R's Object Oriented Programming System: S3 Stats 102A

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Section 1

Object-Oriented Programming

Object-Oriented Programming

Object-oriented programming (OOP) is a style of programming that focuses on defining different types of objects and functions that can be applied to those object types.

OOP can be a bit challenging in R because there are four OOP systems available: S3, S4, RC, and R6.

People have different opinions about which OOP systems are important in R. I am heavily influenced by Hadley Wickham's teachings on the subject.

In this class, we will cover S3 (the most important OOP system in R) and R6.

OOP Paradigms

In R, the S3 and S4 object-oriented programming systems use generic function OOP.

The **RC** and **R6** systems use **encapsulated** OOP.

All OOP systems store information in objects. Actions or functions done with objects are called *methods*.

Encapsulated OOP

Most other programming languages (e.g. Python, C++, Java, etc.) use encapsulated OOP.

In encapsulated OOP, the methods belong to the objects or classes. Calling a specific function often looks like <code>object.method(arg1, etc.)</code>. The object encapsulates both the data (in fields) and behaviors (with methods).

You can think of a "camera" object. The camera has fields to keep information like how many pictures have been taken, how much memory is available. The camera has methods (verbs) it can perform: shoot a photo, focus, record a video.

Encapsulated OOP is available in R via R6 and RC

Generic Function OOP

In generic function OOP, the methods belong to generic functions. Objects or classes store data in fields, but do not keep the method information. Instead, the function looks at the object class and behaves differently based on the object.

You can think of the verb "shoot." This verb takes different meanings or behaviors based on the object we are talking about. If you have a camera, *shoot* means take a picture. If I give you a hockey puck, *shoot* means hitting it towards the goal. If there's a gun, *shoot* means pulling the trigger. With "the breeze," the word *shoot* means to chat idly. In generic function OOP, how the function (verb) behaves depends on the object.

A **polymorphic** function is one that behaves differently for different input types. The **S3** and **S4** systems in R use generic polymorphic functions. In base R, all object oriented behaviors are done in the **S3** system.

Example of a Polymorphic Function

Consider the summary() function, which adapts its output depending on the type of input.

```
library(ggplot2)
summary(diamonds$carat)
```

```
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 0.2000 0.4000 0.7000 0.7979 1.0400 5.0100
```

```
summary(diamonds$cut)
```

```
## Fair Good Very Good Premium Ideal
## 1610 4906 12082 13791 21551
```

The most common polymorphic functions are print(), summary(), and plot().

Classes and Methods

Central to any object-oriented system are the concepts of class and method.

- A class is a definition of an object.
 Typically a class contains several fields that are used to hold class-specific information.
- A method is an implementation (or function) for a specific class.

Intuitively, a class defines what an object is, and methods describe what that object can do.

Inheritance

Another common feature in object-oriented programming languages is the concept of **inheritance**. Classes are usually organized in a hierarchy, so if a method does not exist for a child, then its parent's method is used. We say the child **inherits** behavior from the parent.

For example, an ordered factor inherits from a regular factor, and a tibble inherits from a data frame.

Method Dispatch and Generic Functions

The process of finding the correct method given a class is called **method dispatch**.

In R, **generic functions**, or simply **generics**, are used to determine the appropriate method. The generic function determines the class of its argument(s) and uses that to select the appropriate method.

The print(), summary(), and plot() functions are actually examples of generic functions.

OOP Systems

Base R has three object-oriented systems (plus the base types):

- **S3** is R's first, simplest, and most flexible object-oriented system. S3 is the only object-oriented system used in the base and stats packages, and it is the most commonly used system in CRAN packages.
- **S4** is a formalization of S3 that has much stricter implementation for defining classes, methods, and generic functions. S4 is implemented in the base methods package.
- RC, short for reference classes, is a very different system from S3 and S4. RC implements "message-passing" OOP, so methods belong to classes, not functions.
- With library(R6), you can use the R6 system, which is similar to RC in base R. It
 implements message-passing and encapsulated OOP. Although R6 is not part of Base R,
 it is arguably simpler and more elegant than RC.

