Rangkuman Hands-on Programming with R

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17 December 2020

Chapter 1 The Very Basics

Chapter 1 dari buku **Hands-on Programming with R**. Catatan ini merupakan bagian dari proyek 1: **Weighted Dice**.

The R User Interface

Tidak ada yang penting.

Objects

Tidak ada yang penting.

Functions

round function is used to round a number.

```
round(3.1415)
```

```
## [1] 3
```

factorial function is used to calculate the factorial of a number.

```
factorial(3)
```

```
## [1] 6
```

mean function is used to compute the mean of number(s).

```
mean(1:6)
```

```
## [1] 3.5
```

We can combine round and mean functions as follows:

```
round(mean(1:6))
```

[1] 4

Sample with Replacement

The sample function is used to sample.

```
die <- 1:6
sample(x = die, size=1)</pre>
```

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

args can be used to view the arguments of a function.

```
args(round)
## function (x, digits = 0)
## NULL
args(sample)
## function (x, size, replace = FALSE, prob = NULL)
## NULL
round method can be accompanied with digits, for example:
round(3.1415, digits = 2)
## [1] 3.14
By default, the sample function is without replacement.
sample(die, size=2)
## [1] 5 3
When we want to sample with replacement, we can set replace=TRUE as follows:
sample(die, size=2, replace = TRUE)
## [1] 1 1
If you want to add up the dice, you can feed your result straight into the sum function:
dice <- sample(die, size=2, replace = TRUE)</pre>
dice
## [1] 2 4
sum(dice)
## [1] 6
die <- 1:6
```

Writing Your Own Functions

We are going to write a function which returns the sum of rolling two dice.

```
roll <- function(){
  die <- 1:6
  dice <- sample(die, size=2, replace = TRUE)
  sum(dice)
}
result <- roll()
result</pre>
```

[1] 3

When you run a function in R, R will execute all of the code in the body and then return the result of the last line of code.

The Function Constructor

Arguments

- The name. A user can run the function by typing the name followed by parentheses, e.g., roll2().
- 3. **The arguments**. A user can supply values for these variables, which appear in the body of the function.
- The default values.
 Optional values that R can use for the arguments if a user does not supply a value.

- 2. **The body**. R will run this code whenever a user calls the function.
- roll2 <- function(bones = 1:6) {
 dice <- sample(bones, size = 2,
 replace = TRUE)
 sum(dice)
 }</pre>
- 5. **The last line of code**. The function will return the result of the last line.

```
\#\# Scripts
```

Menggunakan qplot dengan data pakai c.

```
library("ggplot2")

x <- c(-1, -0.8, -0.6, -0.4, -0.2, 0, 0.2, 0.4, 0.6, 0.8, 1)

## [1] -1.0 -0.8 -0.6 -0.4 -0.2 0.0 0.2 0.4 0.6 0.8 1.0

## -1.0 -0.8 -0.6 -0.4 -0.2 0.0 0.2 0.4 0.6 0.8 1.0

y <- x^3

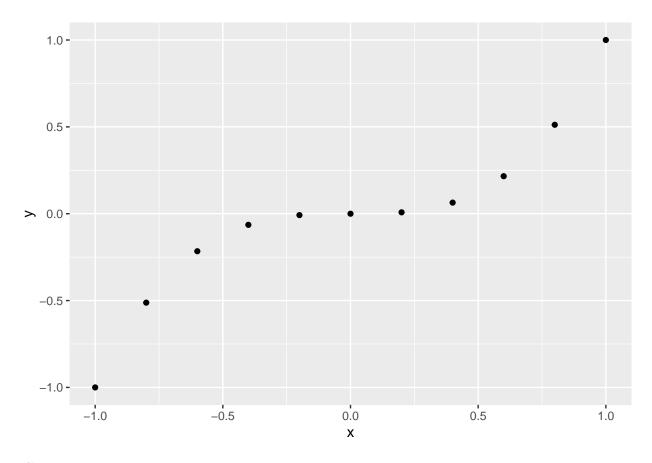
y

## [1] -1.000 -0.512 -0.216 -0.064 -0.008 0.000 0.008 0.064 0.216 0.512

## [11] 1.000

## -1.000 -0.512 -0.216 -0.064 -0.008 0.000 0.008 0.064 0.216 0.512 1.000

qplot(x,y)
```



