Functional programming and iteration with purrr

2019-08-29

purrr: A functional programming toolkit for R

Complete and consistent set of tools for working with functions and vectors

Problems we want to solve:

- Making code clear
- 2 Making code safe
- **3** Working with lists and data frames

Lists, vectors, and data.frames (or tibbles)

```
c(a = "hello", b = 1)

### a b
### "hello" "1"
```

lists can contain any object

```
list(a = "hello", b = 1, c = mean)

### $a

### [1] "hello"

### $b

### [1] 1

### $c

### function (x, ...)

### UseMethod("mean")

### <bytecode: 0x7fdb056a29b0>

### <environment: namespace:base>
```

Your Turn 1

```
number_list <- list(
    a = rnorm(10, mean = 2, sd = 1.4),
    b = rnorm(5, mean = 1, sd = 1),
    c = rnorm(20, mean = 10, sd = 3)
)</pre>
```

There are two ways to subset lists: dollar signs and brackets. Try to subset a from number_list using these approaches. Are they different? What about number_list[["a"]]?

Your Turn 1

```
number_list["a"]

### $a

### [1] 0.310108 2.388401 3.518218 -1.283977 2.600775 2.708478 1.195364

### [8] 1.234715 1.209767 0.753947

number_list$a

### [1] 0.310108 2.388401 3.518218 -1.283977 2.600775 2.708478 1.195364

### [8] 1.234715 1.209767 0.753947

number_list[["a"]]

### [1] 0.310108 2.388401 3.518218 -1.283977 2.600775 2.708478 1.195364

### [1] 0.310108 1.234715 1.209767 0.753947
```

```
library(gapminder)
head(gapminder$pop)
```

```
## [1] 8425333 9240934 10267083 11537966 13079460 14880372
```

```
gapminder[1:6, "pop"]
```

```
gapminder[1:6, "pop"]

### # A tibble: 6 x 1

### pop

### <int>
### 1 8425333

### 2 9240934

### 3 10267083

### 4 11537966

### 5 13079460

### 6 14880372
```

```
head(gapminder[["pop"]])
```

[1] 8425333 9240934 10267083 11537966 13079460 14880372

sum(rnorm(10))

```
sum(rnorm(10))

### [1] -3.831574
```

```
sum(rnorm(10))

### [1] -3.831574

sum(list(x = rnorm(10), y = rnorm(10), z = rnorm(10)))
```

```
sum(rnorm(10))

## [1] -3.831574

sum(list(x = rnorm(10), y = rnorm(10), z = rnorm(10)))

## Error in sum(list(x = rnorm(10), y = rnorm(10), z = rnorm(10)): invaluable
```

map(.x, .f)

.x: a vector, list, or data frame



.f: a function

Returns a list

```
library(purrr)
x_list <- list(x = rnorm(10), y = rnorm(10), z = rnorm(10))
map(x_list, mean)</pre>
```

```
library(purrr)
x_list <- list(x = rnorm(10), y = rnorm(10), z = rnorm(10))
map(x_list, mean)</pre>
```

```
library(purrr)
x_list <- list(x = rnorm(10), y = rnorm(10), z = rnorm(10))
map(x_list, mean)</pre>
```

```
library(purrr)
x_list <- list(x = rnorm(10), y = rnorm(10), z = rnorm(10))
map(x_list, mean)

### $x
### [1] -0.6097971
###
### $y
### [1] -0.2788647
###
### $z
### [1] 0.6165922</pre>
```

```
list()
 c("a", "b", "c")
 c("a", "b", "c")
 c("a", "b", "c")
```

map(| list() , .f) **f** (| c("a", "b", "c") |) **f** (| c("a", "b", "c") |) **f** (| c("a", "b", "c") |)

map (| c("a", "b", "c") |, .f) map(| list() , .f) map (data.frame() , .t)

Your Turn 2

Read the code in the first chunk and predict what will happen.

Run the code in the first chunk. What does it return?

```
list(
  sum_a = sum(number_list$a),
  sum_b = sum(number_list$b),
  sum_c = sum(number_list$c)
)
```

Now, use map() to create the same output.