In this class, we will focus on S3 and R6

Base Types

There is one other system that is not quite object-oriented, but is important to mention:

• Base types are the internal C-level types that are the basis of the other object-oriented systems.

Base types do not form an OOP system, but they provide the building blocks for the other OOP systems.

Base Types

Underlying every R object is a C structure, or struct, that describes how that object is stored in memory. The struct includes the contents of the object, the information needed for memory management, and, most importantly for this section, a **type**. This is the **base type** of an R object.

Base types are not considered an object-oriented system because only the R core team can create new types. As a result, new base types are added very rarely.

We have already seen data structures, which cover the most common base types (the atomic vector types and lists). Base types also encompass functions, environments, and other more exotic objects likes names, calls, and promises.

You can determine an object's base type with typeof().

Base Types

typeof(sum)

Unfortunately, the names of base types are not used consistently throughout R. The type and the corresponding is function may use different names.

```
typeof(mean)

## [1] "closure"

is.function(mean)

## [1] TRUE
```

```
## [1] "builtin"
is.primitive(sum)
```

[1] TRUE

The is.object() Function

Functions that behave differently for different base types are almost always written in C. Even if you never write C code, it is important to understand base types, since everything else is built on top of them:

- S3 objects can be built on top of any base type
- S4 objects use a special base type
- RC objects are a combination of S4 and environments (another base type).

The basic difference between base and object-oriented objects is that object-oriented objects have a class attribute.

To see if an object is a pure base type (i.e., does not have a class attribute), check that is.object(x) returns FALSE.

Section 2

The S3 Object-Oriented System

The S3 Object-Oriented System

S3 is R's first and simplest object-oriented system.

Most objects that you encounter are S3 objects. Unfortunately, there is no simple way to test if an object is an S3 object in base R.

The closest you can come is is.object(x) & !isS4(x), i.e., it is an object but not an S4 object.

The is.object() Function

```
df <- data.frame(x = 1:10, y = letters[1:10])
is.object(df)
## [1] TRUE</pre>
```

[1] FALSE

isS4(df)

is.object(df\$x)

[1] FALSE

is.object(df\$y)

Generic Functions and UseMethod()

In S3, methods belong to functions, called **generic functions**, or **generics** for short. S3 methods do not belong to objects or classes. (In contrast, most OOP systems in other programming languages use **encapsulated OOP**, in which methods belong to objects/classes. S3 uses **functional OOP**, in which methods belong to functions.)

To determine if a function is an S3 generic, you can inspect its source code for a call to UseMethod(), which is the function that figures out the correct method to call, the process of **method dispatch**.

mean

```
## function (x, ...)
## UseMethod("mean")
## <bytecode: 0x0000000015a669d8>
## <environment: namespace:base>
```

Internal Generics

Some S3 generics, like [, sum(), and cbind(), do not call UseMethod() because they are implemented in C. Instead, they call the C functions DispatchGroup() or DispatchGreval().

Functions that do method dispatch in C code are called **internal generics** and are documented in ?"internal generic".

Recognizing S3 Methods

Given a class, the job of an S3 generic is to call the right S3 method. You can recognize S3 methods by their names, which look like generic.class().

For example, the Date method for the mean() generic is called mean.Date(), and the factor method for print() is called print.factor().

Recognizing S3 Methods

The generic.class() method syntax is the reason that most modern style guides discourage the use of . in function names: it makes them look like S3 methods.

For example, is t.test() the t method for objects that are of class test?

Similarly, the use of . in class names can also be confusing: is print.data.frame() the print() method for data.frame class objects, or the print.data() method for frame class objects?

To reduce confusion, most style guides prefer the underscore _ instead of the ., as in read_csv() or as_tibble().

Testing Generic Functions

The pryr package has functions is_s3_generic() and is_s3_method() for testing whether a function is a generic function or a method.

```
library(pryr)
is_s3_generic("t.test")
```

```
## [1] TRUE
```

```
methods(t.test) # there are two versions of t.test
```

```
## [1] t.test.default* t.test.formula*
## see '?methods' for accessing help and source code
```

Testing Generic Functions

t.test

```
## function (x, ...)
## UseMethod("t.test")
## <bytecode: 0x000000013dfe848>
## <environment: namespace:stats>
```

The t.test() function definition calls UseMethod(), which is a clear indicator that it is a generic.

