Types, vectors, and functions in R

2019-08-29

Vectors and Types

c(1, 3, 5) c(TRUE, FALSE, TRUE, TRUE) c("red", "blue")

Vectors have 1 dimension

Vectors have 1 dimension

Vectors have a length.

length(c("blue", "red"))

Vectors have 1 dimension

Vectors have a length.

length(c("blue", "red"))

Some vectors have names.

Vectors have 1 dimension

Vectors have a length.

length(c("blue", "red"))

Some vectors have names.

names(c("x" = 1, "y = 1))

Vectors have types

Types

Numeric/double

Integer

Factor

Character

Logical

Dates

Packages to work with types:

Strings/character: stringr

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Strings/character: stringr

Factors: forcats

Packages to work with types:

Strings/character: stringr

Factors: forcats

Dates: lubridate

Making vectors

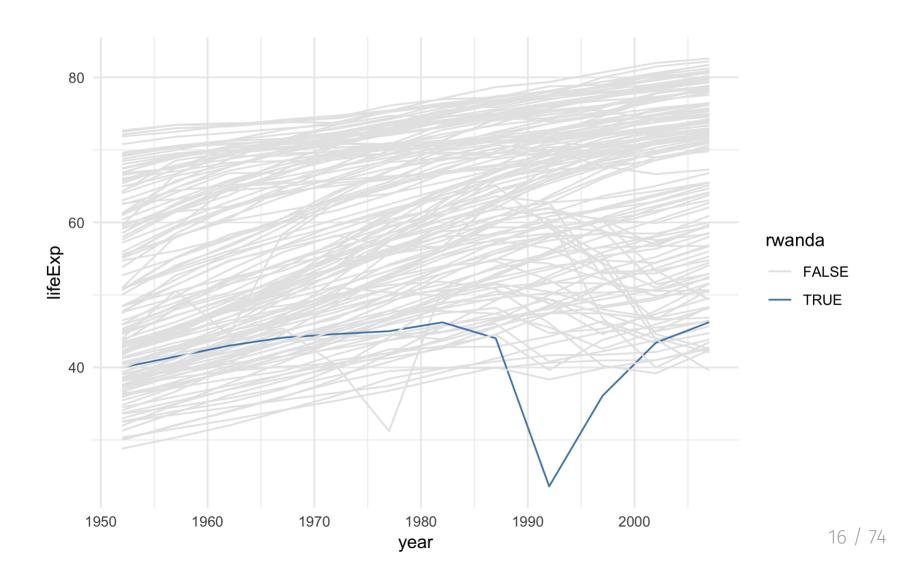
```
1:3
## [1] 1 2 3
c(1, 2, 3)
## [1] 1 2 3
rep(1, 3)
## [1] 1 1 1
seq(from = 1, to = 3, by = .5)
## [1] 1.0 1.5 2.0 2.5 3.0
```

Create a character vector of colors using c(). Use the colors "grey90" and "steelblue". Assign the vector to a name.

Use the vector you just created to change the colors in the plot below using scale_color_manual(). Pass it using the values argument.

```
cols <- c("grey90", "steelblue")

gapminder %>%
  mutate(rwanda = ifelse(country == "Rwanda", TRUE, FALSE)) %>%
  ggplot(aes(year, lifeExp, color = rwanda, group = country)) +
  geom_line() +
  scale_color_manual(values = cols) +
  theme_minimal()
```



Working with vectors

Subset vectors with [] or [[]]

```
x < -c(1, 5, 7)
x[2]
## [1] 5
x[[2]]
## [1] 5
x[c(FALSE, TRUE, FALSE)]
## [1] 5
```

