2 4
```

```
#> 5 5 1 1
```

```
# equivalent to:
```

```
# sample_df[sample_df$a >= 4, ]
```

```
subset(sample_df, b == c)
```

```
#>   a b c
```

```
#> 1 1 5 5
```

```
#> 5 5 1 1
```

```
# equivalent to:
```

```
# sample_df[sample_df$b == sample_df$c, ]
```

`subset()` is special because it implements different scoping rules: the expressions `a >= 4` or `b == c` are evaluated in the specified data frame rather than in the current or global environments. This is the essence of non-standard evaluation.

How does `subset()` work? We've already seen how to capture an argument's expression rather than its result, so we just need to figure out how to evaluate that expression in the right context. Specifically, we want `x` to be interpreted as `sample_df$x`, not `globalenv()$x`. To do this, we need `eval()`. This function takes an expression and evaluates it in the specified environment.

Before we can explore `eval()`, we need one more useful function: `quote()`. It captures an unevaluated expression like `substitute()`, but doesn't do any of the advanced transformations that can make `substitute()` confusing. `quote()` always returns its input as is:

```
quote(1:10)
#> 1:10
quote(x)
#> x
quote(x + y^2)
#> x + y^2
```

We need `quote()` to experiment with `eval()` because `eval()`'s first argument is an expression. So if you only provide one argument, it will evaluate the expression in the current environment. This makes `eval(quote(x))` exactly equivalent to `x`, regardless of what `x` is:

```
eval(quote(x <- 1))
eval(quote(x))
#> [1] 1

eval(quote(y))
#> Error in eval(expr, envir, enclos): object 'y' not found
```

`quote()` and `eval()` are opposites. In the example below, each `eval()` peels off one layer of `quote()`'s.

```
quote(2 + 2)
#> 2 + 2
eval(quote(2 + 2))
#> [1] 4

quote(quote(2 + 2))
#> quote(2 + 2)
eval(quote(quote(2 + 2)))
#> 2 + 2
eval(eval(quote(quote(2 + 2))))
#> [1] 4
```

`eval()`'s second argument specifies the environment in which the code is executed:

```
x <- 10
eval(quote(x))
#> [1] 10
```

```
e <- new.env()
e$x <- 20
eval(quote(x), e)
#> [1] 20
```

Because lists and data frames bind names to values in a similar way to environments, `eval()`'s second argument need not be limited to an environment: it can also be a list or a data frame.

```
eval(quote(x), list(x = 30))
#> [1] 30
eval(quote(x), data.frame(x = 40))
#> [1] 40
```

This gives us one part of `subset()`:

```
eval(quote(a >= 4), sample_df)
#> [1] FALSE FALSE FALSE TRUE TRUE
eval(quote(b == c), sample_df)
#> [1] TRUE FALSE FALSE FALSE TRUE
```

A common mistake when using `eval()` is to forget to quote the first argument. Compare the results below:

```
a <- 10
eval(quote(a), sample_df)
#> [1] 1 2 3 4 5
eval(a, sample_df)
#> [1] 10

eval(quote(b), sample_df)
#> [1] 5 4 3 2 1
eval(b, sample_df)
#> Error in eval(b, sample_df): object 'b' not found
```

We can use `eval()` and `substitute()` to write `subset()`. We first capture the call representing the condition, then we evaluate it in the context of the data frame and, finally, we use the result for subsetting:

```
subset2 <- function(x, condition) {
  condition_call <- substitute(condition)
```

```

  r <- eval(condition_call, x)
  x[r, ]
}
subset2(sample_df, a >= 4)
#>   a b c
#> 4 4 2 4
#> 5 5 1 1

```

### 13.2.1 Exercises

1. Predict the results of the following lines of code:

```

eval(quote(eval(quote(eval(quote(2 + 2))))))
eval(eval(quote(eval(quote(eval(quote(2 + 2)))))))
quote(eval(quote(eval(quote(eval(quote(2 + 2)))))))

```

2. `subset2()` has a bug if you use it with a single column data frame. What should the following code return? How can you modify `subset2()` so it returns the correct type of object?

```

sample_df2 <- data.frame(x = 1:10)
subset2(sample_df2, x > 8)
#> [1] 9 10

```

3. The real `subset` function (`subset.data.frame()`) removes missing values in the condition. Modify `subset2()` to do the same: drop the offending rows.
4. What happens if you use `quote()` instead of `substitute()` inside of `subset2()`?
5. The second argument in `subset()` allows you to select variables. It treats variable names as if they were positions. This allows you to do things like `subset(mtcars, , -cyl)` to drop the cylinder variable, or `subset(mtcars, , disp:drat)` to select all the variables between `disp` and `drat`. How does this work? I've made this easier to understand by extracting it out into its own function.