Testing Generic Functions

```
is_s3_generic("t.data.frame")
## [1] FALSE
is s3 method("t.data.frame")
## [1] TRUE
methods(t)
## [1] t.data.frame t.default
                                   t.gtable*
                                                  t.ts*
                                                                t.vctrs sclr*
## [6] t.vctrs_vctr*
```

see '?methods' for accessing help and source code
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25 / 72

Generic Function Example

The polymorphic behavior of generic functions allows you to use the same function name on different object types.

For example, the function t() can be used on a matrix:

```
x <- matrix(1:12, nrow = 4)
t(x)</pre>
```

```
## [,1] [,2] [,3] [,4]
## [1,] 1 2 3 4
## [2,] 5 6 7 8
## [3,] 9 10 11 12
```

Generic Function Example

The t() function can also be used on a data frame:

```
df \leftarrow data.frame(a = 1:4, b = 5:8, c = 9:12)
t(df)
```

```
## [,1] [,2] [,3] [,4]
## a 1 2 3 4
## b 5 6 7 8
## c 9 10 11 12
```

The functions behave similarly, which is important for the user experience. However, the internal code for each function is different.

Internatl code of t()

t.default

```
## function (x)
## .Internal(t.default(x))
## <bytecode: 0x0000000024ccca68>
## <environment: namespace:base>
```

t.data.frame

When you apply t() to a data.frame object, it will call the function t.data.frame(), which first converts the data.frame into a matrix and then applies the next method available for the object (which will be the default method:

The methods() Function

You can see all the methods that belong to a generic with methods():

[1] t.test.default* t.test.formula*
see '?methods' for accessing help and source code

The methods() Function

You can also list all generics that have a method for a given class:

```
methods(class = "ts") # methods available for time series objects
```

```
##
   Г1] Г
                     Γ<-
                                  aggregate
                                                as.data.frame as tibble
##
    [6] chind
                                                             diffinv
                     coerce
                                  cycle
                                                diff
## [11] filter
                     initialize
                                  kernapply lines
                                                             Math
## [16] Math2
                     monthplot
                                  na.omit
                                                0ps
                                                             plot
## [21] print
                     show
                                  slotsFromS3
                                                              time
## [26] window
                     window<-
## see '?methods' for accessing help and source code
```

Section 3

Defining S3 Classes

Defining Classes and Creating Objects

S3 is a simple and ad hoc system in the sense that it has no formal definition of a class.

To make an object an instance of a class, you only need to set the **class** attribute for a base object.

You can do that during creation with structure(), or after the fact with class<-():

```
# Create and assign class in one step
x <- structure(list("apple"), class = "fruit")

# Create, then set class
y <- list("banana")
class(y) <- "fruit"</pre>
```

The above code has just defined a new class of S3 object called "fruit"

The class() and inherits() Functions

S3 objects are usually built on top of lists or atomic vectors with attributes. Functions can also be turned into S3 objects.

Other base types are either rarely seen in R or have unusual semantics that do not work well with attributes.

You can determine the class of an S3 object with the class() function. You can see if an object inherits from a specific class using inherits(x, "classname").

```
class(x)
```

```
## [1] "fruit"
```

```
inherits(x, "fruit")
```

[1] TRUE

Multiple Classes

The class of an S3 object can be a vector, which describes behavior from most to least specific.

For example, the class of the tibble object is a vector c("tbl_df", "tbl", "data.frame") indicating that tibbles inherit behavior from data frames.

```
class(diamonds)
```

```
## [1] "tbl_df" "tbl" "data.frame"
```

Ordered factors are also examples of objects with multiple classes.

```
class(diamonds$cut)
```

```
## [1] "ordered" "factor"
```

inherits() vs class() == "classname"

When checking to see if an object is of a certain class, use inherits(object, "classname") rather than 'class(object) == "classname"

```
inherits(diamonds$cut, "factor") # returns a single value
```

```
## [1] TRUE
```

```
class(diamonds$cut) == "factor" # returns a vector because class has length 2
```

[1] FALSE TRUE

Constructors

Most S3 classes provide a **constructor** function that creates new objects with the correct structure.

