

Metaprogramming in R

Advanced Statistical Programming for Data Science, Analysis and Finance

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About the Author

Thomas Mailund is an associate professor in bioinformatics at Aarhus University, Denmark. His background is in math and computer science, but for the last decade his main focus has been on genetics and evolutionary studies, particularly comparative genomics, speciation, and gene flow between emerging species.

About the Technical Reviewer



Massimo Nardone has more than 22 years of experience in security, web/mobile development, the cloud, and IT architecture. His true IT passions are security and Android.

He has been programming and teaching how to program with Android, Perl, PHP, Java, VB, Python, C/C++, and MySQL for more than 20 years.

He holds a master of science degree in computing science from the University of Salerno, Italy.

He has worked as a project manager, software engineer, research engineer, chief security architect, information security manager, PCI/SCADA auditor,

and senior lead IT security/cloud/SCADA architect for many years.

He currently works as a chief information security officer (CISO) for Cargotec Oyj.

He was a visiting lecturer and supervisor for exercises at the Networking Laboratory of the Helsinki University of Technology (Aalto University), and he holds four international patents (PKI, SIP, SAML, and proxy areas).

Massimo has reviewed more than 40 IT books for different publishing companies, and he is the coauthor of *Pro Android Games* (Apress, 2015).

Introduction

Welcome to *Metaprogramming in R*. I am writing this book, and my books on R programming in general, to help make more advanced teaching material available beyond the typical introductory level most textbooks on R have. This book covers some of the more advanced techniques used in R programming such as fully exploiting functional programming, writing metaprograms (code for actually manipulating the language structures), and writing domain-specific languages to embed in R.

This book introduces metaprogramming. *Metaprogramming* is when you write programs that manipulate other programs; in other words, you treat code as data that you can generate, analyze, or modify. R is a very high-level language where all operations are functions, and all functions are data that you can manipulate.

There is great flexibility in how function calls and expressions are evaluated. The lazy evaluation semantics of R mean that arguments to functions are passed as unevaluated expressions, and these expressions can be modified before they are evaluated, or they can be evaluated in other environments than the context where a function is defined. This can be exploited to create small domain-specific languages and is a fundamental component in the "tidy verse" in packages such as dplyr or ggplot2 where expressions are evaluated in contexts defined by data frames.

There is some danger in modifying how the language evaluates function calls and expressions, of course. It makes it harder to reason about code. On the other hand, adding small embedded languages for dealing with everyday programming tasks adds expressiveness to the language that far outweighs the risks of programming confusion, as long as such metaprogramming is used sparingly and in well-understood (and well-documented) frameworks.

In this book, you will learn how to manipulate functions and expressions and how to evaluate expressions in nonstandard ways. Prerequisites for reading this book are familiarity with functional programming, at least familiarity with higher-order functions, that is, functions that take other functions as an input or that return functions.

CHAPTER 1



Everything you do in R involves defining functions or calling functions. You cannot do any action without evaluating some function or other. Even assigning values to variables or subscripting vectors or lists involves evaluating functions. But functions are more than just recipes for how to perform different actions; they are also data objects in themselves, and there are ways of probing and modifying them.

Manipulating Functions

If you define a simple function like the following, you can examine the components it consists of:

```
f \leftarrow function(x) x
```

There are three parts to a function: its formal parameters, its body, and the environment it is defined in. The functions formals, body, and environment give you these:

```
formals(f)
## $x
body(f)
## x
environment(f)
## <environment: R_GlobalEnv>
```

Formals

The formal parameters are given as a list where element names are the parameter names and values are default parameters.

```
g <- function(x = 1, y = 2, z = 3) x + y + z
parameters <- formals(g)
for (param in names(parameters)) {
   cat(param, "=>", parameters[[param]], "\n")
}
## x => 1
## y => 2
## z => 3
```

Strictly speaking, it is a so-called pairlist, but that is an implementation detail that has no bearing on how you treat it. You can treat it as if it is a list.

```
g <- function(x = 1, y = 2, z = 3) x + y + z
parameters <- formals(g)
for (param in names(parameters)) {
   cat(param, " => ", '"', parameters[[param]], '"', "\n", sep = "")
}
## x => "1"
## y => "2"
## z => "3"
```

For variables in this list that do not have default values, the list represents the values as the empty name. This is a special symbol that you cannot assign to, so it cannot be confused with a real value. You cannot use the missing function to check for a missing value in a formals function (that function is useful only inside a function call, and in any case there is a difference between a missing parameter and one that doesn't have a default value), but you can always check whether the value is the empty symbol.

Primitive functions (those that call into the runtime system, such as `+`) do not have formals. Only functions that are defined in R.

```
formals(`+`)
## NULL
```