```

select <- function(df, vars) {
  vars <- substitute(vars)
  var_pos <- setNames(as.list(seq_along(df)), names(df))
  pos <- eval(vars, var_pos)
  df[, pos, drop = FALSE]
}

```



```

}
select(mtcars, -cyl)

```

6. What does `evalq()` do? Use it to reduce the amount of typing for the examples above that use both `eval()` and `quote()`.

---

### 13.3 Scoping issues

It certainly looks like our `subset2()` function works. But since we're working with expressions instead of values, we need to test things more extensively. For example, the following applications of `subset2()` should all return the same value because the only difference between them is the name of a variable:

```

y <- 4
x <- 4
condition <- 4
condition_call <- 4

subset2(sample_df, a == 4)
#>   a b c
#> 4 4 2 4
subset2(sample_df, a == y)
#>   a b c
#> 4 4 2 4
subset2(sample_df, a == x)
#>      a b c
#> 1      1 5 5
#> 2      2 4 3
#> 3      3 3 1
#> 4      4 2 4
#> 5      5 1 1
#> NA    NA NA NA
#> NA.1  NA NA NA
subset2(sample_df, a == condition)
#> Error in eval(expr, envir, enclos): object 'a' not found
subset2(sample_df, a == condition_call)
#> Warning in a == condition_call: longer object length is not a
#> multiple of shorter object length

```

```
#> [1] a b c
#> <0 rows> (or 0-length row.names)
```

What went wrong? You can get a hint from the variable names I've chosen: they are all names of variables defined inside `subset2()`. If `eval()` can't find the variable inside the data frame (its second argument), it looks in the environment of `subset2()`. That's obviously not what we want, so we need some way to tell `eval()` where to look if it can't find the variables in the data frame.

The key is the third argument to `eval()`: `enclos`. This allows us to specify a parent (or enclosing) environment for objects that don't have one (like lists and data frames). If the binding is not found in `env`, `eval()` will next look in `enclos`, and then in the parents of `enclos`. `enclos` is ignored if `env` is a real environment. We want to look for `x` in the environment from which `subset2()` was called. In R terminology this is called the **parent frame** and is accessed with `parent.frame()`. This is an example of dynamic scope ([http://en.wikipedia.org/wiki/Scope\\_%28programming%29#Dynamic\\_scoping](http://en.wikipedia.org/wiki/Scope_%28programming%29#Dynamic_scoping)): the values come from the location where the function was called, not where it was defined.

With this modification our function now works:

```
subset2 <- function(x, condition) {
  condition_call <- substitute(condition)
  r <- eval(condition_call, x, parent.frame())
  x[r, ]
}

x <- 4
subset2(sample_df, a == x)
#>   a b c
#> 4 4 2 4
```

Using `enclos` is just a shortcut for converting a list or data frame to an environment. We can get the same behaviour by using `list2env()`. It turns a list into an environment with an explicit parent:

```
subset2a <- function(x, condition) {
  condition_call <- substitute(condition)
  env <- list2env(x, parent = parent.frame())
  r <- eval(condition_call, env)
  x[r, ]
}
```

```

}

x <- 5
subset2a(sample_df, a == x)
#>   a b c
#> 5 5 1 1

```

### 13.3.1 Exercises

1. `plyr::arrange()` works similarly to `subset()`, but instead of selecting rows, it reorders them. How does it work? What does `substitute(order(...))` do? Create a function that does only that and experiment with it.
2. What does `transform()` do? Read the documentation. How does it work? Read the source code for `transform.data.frame()`. What does `substitute(list(...))` do?
3. `plyr::mutate()` is similar to `transform()` but it applies the transformations sequentially so that transformation can refer to columns that were just created:

```

df <- data.frame(x = 1:5)
transform(df, x2 = x * x, x3 = x2 * x)
plyr::mutate(df, x2 = x * x, x3 = x2 * x)

```

How does `mutate` work? What's the key difference between `mutate()` and `transform()`?

4. What does `with()` do? How does it work? Read the source code for `with.default()`. What does `within()` do? How does it work? Read the source code for `within.data.frame()`. Why is the code so much more complex than `with()`?

---

## 13.4 Calling from another function

Typically, computing on the language is most useful when functions are called directly by users and less useful when they are called by other functions. While `subset()` saves typing, it's actually difficult to use non-interactively. For example, imagine we want to create a function that

randomly reorders a subset of rows of data. A nice way to do that would be to compose a function that reorders with another that selects. Let's try that:

```
subset2 <- function(x, condition) {
  condition_call <- substitute(condition)
  r <- eval(condition_call, x, parent.frame())
  x[r, ]
}
```

```
scramble <- function(x) x[sample(nrow(x)), ]
```

```
subscramble <- function(x, condition) {
  scramble(subset2(x, condition))
}
```

But it doesn't work:

```
subscramble(sample_df, a >= 4)
# Error in eval(expr, envir, enclos) : object 'a' not found
traceback()
#> 5: eval(expr, envir, enclos)
#> 4: eval(condition_call, x, parent.frame()) at #3
#> 3: subset2(x, condition) at #1
#> 2: scramble(subset2(x, condition)) at #2
#> 1: subscramble(sample_df, a >= 4)
```

What's gone wrong? To figure it out, let us `debug()` `subset2()` and work through the code line-by-line:

```
debugonce(subset2)
subscramble(sample_df, a >= 4)
#> debugging in: subset2(x, condition)
#> debug at #1: {
#>   condition_call <- substitute(condition)
#>   r <- eval(condition_call, x, parent.frame())
#>   x[r, ]
#> }
n
#> debug at #2: condition_call <- substitute(condition)
n
#> debug at #3: r <- eval(condition_call, x, parent.frame())
```

```

r <- eval(condition_call, x, parent.frame())
#> Error in eval(expr, envir, enclos) : object 'a' not found
condition_call
#> condition
eval(condition_call, x)
#> Error in eval(expr, envir, enclos) : object 'a' not found
Q

```

Can you see what the problem is? `condition_call` contains the expression `condition`. So when we evaluate `condition_call` it also evaluates `condition`, which has the value `a >= 4`. However, this can't be computed because there's no object called `a` in the parent environment. But, if `a` were set in the global environment, even more confusing things can happen:

```

a <- 4
subscramble(sample_df, a == 4)
#>   a b c
#> 1 1 5 5
#> 4 4 2 4
#> 2 2 4 3
#> 5 5 1 1
#> 3 3 3 1

a <- c(1, 1, 4, 4, 4, 4)
subscramble(sample_df, a >= 4)
#>   a b c
#> 4  4 2 4
#> NA NA NA NA
#> 3  3 3 1
#> 5  5 1 1

```

This is an example of the general tension between functions that are designed for interactive use and functions that are safe to program with. A function that uses `substitute()` might reduce typing, but it can be difficult to call from another function.

As a developer, you should always provide an escape hatch: an alternative version of the function that uses standard evaluation. In this case, we could write a version of `subset2()` that takes an already quoted expression:

```

subset2_q <- function(x, condition) {
  r <- eval(condition, x, parent.frame())

```

```
x[r, ]
}
```

Here I use the suffix `_q` to indicate that it takes a quoted expression. Most users won't need this function so the name can be a little longer.

We can then rewrite both `subset2()` and `subscramble()` to use `subset2_q()`:

```
subset2 <- function(x, condition) {
  subset2_q(x, substitute(condition))
}

subscramble <- function(x, condition) {
  condition <- substitute(condition)
  scramble(subset2_q(x, condition))
}

subscramble(sample_df, a >= 3)
#>   a b c
#> 5 5 1 1
#> 4 4 2 4
#> 3 3 3 1
subscramble(sample_df, a >= 3)
#>   a b c
#> 3 3 3 1
#> 5 5 1 1
#> 4 4 2 4
```

Base R functions tend to use a different sort of escape hatch. They often have an argument that turns off NSE. For example, `require()` has `character.only = TRUE`. I don't think it's a good idea to use an argument to change the behaviour of another argument because it makes function calls harder to understand.

### 13.4.1 Exercises

1. The following R functions all use NSE. For each, describe how it uses NSE, and read the documentation to determine its escape hatch.

- `rm()`

- `library()` and `require()`
- `substitute()`
- `data()`
- `data.frame()`

2. Base functions `match.fun()`, `page()`, and `ls()` all try to automatically determine whether you want standard or non-standard evaluation. Each uses a different approach. Figure out the essence of each approach then compare and contrast.
3. Add an escape hatch to `plyr::mutate()` by splitting it into two functions. One function should capture the unevaluated inputs. The other should take a data frame and list of expressions and perform the computation.
4. What's the escape hatch for `ggplot2::aes()`? What about `plyr::()`? What do they have in common? What are the advantages and disadvantages of their differences?
5. The version of `subset2_q()` I presented is a simplification of real code. Why is the following version better?