The constructor should follow two principles:

- Have one argument for the base object, and one for each attribute.
- Check the type of the base object and the types of each attribute.

Constructor functions usually have the same name as the class, much like the built-in ones (e.g., factor() and data.frame()).

Constructors

An example of a constructor:

```
fruit <- function(x) {
   stopifnot(is.character(x)) # checks to see if x is a character vector
   structure(list(x), class = "fruit")
}
# in use:
z <- fruit("pineapple")
z</pre>
```

```
## [[1]]
## [1] "pineapple"
##
## attr(,"class")
## [1] "fruit"
```

Changing Classes

Aside from built-in constructor functions, S3 has no checks for correctness. This means you can change the class of existing objects:

```
# Create a linear model

lm_mtcars <- lm(log(mpg) ~ log(disp), data = mtcars)

class(lm_mtcars)

## [1] "lm"

print(lm_mtcars)</pre>
```

```
##
## Call:
## lm(formula = log(mpg) ~ log(disp), data = mtcars)
##
## Coefficients:
## (Intercept) log(disp)
## 5.3810 -0.4586
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```

Changing Classes

Change the class to data.frame

```
class(lm mtcars) <- "data.frame"</pre>
print(lm mtcars) # No longer prints properly
##
   [1] coefficients residuals
                                   effects
                                            rank
                                                               fitted values
    [6] assign
                     qr
                                   df residual xlevels
                                                               call
   [11] terms
                 model
## <0 rows> (or 0-length row.names)
# But the data is still inside
lm mtcars$coefficients
```

(Intercept) log(disp) 5.3809725 -0.4585683

##

Changing Classes

The lack of built-in validation of classes has the potential to be problematic, but it rarely causes issues in practice.

While you can change the class of an object, you never should. R does not protect you from yourself.

Section 4

Creating S3 Methods and Generics

Creating S3 Generics

The job of an S3 generic is to perform method dispatch, i.e. find the specific implementation for a class.

To create a new generic, create a function that calls UseMethod().

The UseMethod() function takes two arguments: the name of the generic function, and the argument to use for method dispatch. The second argument defaults to the first argument of the function, which is usually what you want.

```
quotation <- function(x) {
   UseMethod("quotation")
}</pre>
```

There is no need to pass any of the arguments of the generic to UseMethod(). The UseMethod() function will pass arguments to the method automatically (using what Hadley Wickham refers to as "deep magic").

Creating S3 Methods

A generic is not useful without some methods.

To add a method to a generic, create a regular function with the correct (generic.class) name:

```
quotation.fruit <- function(x) {
   "Fruits are an important part of a balanced diet."
}

x <- structure(list("banana"), class = "fruit")
class(x)</pre>
```

```
## [1] "fruit"
```

quotation(x)

```
## [1] "Fruits are an important part of a balanced diet."
```

Adding Methods to Existing Generics

Adding a method to an existing generic works in the same way:

```
mean.fruit <- function(x) {
   5
}</pre>
```

```
mean(x)
```

```
## [1] 5
```

As we saw previously with changing classes, there is no check to make sure that the method returns the class compatible with the generic. It is up to you to make sure that your method does not violate the expectations of existing code.

Method Dispatch

How does method dispatch work?

The UseMethod() function creates a vector of function names, like paste0("generic", ".", c(class(x), "default")) and looks for each potential method in turn.

The "default" class makes it possible to set up a fallback method for otherwise unknown classes.

The "default" class is a special **pseudo-class** in that it is not a real class, but we can define methods for it to use for inputs with unknown classes.

Method Dispatch Example

Consider the following generic and methods:

```
quotation <- function(x) {
  UseMethod("quotation")
quotation.fruit <- function(x) {</pre>
  "Fruits are an important part of a balanced diet."
quotation.apple <- function(x) {
  "An apple a day keeps the doctor away."
quotation.default <- function(x) {</pre>
  "The default quotation: Let food be thy medicine and medicine be thy food."
```

Method Dispatch Example

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```
# Dispatches method for apple class
a <- structure(list("Fuji"), class = c("apple", "fruit"))</pre>
quotation(a)
## [1] "An apple a day keeps the doctor away."
# No method for banana class, so uses method for fruit class
b <- structure(list("Chiquita"), class = c("banana", "fruit"))
quotation(b)
## [1] "Fruits are an important part of a balanced diet."
# No method for donut class, so falls back to default
c <- structure(list("Dunkin"), class = "donut")</pre>
quotation(c)
```

[1] "The default quotation: Let food be thy medicine and medicine be thy food."