Working with vectors

Modify elements

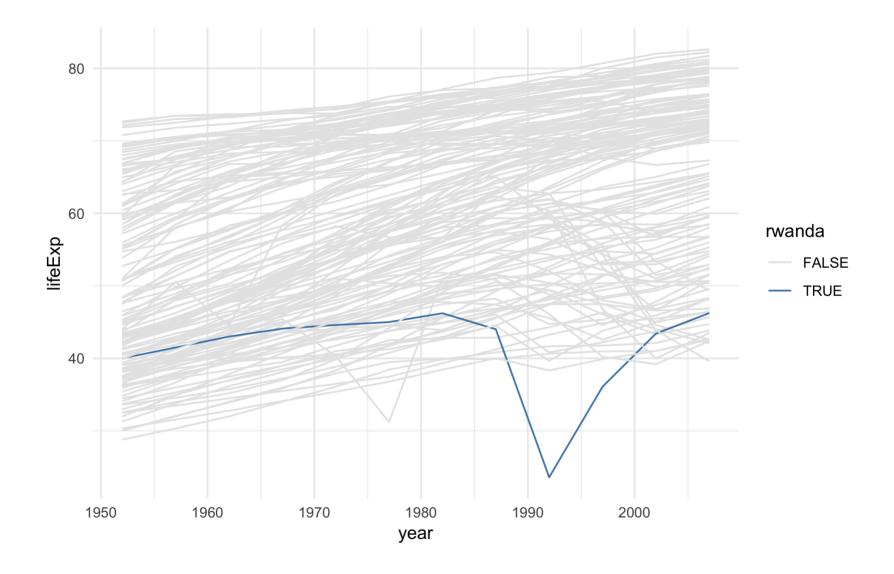
```
x
## [1] 1 5 7
x[2] <- 100
x
## [1] 1 100 7</pre>
```

Modify elements

```
x
### [1] 1 100 7

x[x > 10] <- NA

x
### [1] 1 NA 7</pre>
```



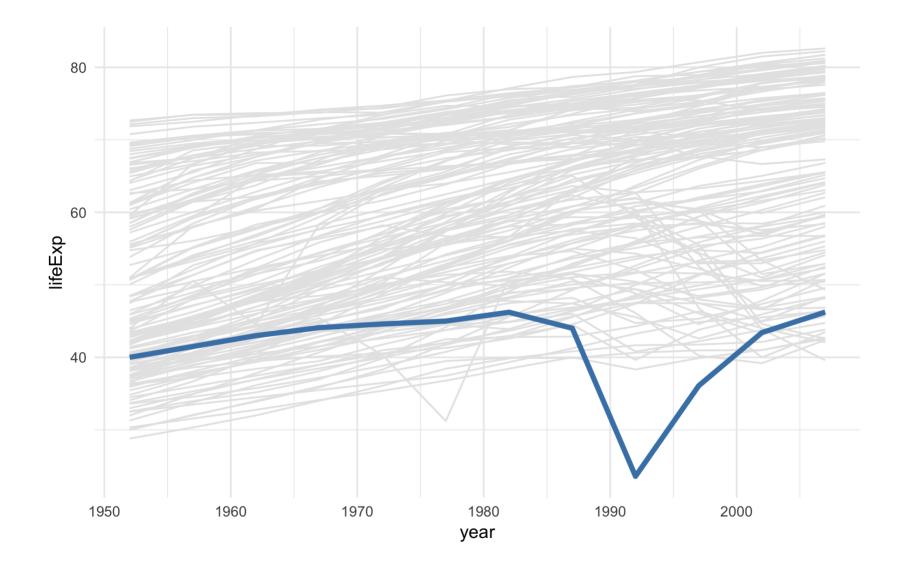
```
cols <- c("grey90", "steelblue")

gapminder %>%
  mutate(rwanda = ifelse(country == "Rwanda", TRUE, FALSE)) %>%
  ggplot(aes(year, lifeExp, color = rwanda, group = country)) +
  geom_line() +
  scale_color_manual(values = cols) +
  theme_minimal()
```

```
cols <- c("grey90", "steelblue")

gapminder %>%
  mutate(rwanda = ifelse(country == "Rwanda", TRUE, FALSE)) %>%
  ggplot(aes(year, lifeExp, group = country)) +
  geom_line(
    data = function(x) filter(x, !rwanda),
    color = cols[1]
  ) +
  theme_minimal()
```

```
cols <- c("grey90", "steelblue")</pre>
gapminder %>%
  mutate(rwanda = ifelse(country == "Rwanda", TRUE, FALSE)) %>%
  ggplot(aes(year, lifeExp, color = rwanda, group = country)) +
  geom line(
    data = function(x) filter(x, !rwanda),
    color = cols[1]
  ) +
  geom line(
    data = function(x) filter(x, rwanda),
    color = cols[2],
    size = 1.5
  ) +
  theme minimal()
```



Create a numeric vector that has the following values: 3, 5, NA, 2, and NA.