Your Turn 2

```
map(number_list, sum)

### $a

### [1] 14.6358

### $b

### [1] 3.77212

###

### $c

### [1] 181.8609
```

```
gapminder %>%
  dplyr::select_if(is.numeric) %>%
  map(sd)
```

```
gapminder %>%
  dplyr::select_if(is.numeric) %>%
  map(sd)
```

```
gapminder %>%
  dplyr::select_if(is.numeric) %>%
  map(sd)
```

```
gapminder %>%
   dplyr::select if(is.numeric) %>%
   map(sd)
排 $year
## [1] 17.26533
4‡4‡
## $lifeExp
## [1] 12.91711
4‡4‡
## $pop
## [1] 106157897
4‡4‡
## $gdpPercap
## [1] 9857.455
```

Your Turn 3

Pass diabetes to map() and map using class(). What are these results telling you?

Your Turn 3

```
head(
  map(diabetes, class),
  3
## $id
## [1] "numeric"
4F4F
排 $chol
## [1] "numeric"
4F4F
## $stab.glu
## [1] "numeric"
```

```
x <- x^2
x <- scale(x)
x <- max(x)</pre>
```

```
x <- x^2
x <- scale(x)
x <- max(x)

y <- x^2
y <- scale(y)
y <- max(y)

z <- z^2
z <- scale(x)
z <- max(z)</pre>
```

```
x <- x^2
x <- scale(x)
x <- max(x)

y <- x^2
y <- scale(y)
y <- max(y)

z <- z^2
z <- scale(x)
z <- max(z)</pre>
```

```
x <- x^3
x <- scale(x)
x <- max(x)

y <- x^2
y <- scale(y)
y <- max(y)

z <- z^2
z <- scale(x)
z <- max(z)</pre>
```

Review: writing functions

```
.f <- function(x) {
    x <- x^3
    x <- scale(x)

    max(x)
}
.f(x)
.f(y)
.f(y)</pre>
```

If you copy and paste your code three times, it's time to write a function

Write a function that returns the mean and standard deviation of a numeric vector.

Give the function a name

Find the mean and SD of x

Map your function to number_list

```
mean_sd <- function(x) {
   x_mean <- mean(x)
   x_sd <- sd(x)
   tibble(mean = x_mean, sd = x_sd)
}
map(number_list, mean_sd)</pre>
```

```
## $a
## # A tibble: 1 x 2
4‡4‡
  mean sd
## <dbl> <dbl>
## 1 1.46 1.39
##
## $b
## # A tibble: 1 x 2
##
  mean sd
## <dbl> <dbl>
## 1 0.754 0.783
##
## $c
## # A tibble: 1 x 2
##
  mean
           sd
## <dbl> <dbl>
## 1 9.09 2.77
```

Three ways to pass functions to map()

- pass directly to map()
- use an anonymous function
- 3 use ~

```
map(
   . X,
  mean,
             TRUE
  na.rm
```

```
map(
  . X,
  function(.x) {
    mean(.x,
    na.rm = TRUE)
```

```
map(
  . X,
  ~mean(.x,
       = TRUE
 na.rm
```

map(gapminder, ~length(unique(.x)))

```
map(gapminder, ~length(unique(.x)))
## $country
## [1] 142
##
## $continent
## [1] 5
##
## $year
## [1] 12
##
## $lifeExp
## [1] 1626
##
## $pop
## [1] 1704
##
## $gdpPercap
```

[1] 1704

Returning types

тар	returns	
map()	list	
map_chr()	character vector	
map_dbl()	double vector (numeric)	
map_int()	integer vector	
map_lgl()	logical vector	
map_dfc()	data frame (by column)	
map_dfr()	data frame (by row)	

Returning types

```
map_int(gapminder, ~length(unique(.x)))
```

Returning types

```
map_int(gapminder, ~length(unique(.x)))

### country continent year lifeExp pop gdpPercap
### 142 5 12 1626 1704 1704
```

