



The Fundamentals of S3 Objects and Methods

Kenny Flagg
useR Bozeman
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<https://github.com/kflagg/useRS3>

- What is object-oriented (OO) programming?
- Some familiar classes of objects
- Writing methods to work with objects
- Case study: extending the `htest` class

What is object-oriented (OO) programming?

- System of programming that models relationships among abstract structures.
- Simplifies development and maintenance of complicated programs. (R Core Team 2017)
- Concepts:
 - *classes* define standard structures to store information.
 - *objects* are realized instances of classes.
 - *methods* are functions that act on objects according to their classes.
- R has several different systems of defining objects.
 - We will work with the S3 system, the least formal and by far most common.

What is object-oriented (OO) programming?

I'm not developing packages. Why should I learn this programming mumbo-jumbo?

- Understanding others' code/debugging.
- Recycling your own code.

Some familiar classes of objects

- You have probably used objects of these classes:
 - `data.frame`
 - `lm`
 - `glm`
 - `htest`
- Let's use the `str()` function to investigate!

```
str(iris)
```

```
versicolor <- glm(I(Species == 'versicolor') ~  
                  Sepal.Length + Sepal.Width + Petal.Length + Petal.Width,  
                  family = binomial, data = iris)  
str(lm(Petal.Length ~ Species, data = iris))
```

```
petal <- t.test(iris$Petal.Length, mu = 4)  
str(petal)
```

Some familiar classes of objects

```
str(iris)
```

```
# 'data.frame': 150 obs. of  5 variables:
#  $ Sepal.Length: num  5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
#  $ Sepal.Width : num  3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
#  $ Petal.Length: num  1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
#  $ Petal.Width : num  0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
#  $ Species      : Factor w/ 3 levels "setosa","versicolor",...: 1 1 1 1 1 1 1 1 1 1
```

Some familiar classes of objects

```
versicolor <- glm(I(Species == 'versicolor') ~
  Sepal.Length + Sepal.Width + Petal.Length + Petal.Width,
  family = binomial, data = iris)
str(versicolor)

# List of 30
# $ coefficients      : Named num [1:5] 7.378 -0.245 -2.797 1.314 -2.778
# ..- attr(*, "names")= chr [1:5] "(Intercept)" "Sepal.Length" "Sepal.Wid
# $ residuals        :Class 'AsIs'   Named num [1:150] -1.09 -1.39 -1.21 -1
# ..- attr(*, "names")= chr [1:150] "1" "2" "3" "4" ...
# $ fitted.values     : Named num [1:150] 0.0849 0.2829 0.172 0.268 0.0671
# ..- attr(*, "names")= chr [1:150] "1" "2" "3" "4" ...
# $ effects           :Class 'AsIs'   Named num [1:150] 2.383 0.739 4.822 -0
# ..- attr(*, "names")= chr [1:150] "(Intercept)" "Sepal.Length" "Sepa
...
# - attr(*, "class")= chr [1:2] "glm" "lm"
```

Some familiar classes of objects

```
petal <- t.test(iris$Petal.Length, mu = 4)
str(petal)
```

```
# List of 9
# $ statistic : Named num -1.68
# ..- attr(*, "names")= chr "t"
# $ parameter : Named num 149
# ..- attr(*, "names")= chr "df"
# $ p.value : num 0.0953
# $ conf.int : atomic [1:2] 3.47 4.04
# ..- attr(*, "conf.level")= num 0.95
# $ estimate : Named num 3.76
# ..- attr(*, "names")= chr "mean of x"
# $ null.value : Named num 4
# ..- attr(*, "names")= chr "mean"
# $ alternative: chr "two.sided"
# $ method : chr "One Sample t-test"
# $ data.name : chr "iris$Petal.Length"
# - attr(*, "class")= chr "htest"
```


Some familiar classes of objects

- Many objects are actually lists with a `class` attribute.
- What happens when you change a data frame's class to "list"?

```
dframe <- data.frame(x = 1:4, y = 5:8, row.names = letters[1:4])  
dframe  
class(dframe) <- 'list'  
dframe
```

Some familiar classes of objects

```
dframe <- data.frame(x = 1:4, y = 5:8, row.names = letters[1:4])  
dframe
```

```
#   x y  
# a 1 5  
# b 2 6  
# c 3 7  
# d 4 8
```

```
class(dframe) <- 'list'  
dframe
```

```
# $x  
# [1] 1 2 3 4  
#  
# $y  
# [1] 5 6 7 8  
#  
# attr("row.names")  
# [1] "a" "b" "c" "d"
```

Writing methods to work with objects

- Objects are nouns.
- Methods are verbs.
 - They tell the computer what to do to objects.
- S3 methods are functions with different versions defined for each class.
 - `print`, `summary`, `plot` all behave according to the class of the object.