Summary

Packages and Help Pages

Packages

- Package qplot berarti quick plot.
- Berikut adalah fungsinya dan qplot is a long function

```
library(ggplot2)
qplot
```

```
## function (x, y, ..., data, facets = NULL, margins = FALSE, geom = "auto",
       xlim = c(NA, NA), ylim = c(NA, NA), log = "", main = NULL,
##
##
       xlab = NULL, ylab = NULL, asp = NA, stat = deprecated(),
       position = deprecated())
##
## {
       deprecate_soft0("3.4.0", "qplot()")
##
       caller_env <- parent.frame()</pre>
##
##
       if (lifecycle::is_present(stat))
##
           lifecycle::deprecate_stop("2.0.0", "qplot(stat)")
##
       if (lifecycle::is_present(position))
           lifecycle::deprecate_stop("2.0.0", "qplot(position)")
##
##
       check_character(geom)
##
       exprs <- enquos(x = x, y = y, ...)
##
       is_missing <- vapply(exprs, quo_is_missing, logical(1))</pre>
##
       is_constant <- (!names(exprs) %in% ggplot_global$all_aesthetics) |</pre>
           vapply(exprs, quo_is_call, logical(1), name = "I")
##
```

```
##
        mapping <- new_aes(exprs[!is_missing & !is_constant], env = parent.frame())</pre>
##
        consts <- exprs[is_constant]</pre>
        aes_names <- names(mapping)</pre>
##
        mapping <- rename_aes(mapping)</pre>
##
##
        if (is.null(xlab)) {
            if (quo_is_missing(exprs$x)) {
##
##
                 xlab <- ""
            }
##
            else {
##
##
                 xlab <- as_label(exprs$x)</pre>
##
        }
##
        if (is.null(ylab)) {
##
##
            if (quo_is_missing(exprs$y)) {
                 ylab <- ""
##
##
            }
##
            else {
##
                 ylab <- as_label(exprs$y)</pre>
##
##
        }
##
        if (missing(data)) {
##
            data <- data frameO()</pre>
            facetvars <- all.vars(facets)</pre>
##
            facetvars <- facetvars[facetvars != "."]</pre>
##
            names(facetvars) <- facetvars</pre>
##
##
            facetsdf <- as.data.frame(mget(facetvars, envir = caller_env))</pre>
##
            if (nrow(facetsdf))
##
                 data <- facetsdf
##
        }
        if ("auto" %in% geom) {
##
##
            if ("sample" %in% aes_names) {
##
                 geom[geom == "auto"] <- "qq"</pre>
            }
##
##
            else if (missing(y)) {
##
                 x <- eval_tidy(mapping$x, data, caller_env)</pre>
##
                 if (is.discrete(x)) {
##
                     geom[geom == "auto"] <- "bar"</pre>
##
                 }
##
                 else {
                     geom[geom == "auto"] <- "histogram"</pre>
##
##
                 }
##
                 if (is.null(ylab))
                     ylab <- "count"
##
            }
##
##
            else {
##
                 if (missing(x)) {
##
                     mapping$x <- quo(seq_along(!!mapping$y))</pre>
##
                 }
##
                 geom[geom == "auto"] <- "point"</pre>
            }
##
##
       p <- ggplot(data, mapping, environment = caller_env)</pre>
##
##
        if (is.null(facets)) {
##
            p <- p + facet_null()</pre>
```

```
}
##
        else if (is.formula(facets) && length(facets) == 2) {
##
            p <- p + facet_wrap(facets)</pre>
##
##
        }
##
        else {
##
            p <- p + facet_grid(rows = deparse(facets), margins = margins)</pre>
##
        if (!is.null(main))
##
##
            p <- p + ggtitle(main)</pre>
##
        for (g in geom) {
##
            params <- lapply(consts, eval_tidy)</pre>
            p <- p + do.call(paste0("geom_", g), params)</pre>
##
##
##
        logv <- function(var) var %in% strsplit(log, "")[[1]]</pre>
##
        if (logv("x"))
##
            p \leftarrow p + scale_x_log10()
##
        if (logv("y"))
##
            p \leftarrow p + scale_y = log10()
##
        if (!is.na(asp))
            p <- p + theme(aspect.ratio = asp)</pre>
##
##
        if (!missing(xlab))
##
            p \leftarrow p + xlab(xlab)
        if (!missing(ylab))
##
##
            p \leftarrow p + ylab(ylab)
##
        if (!missing(xlim) && !all(is.na(xlim)))
            p \leftarrow p + xlim(xlim)
##
##
        if (!missing(ylim) && !all(is.na(ylim)))
##
            p \leftarrow p + ylim(ylim)
##
## }
## <bytecode: 0x564fcd4b5f68>
## <environment: namespace:ggplot2>
```

- If you give qplot two vectors of equal lengths, qplot will draw a scatterplot for you.
- qplot will use the first vector as a set of x values and the second vector as a set of y values.
- Until now, we've been creating sequences of numbers with the : operator; but you can also create vectors of numbers with the c function.
- Give c all of the numbers that you want to appear in the vector, separated by a comma. c stands for *concatenate*, but you can think of it as "collect" or "combine"/