# **Object-Oriented Programming**

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Motivation

- S3 Classes
- S4 Classes
- Reference Classes





Support structural thinking (numeric vectors, factors, data frames, results of fitting a model, ...)





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- Support functional thinking ("generic" functions)
- Allow for simple extensibility (printing, plotting, summarizing, ...)





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### **Basic idea**



Specify the class via the class attribute







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- The "method" of generic foo for class bar is called foo.bar
- Allow for default methods called foo.default





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#### Classes

- Generic and Method Functions
- Group Methods
- Odds and ends
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### Reference Classes



 Function class() gets the class attribute, and there is a replacement function for setting it:

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[1] 1 2 3
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 R> class(x) <- "bar"</li>
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```
[1] 1 2 3
attr(,"class")
[1] "bar"
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 If e.g. class(x) is c("bar", "baz", "grr"), then x belongs to class bar and inherits from baz, then grr, etc.





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- If e.g. class(x) is c("bar", "baz", "grr"), then x belongs to class bar and inherits from baz, then grr, etc.
- Function unclass() removes the class(es)





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- First, using the *GENERIC*.*CLASS* naming convention for each class the primary arg inherits from
- If no such method found, try GENERIC.default (default method)
- If not found either: error.





Typically via UseMethod(). E.g.,

```
R> foo <- function(x, ...) UseMethod("foo")
R> foo
```

function(x, ...) UseMethod("foo")





Simply create functions obeying the *GENERIC*.*CLASS* naming convention. E.g.,

```
R> foo.bar <- function(x, ...) cat("I am here\n") R> foo.bar
```

```
function(x, ...) cat("I am here\n")
```

Basic dispatch by calling the generic:

R > foo(x)

I am here





Function NextMethod() can be used within methods to call the next method. E.g.,

```
R> test <- function(x) UseMethod("test")
R> test.c1 <- function(x) { cat("c1\n"); NextMethod(); x }
R> test.c2 <- function(x) { cat("c2\n"); NextMethod(); x }
R> test.c3 <- function(x) { cat("c3\n"); NextMethod(); x }
R> x <- 1
R> class(x) <- c("c1", "c2")</pre>
```

What will happen?

R > test(x)





Specify argument for dispatch in the UseMethod() call. E.g.,

foo <- function(x, y, ...) UseMethod("foo", y)</pre>





No "formal" testing (predicate) for and coercion to S3 classes. By convention, is.bar() and as.bar(). E.g.,

```
R> is.bar <- function(x) inherits(x, "bar")
R> as.bar <- function(x) if(is.bar(x)) x else bar(x)</pre>
```

(assuming that bar() creates objects of class "bar").





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Issues:

- How can we e.g. add or compare objects of a certain class?
- Or more generally, e.g. compare with objects not in the same class?

Need a dispatch mechanism for "operators" (such as "<") which works in *both* arguments!



# **Group methods**



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Operators grouped in three categories (Math, Ops, Summary)





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Mechanism:

- Operators grouped in three categories (Math, Ops, Summary)
- Invoked if the operands correspond to the same method, or one to a method that takes precedence; otherwise, the default method is used.
- Class methods dominate group methods.
- Dispatch info available: .Method, .Generic, .Group, .Class





Math	abs sign sqrt floor ceiling trunc round signif
	exp log cos sin tan acos asin atan
	cosh sinh tanh acosh asinh atanh
	lgamma gamma gammaCody digamma
	trigamma tetragamma pentagamma
	cumsum cumprod cummax cummin
Ops	+ - * / ^ %% %/%
	&   !
	== != < <= >= >
Summary	all any sum prod min max range





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In addition to UseMethod dispatch and group methods, some functions dispatch "internally" (Dispatch0rEval() in the underlying C code of builtin functions):

- subscripting and subassigning ([, [[, \$)
- length dim dimnames c unlist
- as.character as.vector
- many is.xxx functions for builtins data types xxx





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- No formal class structure (can have objects with class c("foo", "U") and c("foo", "V"), no structural integrity, ...)




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- No formal class structure (can have objects with class c("foo", "U") and c("foo", "V"), no structural integrity, ...)
- No flexible dispatch on several arguments

(Some of these weaknesses disappear for code in namespaces.)



