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

Objects

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)
```

```
## [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] 2 5
```

When we want to `sample` *with replacement*, we can set `replace=TRUE` as follows:

```
sample(die, size=2, replace = TRUE)
```

```
## [1] 4 2
```

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)
dice
```

```
## [1] 2 2
```

```
sum(dice)
```

```
## [1] 4
```

```
die <- 1:6
```

Writing Your Own Functions

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

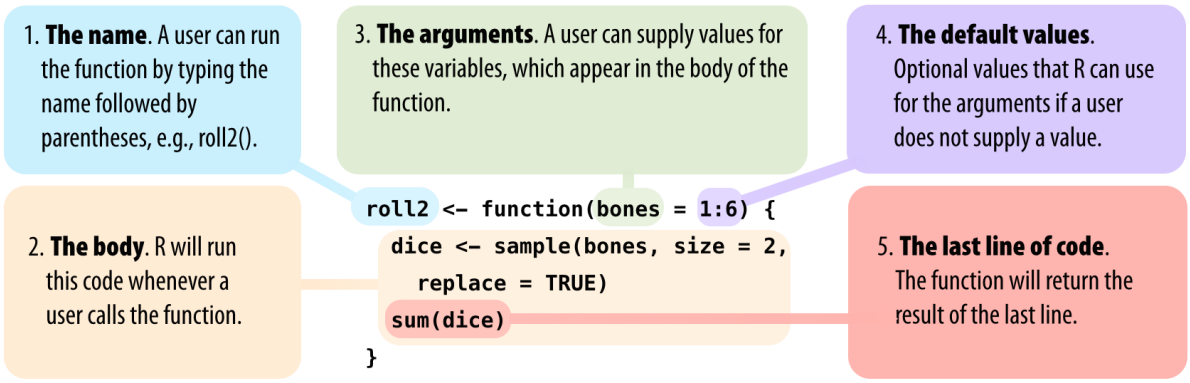
```
roll <- function(){
  die <- 1:6
  dice <- sample(die, size=2, replace = TRUE)
  sum(dice)
}
result <-  roll()
result
```

```
## [1] 11
```

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



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)
x
```

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

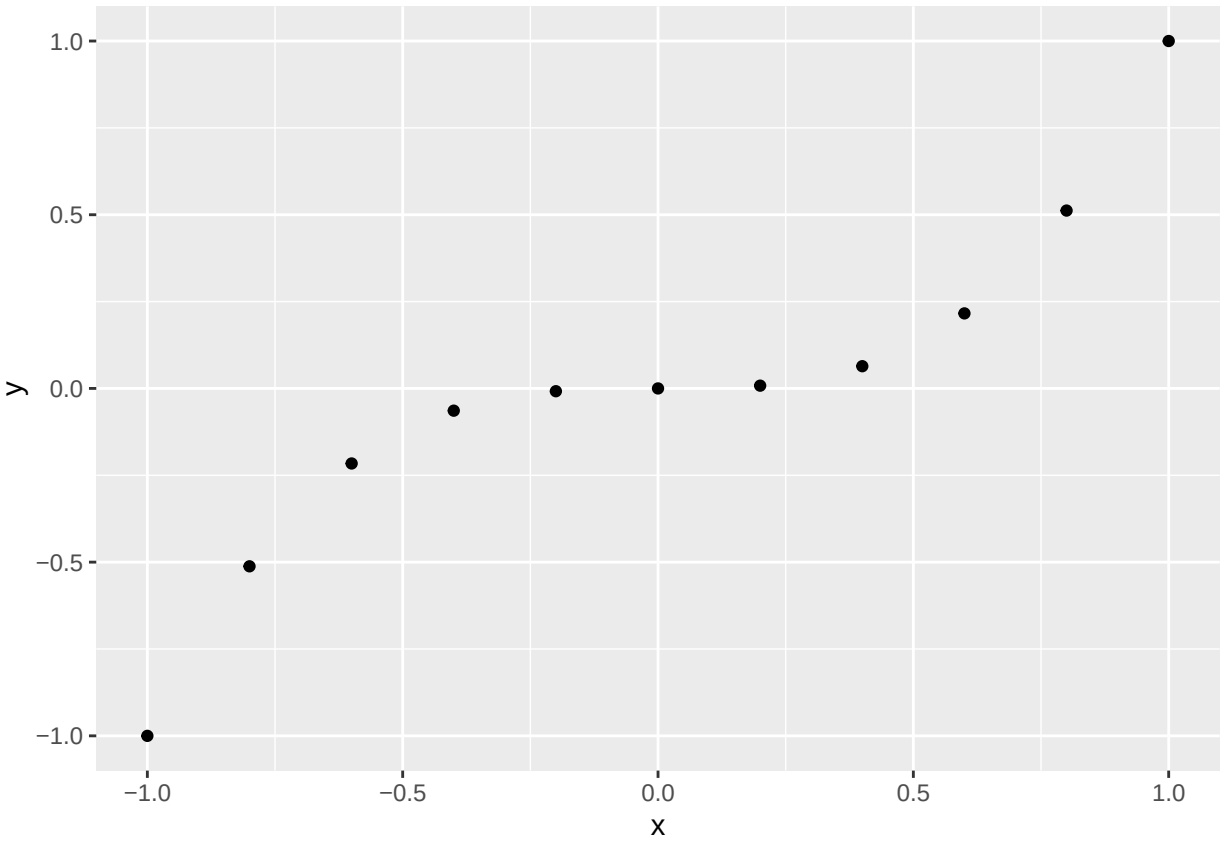
```
##  -1.0 -0.8 -0.6 -0.4 -0.2  0.0  0.2  0.4  0.6  0.8  1.0