```
subset2_q <- function(x, cond, env = parent.frame()) {
  r <- eval(cond, x, env)
  x[r, ]
}
```

Rewrite `subset2()` and `subscramble()` to use this improved version.

---

## 13.5 Substitute

Most functions that use non-standard evaluation provide an escape hatch. But what happens if you want to call a function that doesn't have one? For example, imagine you want to create a lattice graphic given the names of two variables:

```
library(lattice)
xyplot(mpg ~ disp, data = mtcars)

x <- quote(mpg)
```

```
y <- quote(displ)
xyplot(x ~ y, data = mtcars)
#> Error in tmp[sub$]: object of type 'symbol' is not subsetting
```

We might turn to `substitute()` and use it for another purpose: to modify an expression. Unfortunately `substitute()` has a feature that makes modifying calls interactively a bit of a pain. When run from the global environment, it never does substitutions: in fact, in this situation it behaves just like `quote()`:

```
a <- 1
b <- 2
substitute(a + b + z)
#> a + b + z
```

However, if you run it inside a function, `substitute()` does substitute and leaves everything else as is:

```
f <- function() {
  a <- 1
  b <- 2
  substitute(a + b + z)
}
f()
#> 1 + 2 + z
```

To make it easier to experiment with `substitute()`, `pryr` provides the `subs()` function. It works exactly the same way as `substitute()` except it has a shorter name and it works in the global environment. These two features make experimentation easier:

```
a <- 1
b <- 2
subs(a + b + z)
#> 1 + 2 + z
```

The second argument (of both `subs()` and `substitute()`) can override the use of the current environment, and provide an alternative via a list of name-value pairs. The following example uses this technique to show some variations on substituting a string, variable name, or function call:



```
subs(a + b, list(a = "y"))
#> "y" + b
subs(a + b, list(a = quote(y)))
#> y + b
subs(a + b, list(a = quote(y())))
#> y() + b
```

Remember that every action in R is a function call, so we can also replace `+` with another function:

```
subs(a + b, list("+ = quote(f)))
#> f(a, b)
subs(a + b, list("+ = quote(`*`)))
#> a * b
```

You can also make nonsense code:

```
subs(y <- y + 1, list(y = 1))
#> 1 <- 1 + 1
```

Formally, substitution takes place by examining all the names in the expression. If the name refers to:

1. an ordinary variable, it's replaced by the value of the variable.
2. a promise (a function argument), it's replaced by the expression associated with the promise.
3. ..., it's replaced by the contents of ....

Otherwise it's left as is.

We can use this to create the right call to `xyplot()`:

```
x <- quote(mpg)
y <- quote(displ)
subs(xyplot(x ~ y, data = mtcars))
#> xyplot(mpg ~ displ, data = mtcars)
```

It's even simpler inside a function, because we don't need to explicitly quote the `x` and `y` variables (rule 2 above):

```
xyplot2 <- function(x, y, data = data) {
  substitute(xyplot(x ~ y, data = data))
}
xyplot2(mpg, disp, data = mtcars)
#> xyplot(mpg ~ disp, data = mtcars)
```

If we include ... in the call to `substitute`, we can add additional arguments to the call:

```
xyplot3 <- function(x, y, ...) {
  substitute(xyplot(x ~ y, ...))
}
xyplot3(mpg, disp, data = mtcars, col = "red", aspect = "xy")
#> xyplot(mpg ~ disp, data = mtcars, col = "red", aspect = "xy")
```

To create the plot, we'd then `eval()` this call.

### 13.5.1 Adding an escape hatch to `substitute`

`substitute()` is itself a function that uses non-standard evaluation and doesn't have an escape hatch. This means we can't use `substitute()` if we already have an expression saved in a variable:

```
x <- quote(a + b)
substitute(x, list(a = 1, b = 2))
#> x
```

Although `substitute()` doesn't have a built-in escape hatch, we can use the function itself to create one:

```
substitute_q <- function(x, env) {
  call <- substitute(substitute(y, env), list(y = x))
  eval(call)
}

x <- quote(a + b)
substitute_q(x, list(a = 1, b = 2))
#> 1 + 2
```

The implementation of `substitute_q()` is short, but deep. Let's work through the example above: `substitute_q(x, list(a = 1, b = 2))`. It's

a little tricky because `substitute()` uses NSE so we can't use the usual technique of working through the parentheses inside-out.

1. First `substitute(substitute(y, env), list(y = x))` is evaluated. The expression `substitute(y, env)` is captured and `y` is replaced by the value of `x`. Because we've put `x` inside a list, it will be evaluated and the rules of `substitute` will replace `y` with its value. This yields the expression `substitute(a + b, env)`
2. Next we evaluate that expression inside the current function. `substitute()` evaluates its first argument, and looks for name value pairs in `env`. Here, it evaluates as `list(a = 1, b = 2)`. Since these are both values (not promises), the result will be `1 + 2`

A slightly more rigorous version of `substitute_q()` is provided by the `pryr` package.

### 13.5.2 Capturing unevaluated ...

Another useful technique is to capture all of the unevaluated expressions in .... Base R functions do this in many ways, but there's one technique that works well across a wide variety of situations:

```
dots <- function(...) {
  eval(substitute(alist(...)))
}
```

This uses the `alist()` function which simply captures all its arguments. This function is the same as `pryr::dots()`. `pryr` also provides `pryr::named_dots()`, which, by using deparsed expressions as default names, ensures that all arguments are named (just like `data.frame()`).

### 13.5.3 Exercises

1. Use `subs()` to convert the LHS to the RHS for each of the following pairs:
  - `a + b + c -> a * b * c`
  - `f(g(a, b), c) -> (a + b) * c`
  - `f(a < b, c, d) -> if (a < b) c else d`

2. For each of the following pairs of expressions, describe why you can't use `subs()` to convert one to the other.
  - `a + b + c -> a + b * c`
  - `f(a, b) -> f(a, b, c)`
  - `f(a, b, c) -> f(a, b)`
3. How does `pryr::named_dots()` work? Read the source.

---

## 13.6 The downsides of non-standard evaluation

The biggest downside of NSE is that functions that use it are no longer referentially transparent ([http://en.wikipedia.org/wiki/Referential\\_transparency\\_\(computer\\_science\)](http://en.wikipedia.org/wiki/Referential_transparency_(computer_science))). A function is **referentially transparent** if you can replace its arguments with their values and its behaviour doesn't change. For example, if a function, `f()`, is referentially transparent and both `x` and `y` are 10, then `f(x)`, `f(y)`, and `f(10)` will all return the same result. Referentially transparent code is easier to reason about because the names of objects don't matter, and because you can always work from the innermost parentheses outwards.

There are many important functions that by their very nature are not referentially transparent. Take the assignment operator. You can't take `a <- 1` and replace `a` by its value and get the same behaviour. This is one reason that people usually write assignments at the top-level of functions. It's hard to reason about code like this:

```
a <- 1
b <- 2
if ((b <- a + 1) > (a <- b - 1)) {
  b <- b + 2
}
```

Using NSE prevents a function from being referentially transparent. This makes the mental model needed to correctly predict the output much more complicated. So, it's only worthwhile to use NSE if there is significant gain. For example, `library()` and `require()` can be called either with or without quotes, because internally they use `deparse(substitute(x))` plus some other tricks. This means that these two lines do exactly the same thing:

```
library(ggplot2)
library("ggplot2")
```

Things start to get complicated if the variable is associated with a value. What package will this load?

```
ggplot2 <- "plyr"
library(ggplot2)
```

There are a number of other R functions that work in this way, like `ls()`, `rm()`, `data()`, `demo()`, `example()`, and `vignette()`. To me, eliminating two keystrokes is not worth the loss of referential transparency, and I don't recommend you use NSE for this purpose.

One situation where non-standard evaluation is worthwhile is `data.frame()`. If not explicitly supplied, it uses the input to automatically name the output variables:

```
x <- 10
y <- "a"
df <- data.frame(x, y)
names(df)
#> [1] "x" "y"
```

I think it's worthwhile because it eliminates a lot of redundancy in the common scenario when you're creating a data frame from existing variables. More importantly, if needed, it's easy to override this behaviour by supplying names for each variable.

Non-standard evaluation allows you to write functions that are extremely powerful. However, they are harder to understand and to program with. As well as always providing an escape hatch, carefully consider both the costs and benefits of NSE before using it in a new domain.

### 13.6.1 Exercises

1. What does the following function do? What's the escape hatch? Do you think that this is an appropriate use of NSE?

```
nl <- function(...) {
  dots <- named_dots(...)
  lapply(dots, eval, parent.frame())
}
```

2. Instead of relying on promises, you can use formulas created with `~` to explicitly capture an expression and its environment. What are the advantages and disadvantages of making quoting explicit? How does it impact referential transparency?
3. Read the standard non-standard evaluation rules found at <http://developer.r-project.org/nonstandard-eval.pdf>.