47 / 72

Methods Are Functions

Because methods are normal R functions, they can also be called directly:

```
b <- structure(list("Chiquita"), class = c("banana", "fruit"))
# Call the correct method with dispatch:
quotation(b)</pre>
```

```
## [1] "Fruits are an important part of a balanced diet."
```

```
# Force R to call the wrong method:
quotation.apple(b)
```

```
## [1] "An apple a day keeps the doctor away."
```

However, this is just as dangerous as changing the class of an object, so you should not do it.

Method Dispatch

If there is no default method and you attempt to use a function on an object for which no method exists. R will throw an error.

```
rm(quotation.default) # we remove the default method
c <- structure(list("Dunkin"), class = "donut")
quotation(c) # we call the function on an object with class donut</pre>
```

```
## Error in UseMethod("quotation"): no applicable method for 'quotation' appl
```

Section 5

Inheritance in S3 Classes

Inheritance in S3 Classes

S3 classes can share behavior through **inheritance**.

Inheritance in S3 follows three principles:

- The class of an object can be a character vector.
- If a method is not found for the class in the first element of the vector, R looks for a method for the second class (and so on).
- A method can delegate work by calling NextMethod().

We will expand on these principles in this section.

Method Dispatch Hierarchy

The s3_dispatch() function in the sloop package inputs a function call and outputs the list of all possible function names that are considered for method dispatch.

- The lack of a symbol means the method does not exist.
- The => arrow means the method exists and is found by UseMethod().
- The -> arrow means the method exists and is used through NextMethod().
- The * means the method exists but is not used.

Method Dispatch Hierarchy

For example, consider calling the generic print() function on an ordered factor:

```
library(sloop)
s3_dispatch(print(factor(letters, ordered = TRUE)))
```

```
## print.ordered
## => print.factor
## * print.default
```

There is no print() method for the ordered class, so the print() generic dispatches the print.factor() method for ordered objects.

Subclasses and Superclasses

The ordered class is said to be a **subclass** of factor because it always appears before it in the class vector.

Conversely, the factor class is a superclass of ordered.

S3 has no formal restrictions on the relationship between sub- and superclasses, but there are two principles to follow when creating a subclass:

- The base type of the subclass should be that same as the superclass.
- The attributes of the subclass should be a superset of the attributes of the superclass.

Section 6

Method Dispatch Self-Quiz

Method Dispatch Self-Quiz

The following slides can be seen as a self-quiz for the topic of method dispatch.

The functions and methods are very simple, usually adding a constant like 2 or 10. Becasue they are so simple, I've opted to leave them as a single line.

With each slide, try to predict the output. The full details of what methods R searches are provided with the s3_dispatch() function.

As far as studying, you need to learn which method will be dispatched. You do not need to learn the exact output of the s3_dispatch() function itself.

Method Dispatch Self Quiz

```
rm(list = ls())
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10 # the f method for class k
k <- 1</pre>
```

Method Dispatch Self Quiz

```
rm(list = ls())
f  <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10 # the f method for class k
k <- 1</pre>
```

f(k)

Method Dispatch Self Quiz

```
rm(list = ls())
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10 # the f method for class k
k <- 1</pre>
```

```
## Error in UseMethod("f"): no applicable method for 'f' applied to an object of class "c('dou
```

```
s3_dispatch(f(k)) # full details of the result
```

```
## f.double
## f.numeric
## f.default
```

f(k)

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
k <- 1</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
k <- 1</pre>
```

```
class(k) <- "j" # object k has a class of "j"
f(k)</pre>
```

=> f.j ## f.default

