

R programming: generics

2025-03-21

M1 MIDS/MFA/LOGOS

[Université Paris Cité](#)

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```
stopifnot(  
  require(Hmisc),  
  require(skimr),  
  require(patchwork),  
  require(ggforce),  
  require(glue),  
  require(ggfortify),  
  require(broom),  
  require(tidyverse)  
)  
  
tidymodels::tidymodels_prefer(quiet = TRUE)  
  
old_theme <- theme_set(  
  theme_minimal(base_size=9,  
                base_family = "Helvetica")  
)
```

```
gc <- options(ggplot2.discrete.colour="viridis")  
gc <- options(ggplot2.discrete.fill="viridis")  
gc <- options(ggplot2.continuous.fill="viridis")  
gc <- options(ggplot2.continuous.colour="viridis")
```

Objectives

```
stopifnot(  
  require(rlang),  
  require(lobstr),  
  require(sloop),  
  require(devtools),  
  require(usethis),  
  require(testthat),  
  require(generics)  
)
```

Loading required package: rlang
Loading required package: lobster
Loading required package: sloop
Loading required package: devtools
Loading required package: usethis
Loading required package: testthat
Loading required package: generics

Generics and S3 classes

Let us first create an instance of class `lm`.

```
lm0 <- lm(Gas ~ Insul * Temp, MASS::whiteside)
```

i Question

- What does function `class()` do?
- Is it possible to belong to type `list` and to class `lm` simultaneously?
- In R what is an *attribute*?
- How do we *set* and *get* attributes?
- What does function `inherits()` do?

i Question

Load package `sloop`.

- What does `sloop::otype()` do? Apply it to an object of class `lm`.
- What happens when we first `unclass()` the object?

i Question

`sloop` exports functions `s3_class()` and `s3_get_method()`

- Apply `s3_class()` to all members of `lm0`
- What is the `otype` of `autoplot()` applied to an object of class `lm`?
-

i Question

i Question

i Question

i Question

- [OO in Advanced R Programming 1st Edition](#)

- [S3 classes Advanced R 2nd Edition](#)

Programming with dplyr and ggplot2

We first aim at programming a function that takes as input a dataframe `df`, a column name `col`, and that, depending on the type of the column denoted by `col`, plots a histogram (for numerical column), a barplot (for factors), or raise an error if the column is neither categorical, nor numerical.

The function should return a ggplot object.

Here is a first attempt.

Let us first build a toy tibble.

```
tb <- tibble(
  col_num = rnorm(100),
  col_fac = as_factor(sample(letters, 100, replace = T)),
  col_ts = Sys.time() + duration(sample(1:20, 100, replace=T), units="days")
)

tb |>
  head()
```

```
# A tibble: 6 x 3
  col_num col_fac col_ts
  <dbl> <fct> <dtm>
1  0.139 d      2025-04-02 16:52:06
2  1.65 v      2025-04-07 16:52:06
3 -0.715 k      2025-04-05 16:52:06
4 -2.02 q      2025-04-10 16:52:06
5 -0.385 o      2025-04-03 16:52:06
6  0.0715 g      2025-03-24 15:52:06
```

i Question

- Pass more optional arguments to `geom_...` (use ellipsis `...`)
- Avoid quoting the column name

i Question

i Question

How could you add a `geom_point()` layer to each element of the following list?

```
plots <- list(
  ggplot(mpg, aes(displ, hwy)),
  ggplot(diamonds, aes(carat, price)),
  ggplot(faithfuld, aes(waiting, eruptions, size = density))
)
```

From [R Advanced Programming](#)

Inside `lm()`**i** Question

In classes like `lm`, `prcomp`, ... we have a member called `call`. What does it represent?
How is it constructed?

👉 First, read the code of `lm`.

```
> lm
function (formula, data, subset, weights, na.action, method = "qr",
  model = TRUE, x = FALSE, y = FALSE, qr = TRUE, singular.ok = TRUE,
  contrasts = NULL, offset, ...)
{
  ret.x <- x
  ret.y <- y
  cl <- match.call()
  mf <- match.call(expand.dots = FALSE)
  m <- match(c("formula", "data", "subset", "weights", "na.action",
    "offset"), names(mf), 0L)
  mf <- mf[c(1L, m)]
  mf$drop.unused.levels <- TRUE
  mf[[1L]] <- quote(stats::model.frame)
  mf <- eval(mf, parent.frame())
  if (method == "model.frame")
    return(mf)
  else if (method != "qr")
    warning(gettextf("method = '%s' is not supported. Using 'qr'",
      method), domain = NA)
  mt <- attr(mf, "terms")
  y <- model.response(mf, "numeric")
  w <- as.vector(model.weights(mf))
  if (!is.null(w) && !is.numeric(w))
    stop("'weights' must be a numeric vector")
  offset <- model.offset(mf)
  mlm <- is.matrix(y)
  ny <- if (mlm)
    nrow(y)
  else length(y)
  if (!is.null(offset)) {
    if (!mlm)
      offset <- as.vector(offset)
    if (NROW(offset) != ny)
      stop(gettextf("number of offsets is %d, should equal %d (number of observation",
        NROW(offset), ny), domain = NA)
  }
  if (is.empty.model(mt)) {
    x <- NULL
    z <- list(coefficients = if (mlm) matrix(NA_real_, 0,
      ncol(y)) else numeric(),
      residuals = y,
      fitted.values = 0 * y,
      weights = w,
      rank = 0L,
```

```
      df$residual = if (!is.null(w)) sum(w != 0) else ny
    )
    if (!is.null(offset)) {
      z$fitted.values <- offset
      z$residuals <- y - offset
    }
  }
  else {
    x <- model.matrix(mt, mf, contrasts)
    z <- if (is.null(w))
      lm.fit(x, y, offset = offset, singular.ok = singular.ok,
            ...)
    else lm.wfit(x, y, w, offset = offset, singular.ok = singular.ok,
                ...)
  }
  class(z) <- c(if (mlm) "mlm", "lm")
  z$na.action <- attr(mf, "na.action")
  z$offset <- offset
  z$contrasts <- attr(x, "contrasts")
  z$xlevels <- .getXlevels(mt, mf)
  z$call <- cl
  z$terms <- mt
  if (model)
    z$model <- mf
  if (ret.x)
    z$x <- x
  if (ret.y)
    z$y <- y
  if (!qr)
    z$qr <- NULL
  z
}
<bytecode: 0x55564224e930>
<environment: namespace:stats>
```

i Question

Have a look at function `match.call()`

Data masking and environments**i Question****Tidy evaluation****i Question**

What is *quasi-quotation*?
Keep the `rlang` cheatsheet around.

i Question

Explain the difference between an *expression* and a *quosure*

i Question

References

[Programming with ggplot](#)