Writing methods to work with objects

- You can see what methods are available for a class.

```
methods(class = 'lm')
```

- Are there things you didn't know you could do with `lm` objects?

Writing methods to work with objects

- What do methods look like? (Recall you can usually type the name of a function to see its source code.)

```
summary  
summary.data.frame  
summary.lm  
summary.default
```

- Some source code is hidden, but you can often find it with `getS3method`.

```
plot.lm  
getS3method('plot', 'lm')
```

Writing methods to work with objects

- Generic methods are defined with UseMethod.

```
method <- function(x, ...) UseMethod('method')
```

- Methods for specific classes are named method.class.
- If a method isn't defined for a class, R uses method.default.

Case study

Let's extend the `htest` class to handle a one-sample simulation test!
Then we'll write a method to plot a histogram of the sampling distribution.

```
result <- sim.test(iris$Petal.Length, mu = 4)

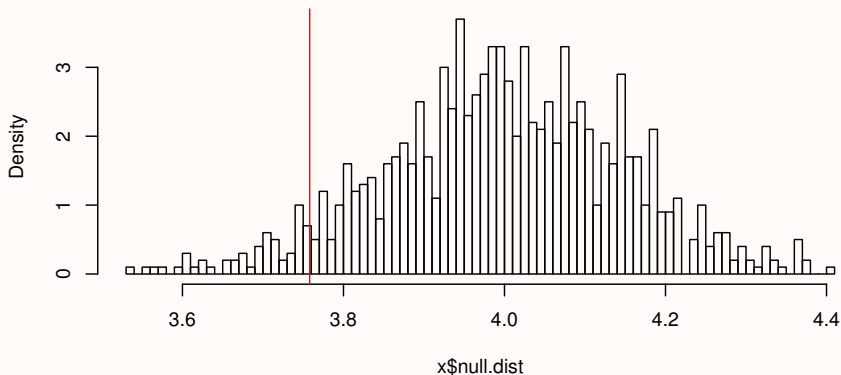
# This uses the print.htest function.
result

#
# One Sample Simulation Test
#
# data:  iris$Petal.Length
# number of simulations = 1000, p-value = 0.944
# alternative hypothesis: true  is less than 4
# sample estimates:
# mean of x
#      3.758
```

Case study

```
# This uses the plot.simtest function.  
plot(result, breaks = 100, freq = FALSE)
```

Histogram of x\$null.dist



Case study

- Write a function that does a one sample simulation test. Try it out on your favorite dataset.
- I put some example code to do a simulation test on GitHub:
<https://github.com/kflagg/useRS3/blob/master/code.r>
- At the end of the function, combine the variables into a list (use the variable names in the code below).
- Set the class to a vector containing "htest" and a second class like "simtest". Try running your function with and without setting the class.

```
result <- list(  
  # These are standard elements of htest objects.  
  estimate = ...,      # An estimate of the mean.  
  parameter = ...,     # Optional. I put the number of simulations here.  
  p.value = ...,       # Your p-value.  
  null.value = ...,    # The value of the mean under the null hypothesis.  
  alternative = ...,    # The text "less" for a left-tailed test.  
  method = ...,        # A name for the method, like "Simulation Test"  
  data.name = ...,     # The name of the data vector (text).  
  
  # This is something new for the simtest class.  
  null.dist = ...      # The vector of simulated sample means.  
)  
  
class(result) <- c(...) # Make the class "htest" and "simtest".
```

Case study

- Write a function that
 - takes at least one argument (a `simtest` object),
 - plots a histogram of the object's `null.dist` variable,
 - and draws a vertical line at the observed value (the estimate).
- Name the function `plot.simtest`.

```
plot.simtest <- function(...){  
  ...  
}  
  
plot(result)
```

Today we learned...

- How object-oriented programming works in R.
 - Classes are created by changing `class(object) <-`
 - Methods are functions named like `method.class`.
 - You run `method(object)` and R will automatically use `method.class(object)`.
- How some familiar objects are structured.
 - Most are just lists.
- How to create a new class and write a method.

R Core Team (2017). *R Language Definition*. Version 3.4.2. URL: <https://cran.r-project.org/doc/manuals/r-release/R-lang.pdf>.