Try using sum(). Then add na.rm = TRUE.

Check which values are missing with is.na(); save the results to a new object and take a look

Chance all missing values of x to 0

Try sum() again without na.rm = TRUE.

```
x <- c(3, 5, NA, 2, NA)
sum(x)
```

[1] NA

```
sum(x, na.rm = TRUE)
```

[1] 10

```
x_missing <- is.na(x)</pre>
x_missing
## [1] FALSE FALSE TRUE FALSE TRUE
x[x_missing] <- 0</pre>
X
## [1] 3 5 0 2 0
sum(x)
## [1] 10
```

Function arguments

```
add_one <- function(x) {
    x <- x + 1
    x
}
add_one(1)
#> 2
Create function
```

```
function name
 add_one <- function(x) {</pre>
    x < - x + 1
                function
                    body
- add_one(1)
#> 2
```

```
add_one <- function(x) {
    x <- x + 1
    X Output
}
add_one(1) — input
#> 2
```

Create a function called sim_data that doesn't take any arguments.

In the function body, we'll return a tibble.

For x, have rnorm() return 50 random numbers.

For sex, use rep() to create 50 values of "male" and "female". Hint: You'll have to give rep() a character vector. for the first argument. The times argument is how many times rep() should repeat the first argument, so make sure you 3. account for that.

For age() use the sample() function to sample 50 numbers from 25 to 50 with replacement.

Call sim_data()

```
sim_data <- function() {
  tibble(
    x = rnorm(50),
    sex = rep(c("male", "female"), times = 25),
    age = sample(25:50, size = 50, replace = TRUE)
  )
}
sim_data()</pre>
```

```
sim_data <- function() {
   tibble(
        x = rnorm(50),
        sex = rep(c("male", "female"), times = 25),
        age = sample(25:50, size = 50, replace = TRUE)
   )
}
sim_data()</pre>
```

```
sim_data <- function() {
   tibble(
        x = rnorm(50),
        sex = rep(c("male", "female"), times = 25),
        age = sample(25:50, size = 50, replace = TRUE)
   )
}
sim_data()</pre>
```

```
sim_data <- function() {
  tibble(
    x = rnorm(50),
    sex = rep(c("male", "female"), times = 25),
    age = sample(25:50, size = 50, replace = TRUE)
  )
}
sim_data()</pre>
```

```
## # A tibble: 50 x 3
##
           x sex
                      age
##
   <dbl> <chr>
                    <int>
##
      0.0202 male
                       34
   2 -1.10 female
                       42
##
##
   3 0.200 male
                       40
##
   4 0.357 female
                       40
##
  5 -0.0356 male
                       27
  6 -0.248 female
                       44
##
   7 0.233 male
                       29
##
## 8 -0.760 female
                       34
##
   9 -0.0736 male
                       50
     1.06
             female
## 10
                       46
排排 排 ... with 40 more rows
```

E-Values

The strength of unmeasured confounding required to explain away a value

E-Values

The strength of unmeasured confounding required to explain away a value

Rate ratio: 3.9 = E-value: 7.3

Write a function to calculate an E-Value given an RR.

Call the function evalue and give it an argument called estimate. In the body of the function, calculate the E-Value using estimate + sqrt(estimate * (estimate - 1))

Call evalue() for a risk ratio of 3.9

```
evalue <- function(estimate) {
  estimate + sqrt(estimate * (estimate - 1))
}
evalue(3.9)
## [1] 7.263034</pre>
```

Control Flow

```
if (PREDICATE) {
  true_result
3
if (PREDICATE) {
  true_result
} else {
  default_result
3
if (PREDICATE) {
  true result
} else if (ANOTHER_PREDICATE) {
  true_result
} else {
  default_result
3
```

Other functions to control flow

```
ifelse(PREDICATE, true_result, false_result)
dplyr::case_when(
    PREDICATE ~ true_result,
    PREDICATE ~ true_result,
    TRUE ~ default_result
)
switch(
    X,
    value1 = result,
    value2 = result
)
```

Validation and stopping if (is.numeric(x)) stop(), warn()

```
function(x) {
  if (is.numeric(x)) stop("x must be a character")
  # do something with a character
}
```

Use if () together with is.numeric() to make sure estimate is a number. Remember to use! for not.