```
f <- function(x) UseMethod("f") # the generic f function</pre>
f.j \leftarrow function(x) x + 2 # the f method for class j
f.k \leftarrow function(x) x + 10
k <- 1
class(k) <- "j" # object k has a class of "j"</pre>
f(k)
## [1] 3
## attr(,"class")
## [1] "j"
s3_dispatch(f(k))
```

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```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
k <- 1</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
k <- 1</pre>
```

```
class(k) <- "k" # object k has a class of "k"
f(k)</pre>
```

=> f.k ## f.default

```
f <- function(x) UseMethod("f") # the generic f function</pre>
f.j \leftarrow function(x) x + 2 # the f method for class j
f.k \leftarrow function(x) x + 10
k <- 1
class(k) <- "k" # object k has a class of "k"</pre>
f(k)
## [1] 11
## attr(,"class")
## [1] "k"
s3_dispatch(f(k))
```

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```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
k <- 1</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
k <- 1

class(k) <- "k"
f.default <- function(x) x + 100
f(k)</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function</pre>
f.j \leftarrow function(x) x + 2 # the f method for class j
f.k \leftarrow function(x) x + 10
k <- 1
class(k) \leftarrow "k"
f.default \leftarrow function(x) x + 100
f(k)
## [1] 11
## attr(,"class")
## [1] "k"
```

=> f.k

* f.default
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s3_dispatch(f(k)) # full details of the result

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
k <- 1</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
k <- 1

class(k) <- c("a", "b")
f(k)</pre>
```

##

f.b

```
f <- function(x) UseMethod("f") # the generic f function</pre>
f.j \leftarrow function(x) x + 2 # the f method for class j
f.k \leftarrow function(x) x + 10
f.default \leftarrow function(x) x + 100
k <- 1
class(k) <- c("a", "b")
f(k)
## [1] 101
## attr(,"class")
## [1] "a" "b"
s3_dispatch(f(k))
## f.a
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
k <- 1</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
k <- 1</pre>
```

```
class(k) <- NULL
f(k)</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
k <- 1

class(k) <- NULL
f(k)</pre>
```

```
s3_dispatch(f(k))
```

[1] 101

```
## f.double
## f.numeric
## => f.default
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 50</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50

l <- structure(10, class = c("k", "l"))
f(1)</pre>
```

* f.1

```
f <- function(x) UseMethod("f") # the generic f function</pre>
f.j \leftarrow function(x) x + 2 # the f method for class j
f.k \leftarrow function(x) x + 10
f.default <- function(x) x + 100
f.1 \leftarrow function(x) x + 50
1 <- structure(10, class = c("k", "l"))</pre>
f(1)
## [1] 20
## attr(,"class")
## [1] "k" "l"
s3_dispatch(f(1))
## => f.k
```

* f.default
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```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50
l <- 10</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50
l <- 10

class(l) <- c("m","l")
f(l)</pre>
```

f.m

=> f.1

```
f <- function(x) UseMethod("f") # the generic f function</pre>
f.j \leftarrow function(x) x + 2 # the f method for class j
f.k \leftarrow function(x) x + 10
f.default <- function(x) x + 100
f.1 \leftarrow function(x) x + 50
1 <- 10
class(1) <- c("m","1")
f(1)
## [1] 60
## attr(,"class")
## [1] "m" "l"
s3_dispatch(f(1))
```

* f.default Copyright Miles Chen. For personal use only. Do not distribute.

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 50
1 <- 10</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50
l <- 10

class(l) <- c("m", "n")
f(l)</pre>
```

f.m

f n

##

```
f <- function(x) UseMethod("f") # the generic f function</pre>
f.j \leftarrow function(x) x + 2 # the f method for class j
f.k \leftarrow function(x) x + 10
f.default <- function(x) x + 100
f.1 \leftarrow function(x) x + 50
1 <- 10
class(1) <- c("m", "n")
f(1)
## [1] 110
## attr(,"class")
## [1] "m" "n"
s3_dispatch(f(1))
```

=> f.default Copyright Miles Chen. For personal use only. Do not distribute.