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### **Basics**



Every object has exactly one class





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- Every object has exactly one class
- All objects in a class must have the same structure
- All methods for a new-style generic must have exactly the same formal arguments (could be ...)
- Can dispatch according to the signature of arbitrarily many arguments



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Classes are created by setClass():





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- First arg is the name of the class
- Arg representation specifies the slots

```
R> setClass("fungi",
+ representation(x = "numeric", y = "numeric",
+ species = "character"))
```





Creation can use existing classes:

```
R> setClass("xyloc",
+ representation(x = "numeric",
+ y = "numeric"))
R> setClass("fungi",
+ representation("xyloc",
+ species = "character"))
```





Existing classes can be examined using getClass():

```
R> getClass("fungi")
```

```
Class "fungi" [in ".GlobalEnv"]
```

Slots:

Name: species x y Class: character numeric numeric

Extends: "xyloc"





New instances can be created using new()

```
R> f1 <- new("fungi", x = runif(5), y = runif(5),
+ species = sample(letters[1:3], 5, replace = TRUE))
```





Such instances can be inspected by typing their name (which calls show() instead of print())

R> f1

```
An object of class "fungi"
Slot "species":
[1] "c" "a" "c" "b" "a"
```

Slot "x": [1] 0.53827758 0.72234531 0.04228939 0.46204104 0.20706792

Slot "y": [1] 0.69083106 0.07209983 0.51760378 0.12305639 0.45677010





The slots can be accessed using @:

R> f1@x

 $[1] \ 0.53827758 \ 0.72234531 \ 0.04228939 \ 0.46204104 \ 0.20706792 \\$ 





Typically, there would be a *creator* function:

```
R> fungi <- function(x, y, species)
+ new("fungi", x = x, y = y, species = species)</pre>
```

so that users do not need to call new() themselves.





Using the creator:

```
R> f2 <- fungi(runif(3), runif(3), letters[1 : 3])
R> f2
```

```
An object of class "fungi"
Slot "species":
[1] "a" "b" "c"
```

```
Slot "x":
[1] 0.3760097 0.0144067 0.6897521
```

```
Slot "y":
[1] 0.6598063 0.6988959 0.7950575
```





Can specify restrictions for "valid" objects via the validity argument to setClass or via setValidity (note: sets to the value of the current version.)

```
R> ## A valid "xyloc" object has the same number
R> ## of x and y values:
R> .valid_xyloc_object <- function(object) {</pre>
      nx <- length(object@x)</pre>
+
      ny <- length(object@y)</pre>
+
      if(nx == ny)
+
           TRUE
+
      else
+
           sprintf("Unequal x, y lengths: %d, %d", nx, ny)
+
+
  }
```





```
R> setValidity("xyloc", .valid_xyloc_object)
```

```
Class "xyloc" [in ".GlobalEnv"]
```

Slots:

```
Name: x y
Class: numeric numeric
```

```
Known Subclasses: "fungi"
```

[1] "invalid class \"xyloc\" object: Unequal x, y lengths: 5, 3"





```
R> ## We can still do bad things "directly":
R> xyl@y <- 3
R> ## However, now validObject() will throw an error ...
R> tc <- tryCatch(validObject(xy1), error = identity)
R> tc
```

<simpleError in validObject(xy1): invalid class "xyloc" object: Unequal</pre>

```
R> conditionMessage(tc)
```

[1] "invalid class \"xyloc\" object: Unequal x, y lengths: 5, 1"







Classes have prototypes stored with their definition







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- Default prototype: for each slot a new object of that class
- Other prototypes might be more appropriate; can be specified via prototype arg to setClass()





Usually have neither representation nor prototype





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- Useful by inheritance between classes





- Usually have neither representation nor prototype
- Useful by inheritance between classes
- Can be created using "VIRTUAL" when specifying the representation



### Inheritance



# Class A *inherits from* (or *extends*) class B if is(x, "B") is true whenever is(x, "A") is.







```
Class A inherits from (or extends) class B if is(x, "B") is true whenever is(x, "A") is.
```

Can be determined using function extends():

```
R> extends("fungi", "xyloc")
```

[1] TRUE





Use general-purpose functions is() and as() for testing and coercion:
R> is(f1, "xyloc")
[1] TRUE
R> as(c(1, 2, 3, 4.4), "integer")
[1] 1 2 3 4





Use general-purpose functions is() and as() for testing and coercion:
 R> is(f1, "xyloc")
 [1] TRUE

```
R> as(c(1, 2, 3, 4.4), "integer")
```

```
[1] 1 2 3 4
```

 Corresponding information obtained from the class definition, or via setIs() and setAs()



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- Explicit setting of methods for certain *signatures* (the classes of the arguments used in the dispatch)
- Can use several arguments in the dispatch
- Can determine the method dispatched to with selectMethod()





Typically by setting a method (which automatically makes a function generic with the old definition as the default method).