If the estimate is less than 1, set estimate to be equal to 1 / estimate.

Call evalue() for a risk ratio of 3.9. Then try 0.80. Then try a character value.

```
evalue <- function(estimate) {
  if (!is.numeric(estimate)) stop("`estimate` must be numeric")
  if (estimate < 1) estimate <- 1 / estimate
  estimate + sqrt(estimate * (estimate - 1))
}</pre>
```

```
evalue(3.9)

## [1] 7.263034

evalue(.80)

## [1] 1.809017

evalue("3.9")

## Error in evalue("3.9"): `estimate` must be numeric
```

Add a new argument called type. Set the default value to "rr"

Check if type is equal to "or". If it is, set the value of estimate to be sqrt(estimate)

Call evalue() for a risk ratio of 3.9. Then try it again
with type = "or".

```
evalue <- function(estimate, type = "rr") {
  if (!is.numeric(estimate)) stop("`estimate` must be numeric")
  if (type == "or") estimate <- sqrt(estimate)
  if (estimate < 1) estimate <- 1 / estimate
  estimate + sqrt(estimate * (estimate - 1))
}</pre>
```

```
evalue(3.9)

### [1] 7.263034

evalue(3.9, type = "or")

### [1] 3.362342
```

Create a new function called transform_to_rr with arguments estimate and type.

Use the same code above to check if type == "or" and transform if so. Add another line that checks if type == "hr". If it does, transform the estimate using this formula: (1 - 0.5^sqrt(estimate)) / (1 - 0.5^sqrt(1 / estimate)).

Move the code that checks if estimate < 1 to transform_to_rr (below the OR and HR transformations)

Return estimate

In evalue(), change the default argument of type to be a character vector containing "rr", "or", and "hr".

Get and validate the value of type **using** match.arg(). **Follow the pattern** argument_name <- match.arg(argument_name)

Transform estimate **using** transform_to_rr(). **Don't forget to pass it both** estimate **and** type!

```
transform to rr <- function(estimate, type) {
 if (type == "or") estimate <- sqrt(estimate)</pre>
  if (type == "hr") {
    estimate <-
      (1 - 0.5^{sqrt}(estimate)) / (1 - 0.5^{sqrt}(1 / estimate))
  7
 if (estimate < 1) estimate <- 1 / estimate</pre>
 estimate
3
evalue <- function(estimate, type = c("rr", "or", "hr")) {
 # validate arguments
 if (!is.numeric(estimate)) stop("`estimate` must be numeric")
 type <- match.arg(type)</pre>
 # calculate evalue
 estimate <- transform_to_rr(estimate, type)</pre>
 estimate + sqrt(estimate * (estimate - 1))
3
```

```
transform to rr <- function(estimate, type) {
  if (type == "or") estimate <- sqrt(estimate)</pre>
  if (type == "hr") {
    estimate <-
      (1 - 0.5^sqrt(estimate)) / (1 - 0.5^sqrt(1 / estimate))
  3
  if (estimate < 1) estimate <- 1 / estimate</pre>
  estimate
3
evalue <- function(estimate, type = c("rr", "or", "hr")) {
  # validate arguments
  if (!is.numeric(estimate)) stop("`estimate` must be numeric")
  type <- match.arg(type)</pre>
  # calculate evalue
  estimate <- transform_to_rr(estimate, type)</pre>
  estimate + sqrt(estimate * (estimate - 1))
3
```

```
transform to rr <- function(estimate, type) {
  if (type == "or") estimate <- sqrt(estimate)</pre>
  if (type == "hr") {
    estimate <-
      (1 - 0.5^{sqrt}(estimate)) / (1 - 0.5^{sqrt}(1 / estimate))
  3
  if (estimate < 1) estimate <- 1 / estimate</pre>
  estimate
3
evalue <- function(estimate, type = c("rr", "or", "hr")) {
  # validate arguments
  if (!is.numeric(estimate)) stop("`estimate` must be numeric")
  type <- match.arg(type)</pre>
  # calculate evalue
  estimate <- transform_to_rr(estimate, type)</pre>
  estimate + sqrt(estimate * (estimate - 1))
3
```

```
evalue(3.9)
## [1] 7.263034
evalue(3.9, type = "or")
## [1] 3.362342
evalue(3.9, type = "hr")
## [1] 4.474815
evalue(3.9, type = "rd")
## Error in match.arg(type): 'arg' should be one of "rr", "or", "hr"
```