Do the same as #3 above but return a vector instead of a list.

```
map_chr(diabetes, class)
```

```
chol
                               stab.glu
                                                hdl
                                                           ratio
                                                                       glyhb
4F4F
            id
     "numeric"
                 "numeric" "numeric"
                                          "numeric"
                                                       "numeric"
                                                                   "numeric"
##
41:41:
      location
                                 gender
                                             height
                                                         weight
                                                                       frame
                       age
## "character"
                 "numeric" "character"
                                          "numeric"
                                                       "numeric" "character"
##
         bp.1s
                     bp.1d
                                  bp.2s
                                              bp.2d
                                                           waist
                                                                         hip
     "numeric"
                 "numeric" "numeric"
                                          "numeric"
                                                       "numeric"
41:41:
                                                                   "numeric"
##
     time.ppn
     "numeric"
##
```

Check diabetes for any missing data.

Using the ~.f(.x) shorthand, check each column for any missing values using is.na() and any()

Return a logical vector. Are any columns missing data? What happens if you don't include any()? Why?

Try counting the number of missing, returning an integer vector

```
map_lgl(diabetes, ~any(is.na(.x)))
                                                    glyhb location
         id
                chol stab.glu
                                    hdl
                                           ratio
##
                                                                         age
##
      FALSE
                TRUE
                        FALSE
                                   TRUE
                                            TRUE
                                                     TRUE
                                                              FALSE
                                                                       FALSE
##
     gender
              height
                       weight
                                  frame
                                           bp.1s
                                                    bp.1d
                                                              bp.2s
                                                                       bp.2d
##
     FALSE
                TRUE
                         TRUE
                                   TRUE
                                            TRUE
                                                     TRUE
                                                              TRUE
                                                                        TRUE
4F4F
      waist
                 hip time.ppn
##
       TRUE
                TRUE
                         TRUE
```

```
map_int(diabetes, ~sum(is.na(.x)))
                                                 glyhb location
##
        id
               chol stab.glu
                                 hdl
                                        ratio
                                                                    age
                  1
                                   1
##
         0
                                                    13
                                                             0
                                                                      0
            height
                    weight
##
    gender
                               frame
                                        bp.1s
                                                 bp.1d
                                                        bp.2s
                                                                   bp.2d
##
                  5
                                   12
                                                            262
                                                                    262
4F4F
     waist
           hip time.ppn
##
```

Turn diabetes **into a list split by** location **using the** split() **function. Check its length.**

Fill in the model_lm function to model chol (the outcome) with ratio and pass the .data argument to lm()

map model_lm to diabetes_list so that it returns a data frame (by row).

```
diabetes_list <- split(diabetes, diabetes$location)
length(diabetes_list)
model_lm <- function(.data) {
   mdl <- lm(chol ~ ratio, data = .data)
   # get model statistics
   broom::glance(mdl)
}
map(diabetes_list, model_lm)</pre>
```

```
## [1] 2
## $Buckingham
## # A tibble: 1 x 11
## r.squared adj.r.squared sigma statistic p.value df logLik AIC BIC
              <dbl> <dbl> <dbl> <int> <dbl> <dbl> <dbl> <dbl> <dbl> <</pre>
        <fdbl>
4F4F
                   0.248 38.8 66.4 4.11e-14 2 -1009. 2025. 2034.
        0.252
## 1
## # ... with 2 more variables: deviance <dbl>, df.residual <int>
41:41:
排 $Louisa
## # A tibble: 1 x 11
## r.squared adj.r.squared sigma statistic p.value df logLik AIC BIC
4F4F
        <dbl>
              <dbl> <dbl> <dbl> <dbl> <int> <dbl> <dbl> <dbl> <dbl> <</pre>
                     0.201 39.4 51.7 1.26e-11 2 -1033. 2072. 2082.
## 1 0.204
## # ... with 2 more variables: deviance <dbl>, df.residual <int>
```

map2(.x, .y, .f)