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()
##     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))
##     is_constant <- (!names(exprs) %in% ggplot_global$all_aesthetics) |
##         vapply(exprs, quo_is_call, logical(1), name = "I")
```

```
##     mapping <- new_aes(exprs[!is_missing & !is_constant], env = parent.frame())
##     consts <- exprs[is_constant]
##     aes_names <- names(mapping)
##     mapping <- rename_aes(mapping)
##     if (is.null(xlab)) {
##         if (quo_is_missing(exprs$x)) {
##             xlab <- ""
##         }
##         else {
##             xlab <- as_label(exprs$x)
##         }
##     }
##     if (is.null(ylab)) {
##         if (quo_is_missing(exprs$y)) {
##             ylab <- ""
##         }
##         else {
##             ylab <- as_label(exprs$y)
##         }
##     }
##     if (missing(data)) {
##         data <- data_frame0()
##         facetvars <- all.vars(facets)
##         facetvars <- facetvars[facetvars != "."]
##         names(facetvars) <- facetvars
##         facetsdf <- as.data.frame(mget(facetvars, envir = caller_env))
##         if (nrow(facetsdf))
##             data <- facetsdf
##     }
##     if ("auto" %in% geom) {
##         if ("sample" %in% aes_names) {
##             geom[geom == "auto"] <- "qq"
##         }
##         else if (missing(y)) {
##             x <- eval_tidy(mapping$x, data, caller_env)
##             if (is.discrete(x)) {
##                 geom[geom == "auto"] <- "bar"
##             }
##             else {
##                 geom[geom == "auto"] <- "histogram"
##             }
##             if (is.null(ylab))
##                 ylab <- "count"
##         }
##         else {
##             if (missing(x)) {
##                 mapping$x <- quo(seq_along(!!mapping$y))
##             }
##             geom[geom == "auto"] <- "point"
##         }
##     }
##     p <- ggplot(data, mapping, environment = caller_env)
##     if (is.null(facets)) {
##         p <- p + facet_null()
```

```
##     }
##     else if (is.formula(facets) && length(facets) == 2) {
##         p <- p + facet_wrap(facets)
##     }
##     else {
##         p <- p + facet_grid(rows = deparse(facets), margins = margins)
##     }
##     if (!is.null(main))
##         p <- p + ggtitle(main)
##     for (g in geom) {
##         params <- lapply(consts, eval_tidy)
##         p <- p + do.call(paste0("geom_", g), params)
##     }
##     logv <- function(var) var %in% strsplit(log, "")[[1]]
##     if (logv("x"))
##         p <- p + scale_x_log10()
##     if (logv("y"))
##         p <- p + scale_y_log10()
##     if (!is.na(asp))
##         p <- p + theme(aspect.ratio = asp)
##     if (!missing(xlab))
##         p <- p + xlab(xlab)
##     if (!missing(ylab))
##         p <- p + ylab(ylab)
##     if (!missing(xlim) && !all(is.na(xlim)))
##         p <- p + xlim(xlim)
##     if (!missing(ylim) && !all(is.na(ylim)))
##         p <- p + ylim(ylim)
##     p
## }
## <bytecode: 0x560e0108f918>
## <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"/