Pass the dots: ...

```
select_gapminder <- function(...) {
   gapminder %>%
    select(...)
}
select_gapminder(pop, year)
```

Pass the dots: ...

```
select_gapminder <- function(...) {
  gapminder %>%
    select(...)
}
select_gapminder(pop, year)
```

Pass the dots: ...

```
## # A tibble: 1,704 x 2
##
           pop year
         <int> <int>
##
4⊧4⊧
   1 8425333 1952
4‡4‡
       9240934 1957
4‡4‡
   3 10267083 1962
4‡4‡
   4 11537966 1967
   5 13079460 1972
###
4‡4‡
   6 14880372 1977
   7 12881816 1982
4⊧4⊧
   8 13867957 1987
4⊧4⊧
   9 16317921 1992
##
## 10 22227415 1997
## # ... with 1,694 more rows
```

Use ... to pass the arguments of your function, filter_summarize(), to filter().

In summarize, get the n and mean life expectancy for the data set

Check filter_summarize() with year == 1952.

Try filter_summarize() again for 2002, but also filter countries that have "and" in the country name. Use str_detect() from the stringr package.

```
filter_summarize <- function(...) {
  gapminder %>%
  filter(...) %>%
  summarize(n = n(), mean_lifeExp = mean(lifeExp))
}
```

```
filter summarize(year == 1952)
## # A tibble: 1 x 2
### n mean_lifeExp
## <int> <dbl>
## 1 142 49.1
filter_summarize(year == 2002, str_detect(country, " and "))
## # A tibble: 1 x 2
### n mean_lifeExp
## <int> <dbl>
## 1 4 69.9
```

Programming with dplyr, ggplot2, and friends

```
plot_hist <- function(x) {
  ggplot(gapminder, aes(x = x)) + geom_histogram()
}</pre>
```

Programming with dplyr, ggplot2, and friends

```
plot_hist <- function(x) {
   ggplot(gapminder, aes(x = x)) + geom_histogram()
}

plot_hist(lifeExp)

## Error in FUN(X[[i]], ...): object 'lifeExp' not found</pre>
```

Programming with dplyr, ggplot2, and friends

```
plot_hist <- function(x) {
    ggplot(gapminder, aes(x = x)) + geom_histogram()
}

plot hist("lifeExp")</pre>
```

```
## Error: StatBin requires a continuous x variable: the x variable is dis
```

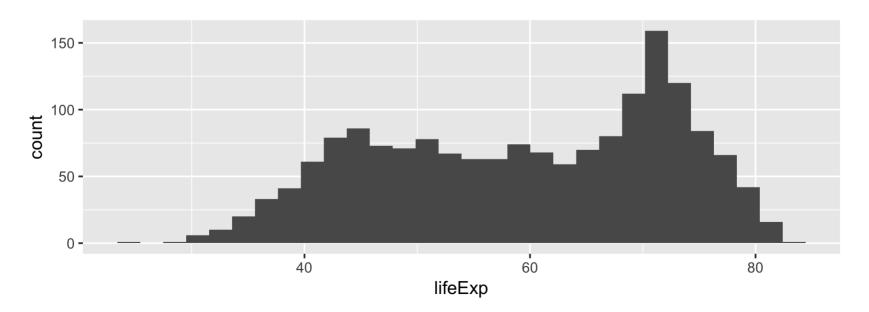
Curly-curly

```
plot_hist <- function(x) {
   ggplot(gapminder, aes(x = {{x}})) + geom_histogram()
}</pre>
```

Curly-curly

```
plot_hist <- function(x) {
  ggplot(gapminder, aes(x = {{x}})) + geom_histogram()
}</pre>
```

```
plot_hist(lifeExp)
```



Filter gapminder by year using the value of .year (notice the period before hand!). You do NOT need curly-curly for this. (Why is that?)