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50
l <- 10</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50
l <- 10</pre>
```

```
class(1) <- c("m","n")
f.j(1)</pre>
```

class(1) <- c("m", "n")

[1] "m" "n"

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50
l <- 10</pre>
```

```
f.j(1)
## [1] 12
## attr(,"class")
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50</pre>
```

f(7)

##

##

f.double

=> f.default

```
f <- function(x) UseMethod("f") # the generic f function</pre>
f.j \leftarrow function(x) x + 2  # the f method for class j
f.k \leftarrow function(x) x + 10
f.default \leftarrow function(x) x + 100
f.1 \leftarrow function(x) x + 50
f(7)
## [1] 107
s3_dispatch(f(7))
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50</pre>
```

```
f.j(7)
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50</pre>
```

```
f.j(7)
```

```
## [1] 9
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 50
f.integer <- function(x) 100 * x</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50
f.integer <- function(x) 100 * x</pre>
```

f(7)

```
f <- function(x) UseMethod("f") # the generic f function</pre>
f.j \leftarrow function(x) x + 2 # the f method for class j
f.k \leftarrow function(x) x + 10
f.default \leftarrow function(x) x + 100
f.1 \leftarrow function(x) x + 50
f.integer \leftarrow function(x) 100 * x
f(7)
## [1] 107
s3_dispatch(f(7))
```

```
## => f.default
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```

f.double

f numeric

##

##

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 50
f.integer <- function(x) 100 * x</pre>
```

```
f <- function(x) UseMethod("f") # the generic f function
f.j <- function(x) x + 2 # the f method for class j
f.k <- function(x) x + 10
f.default <- function(x) x + 100
f.l <- function(x) x + 50
f.integer <- function(x) 100 * x</pre>
```

```
f(7L)
```

```
f <- function(x) UseMethod("f") # the generic f function</pre>
f.j \leftarrow function(x) x + 2  # the f method for class j
f.k \leftarrow function(x) x + 10
f.default \leftarrow function(x) x + 100
f.1 \leftarrow function(x) x + 50
f.integer \leftarrow function(x) 100 * x
f(7L)
## [1] 700
s3_dispatch(f(7L))
```

```
## f.numeric
## * f.default
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```

=> f.integer

```
# Note the name in UseMethod is "g" instead of "f". You should never do this.
f <- function(x) UseMethod("g")
f.j <- function(x) x + 2
g.j <- function(x) -1 * (x + 2)
f.default <- function(x) x + 100
g.default <- function(x) -1 * (x + 100)</pre>
```

```
# Note the name in UseMethod is "g" instead of "f". You should never do this.
f <- function(x) UseMethod("g")
f.j <- function(x) x + 2
g.j <- function(x) -1 * (x + 2)
f.default <- function(x) x + 100
g.default <- function(x) -1 * (x + 100)</pre>
```

```
k <- structure(10, class = "j")
f(k)</pre>
```

```
# Note the name in UseMethod is "g" instead of "f". You should never do this.
f <- function(x) UseMethod("g")
f.j <- function(x) x + 2
g.j <- function(x) -1 * (x + 2)
f.default <- function(x) x + 100
g.default <- function(x) -1 * (x + 100)</pre>
```

```
## [1] -12
## attr(,"class")
## [1] "j"
```

f(k)

k <- structure(10, class = "j")</pre>

```
# Note the name in UseMethod is "g" instead of "f". You should never do this.
f <- function(x) UseMethod("g")
f.j <- function(x) x + 2
g.j <- function(x) -1 * (x + 2)
f.default <- function(x) x + 100
g.default <- function(x) -1 * (x + 100)</pre>
```

```
# Note the name in UseMethod is "g" instead of "f". You should never do this.
f <- function(x) UseMethod("g")
f.j <- function(x) x + 2
g.j <- function(x) -1 * (x + 2)
f.default <- function(x) x + 100
g.default <- function(x) -1 * (x + 100)</pre>
```

```
m <- 10 f(m)
```

```
# Note the name in UseMethod is "g" instead of "f". You should never do this.
f <- function(x) UseMethod("g")
f.j <- function(x) x + 2
g.j <- function(x) -1 * (x + 2)
f.default <- function(x) x + 100
g.default <- function(x) -1 * (x + 100)</pre>
```

```
## [1] -110
```

m <- 10 f(m)