.x, .y: a vector, list, or data frame

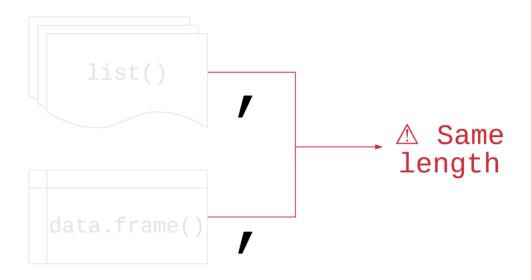
.f: a function

Returns a list

map2(

```
list()
data.frame()
```

map2(



.f

map2(

```
data.frame()
~.f(.x,.y)
```

map2()

```
means <-c(-3, 4, 2, 2.3)

sds <-c(.3, 4, 2, 1)

map2_dbl(means, sds, rnorm, n = 1)
```

map2()

```
means <-c(-3, 4, 2, 2.3)

sds <-c(.3, 4, 2, 1)

map2_dbl(means, sds, rnorm, n = 1)
```

map2()

```
means <-c(-3, 4, 2, 2.3)

sds <-c(.3, 4, 2, 1)

map2_dbl(means, sds, rnorm, n = 1)

## [1] -2.997932 2.178125 1.266952 2.948287
```

Split the gapminder dataset into a list by country

Create a list of models using map(). For the first argument, pass gapminder_countries. For the second, use the ~.f() notation to write a model with lm(). Use lifeExp on the left hand side of the formula and year on the second. Pass .x to the data argument.

Use map2() to take the models list and the data set list and map them to predict(). Since we're not adding new arguments, you don't need to use ~.f().

```
gapminder_countries <- split(gapminder, gapminder$country)
models <- map(gapminder_countries, ~ lm(lifeExp ~ year, data = .x))
preds <- map2(models, gapminder_countries, predict)
head(preds, 3)</pre>
```

```
gapminder_countries <- split(gapminder, gapminder$country)
models <- map(gapminder_countries, ~ lm(lifeExp ~ year, data = .x))
preds <- map2(models, gapminder_countries, predict)
head(preds, 3)</pre>
```

```
gapminder_countries <- split(gapminder, gapminder$country)
models <- map(gapminder_countries, ~ lm(lifeExp ~ year, data = .x))
preds <- map2(models, gapminder_countries, predict)
head(preds, 3)</pre>
```

```
## $Afghanistan
## 1 2 3 4 5 6
## 29.90729 31.28394 32.66058 34.03722 35.41387 36.79051
## $Albania
## 1 2 3 4 5 6
## 59.22913 60.90254 62.57596 64.24938 65.92279 67.59621
## $Algeria
## $Algeria
## 43.37497 46.22137 49.06777 51.91417 54.76057 57.60697
```

input 1	input 2	returns
map()	map2()	list
map_chr()	map2_chr()	character vector
map_dbl()	map2_dbl()	double vector (numeric)
map_int()	map2_int()	integer vector
map_lgl()	map2_lgl()	logical vector
map_dfc()	map2_dfc()	data frame (by column)
map_dfr()	map2_dfr()	data frame (by row)

Other mapping functions

pmap() and friends: take n lists or data frame with argument names

Other mapping functions

pmap() and friends: take n lists or data frame with argument names

walk() and friends: for side effects like plotting; returns input invisibly

Other mapping functions

pmap() and friends: take n lists or data frame with argument names

walk() and friends: for side effects like plotting; returns input invisibly

imap() and friends: includes counter i

Other mapping functions

pmap() and friends: take n lists or data frame with argument names

walk() and friends: for side effects like plotting; returns input invisibly

imap() and friends: includes counter i

map_if(), map_at(): Apply only to certain
elements

input 1	input 2	input n	returns
map()	map2()	pmap()	list
map_chr()	map2_chr()	pmap_chr()	character vector
map_dbl()	map2_dbl()	pmap_dbl()	double vector (numeric)
map_int()	map2_int()	pmap_int()	integer vector
map_lgl()	map2_lgl()	pmap_lgl()	logical vector
map_dfc()	map2_dfc()	pmap_dfc()	data frame (by column)
map_dfr()	map2_dfr()	pmap_dfr()	data frame (by row)
walk()	walk2()	pwalk()	input (side effects!)