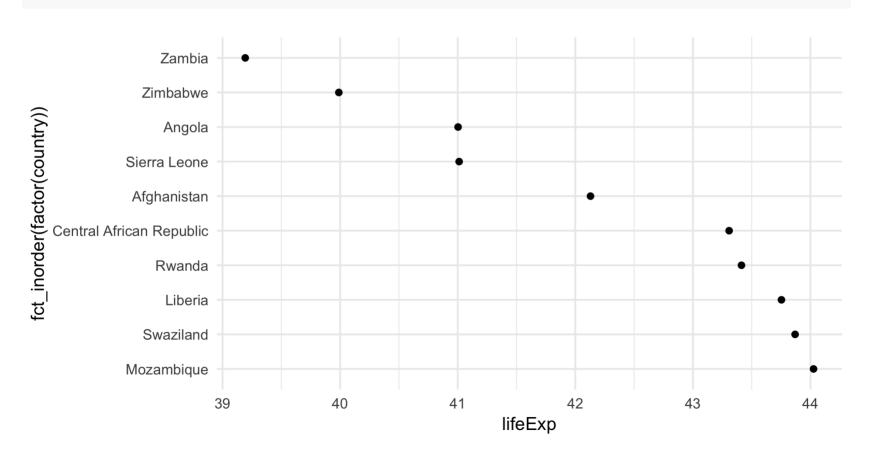
Arrange it by the variable. Don't forget to wrap it in curly-curly!

Make a scatter plot. Use variable for x. For y, we'll use country, but to keep it in the order we arranged it by, we'll turn it into a factor. Wrap the the factor() call with fct_inorder(). Check the help page if you want to know more about what this is doing.

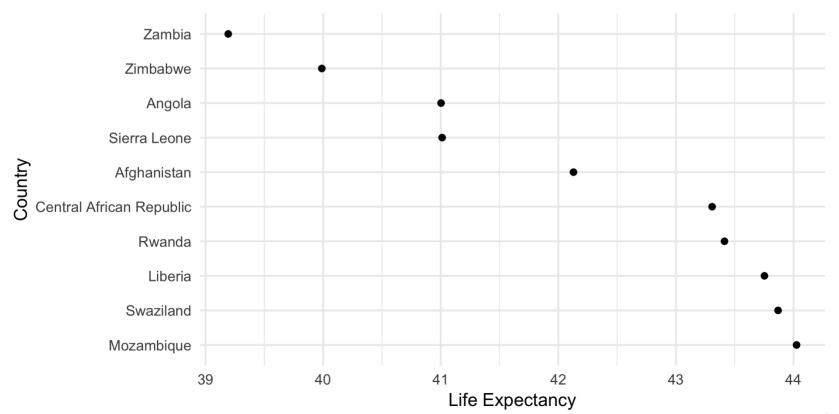
```
top_barplot <- function(variable, .year) {
   gapminder %>%
   filter(year == .year) %>%
   arrange(desc({{variable}})) %>%
   # take the 10 lowest values
   tail(10) %>%
   ggplot(aes(x = {{variable}}, y = fct_inorder(factor(country))))
      geom_point() +
      theme_minimal()
}
```

```
top_barplot <- function(variable, .year) {
  gapminder %>%
  filter(year == .year) %>%
  arrange(desc({{variable}})) %>%
  # take the 10 lowest values
  tail(10) %>%
  ggplot(aes(x = {{variable}}, y = fct_inorder(factor(country))))
   geom_point() +
   theme_minimal()
}
```

top_barplot(lifeExp, 2002)



```
top_barplot(lifeExp, 2002) +
  labs(x = "Life Expectancy", y = "Country")
```



Resources

R for Data Science: A comprehensive but friendly introduction to the tidyverse.

Free online.

Advanced R, 2nd ed.: Detailed guide to how R works and how to make your code better. Free online.

RStudio Primers: Free interactive courses in the Tidyverse