Or explicitly using setGeneric().





```
By using setMethod():
```

```
R> setMethod("show", "fungi",
+ function(object) cat("I am just a fungus.\n"))
R> f1
```

```
I am just a fungus.
```

(Note: sets to the value of the current version if a named function is used.)

For more complicated signatures:

```
R> setMethod("plot", c("numeric", "factor"), .....)
```


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- Partially different from the Green Book (e.g., currently no getGroupMembers())
- Mechanism "similar" to S3 group methods
- Can construct arbitrary groups using setGroupGeneric()





Math	log, sqrt, log10, cumprod, abs, acos, acosh, asin, asinh, atan, atanh, ceiling, cos, cosh, cumsum, exp, floor, gamma, lgamma, sin, sinh, tan, tanh, trunc
Math2	round signif
Ops	Arith Compare [Logic]
Arith	+ - * / ^ %% %/%
Compare	== != < <= >= >
Summary	max, min, range, prod, sum, any, all
Complex	Arg, Conj, Im, Mod, Re



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Main differences between reference classes and S3 and S4:

- Refclass objects use message-passing OO
- Refclass objects are mutable: the usual R copy on modify semantics do not apply

Thus, the refclass object system behaves more like Java or C#.

Use them only where mutable state is absolutely required!





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A reference class has three main components, given by three arguments to setRefClass:

- contains, the classes which the class inherits from.
- fields are the equivalent of slots in S4. They can be specified as a vector of field names, or a named list of field types.
- methods are functions that operate within the context of the object and can modify its fields.





A simple reference class for bank accounts with methods for withdrawal and deposit:

```
R> setRefClass("Account",
               fields = list(balance = "numeric"),
+
              methods =
+
               list(deposit =
+
                    function(amount) {
+
                        ## The following string documents the method
+
                        "Deposit the given amount from the account"
+
                        balance <<- balance + amount
+
+
                    }.
                    withdraw =
+
                    function(amount) {
+
                        "Withdraw the given amount from the account"
+
                        balance <<- balance - amount
+
                    }))
+
```







What is returned by setRefClass, or can be retrieved by getRefClass, is a *reference class generator*:

```
R> (Account <- getRefClass("Account"))</pre>
```

```
Generator for class "Account":
```

Class fields:

```
Name: balance
Class: numeric
```

```
Class Methods:
    "withdraw", "deposit", "field", "trace", "getRefClass",
    "initFields", "copy", "callSuper", ".objectPackage", "export",
    "untrace", "getClass", "show", "usingMethods", ".objectParent",
    "import"
```

#### Reference Superclasses: "envRefClass"





 new() for creating new objects of that class. Takes named arguments specifying initial values for the fields.





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- new() for creating new objects of that class. Takes named arguments specifying initial values for the fields.
- methods() for modifying existing or adding new methods
- help() for getting help about methods
- fields() to get a list of fields defined for class
- lock() locks the named fields so that their value can only be set once





We can use the generator to create an account object:

```
R> a1 <- Account$new(balance = 100)
R> a1
```

```
Reference class object of class "Account"
Field "balance":
[1] 100
```

We can ask the generator for help:

```
R> Account$help("withdraw")
```

```
Call:
$withdraw(amount)
```

#### Withdraw the given amount from the account





We can then make deposits and withdrawals:

```
R> al$withdraw(50)
R> al
```

```
Reference class object of class "Account"
Field "balance":
[1] 50
```

To access the balance:

R> a1\$balance

[1] 50

R> a1\$field("balance")

[1] 50 Slide 54





The most important property of refclass objects is that they are mutable, or equivalently they have reference semantics:

```
R> a2 <- a1
R> a2$deposit(10)
R> a1
Reference class object of class "Account"
Field "balance":
[1] 60
```

(In the example, copies will always refer to the same account.)



## **Cool stuff**



```
R> setRefClass("Dist")
R> setRefClass("DistUniform",
              c("a", "b"),
+
               "Dist",
+
              methods =
+
               list(mean =
+
                    function() {
+
                        (a + b) / 2
+
                    }))
+
R> DistUniform <- getRefClass("DistUniform")</pre>
R > U01 <- DistUniform (a = 0, b = 1)
R > U01 ()
```

[1] 0.5

(Could add other sub-classes for observed or simulated values.)