Your turn 9

Create a new directory using the fs package. Call it "figures".

Write a function to plot a line plot of a given variable in gapminder over time, faceted by continent. Then, save the plot (how do you save a ggplot?). For the file name, paste together the folder, name of the variable, and extension so it follows the pattern "folder/variable_name.png"

Create a character vector that has the three variables we'll plot: "lifeExp", "pop", and "gdpPercap".

Use walk() to save a plot for each of the variables

Your turn 9

```
fs::dir create("figures")
ggsave gapminder <- function(variable) {</pre>
  # we're using `aes string()` so we don't need the curly-curly syn
  p <- ggplot(</pre>
    gapminder,
    aes_string(x = "year", y = variable, color = "country")
  ) +
    geom line() +
    scale color manual(values = country colors) +
    facet wrap(vars(continent.)) +
    theme(legend.position = "none")
  ggsave(
    filename = paste0("figures", variable, ".png"),
    plot = p,
    dpi = 320
```

Your turn 9

```
vars <- c("lifeExp", "pop", "gdpPercap")
walk(vars, ggsave_gapminder)</pre>
```

Base R

base R	purrr		
lapply()	map()		
vapply()	map_*()		
sapply()	?		
x[] <- lapply()	map_dfc()		
mapply()	map2(), pmap()		

Benefits of purrr

- Consistent
- 2 Type-safe
- 3 ~f(.x)

```
x <- rnorm(10)
y <- map(x, mean)

x <- rnorm(10)
y <- vector("list", length(x))
for (i in seq_along(x)) {
   y[[i]] <- mean(x[[i]])
}</pre>
```

```
x <- rnorm(10)
y <- map(x, mean)

x <- rnorm(10)
y <- vector("list", length(x))
for (i in seq_along(x)) {
   y[[i]] <- mean(x[[i]])
}</pre>
```

```
x <- rnorm(10)
y <- map(x, mean)

x <- rnorm(10)
y <- vector("list", length(x))
for (i in seq_along(x)) {
   y[[i]] <- mean(x[[i]])
}</pre>
```

```
x <- rnorm(10)
y <- map(x, mean)

x <- rnorm(10)
y <- vector("list", length(x))
for (i in seq_along(x)) {
   y[[i]] <- mean(x[[i]])
}</pre>
```

Of course someone has to write loops. It doesn't have to be you.

--Jenny Bryan

Working with lists and nested data

Work with Lists

FILTER LISTS



pluck(.x, ..., .default=NULL)
Select an element by name
or index, pluck(x,"b") , or its
attribute with attr_getter.
pluck(x,"b",attr_getter("n"))



keep(.x, .p, ...) Select elements that pass a logical test. *keep(x, is.na)*



discard(.x, .p, ...) Select elements that do not pass a logical test. *discard*(x, is.na)



compact(.x, .p = identity)
Drop empty elements.
compact(x)



head_while(.x, .p, ...)
Return head elements
until one does not pass.
Also tail_while.
head_while(x, is.character)

RESHAPE LISTS



flatten(.x) Remove a level of indexes from a list. Also flatten_chr, flatten_dbl, flatten_dfc, flatten_dfr, flatten_int, flatten_lgl. flatten(x)



transpose(.l, .names = NULL) Transposes the index order in a multi-level list. transpose(x)

SUMMARISE LISTS



every(.x, .p, ...) Do all elements pass a test? every(x, is.character)



b ■ с ■ some(x, is.character)
has_element(.x, .y) Does a
list contain an element?
has element(x, "foo")

some(.x, .p, ...) Do some

elements pass a test?



detect(.x, .f, ..., .right=FALSE, .p) Find first element to pass. detect(x, is.character)



detect_index(.x, .f, ..., .right = FALSE, .p) Find index of first element to pass. detect_index(x, is.character)



vec_depth(x) Return depth (number of levels of indexes). *vec_depth(x)*

JOIN (TO) LISTS



append(x, values, after =
length(x)) Add to end of list.
append(x, list(d = 1))



prepend(x, values, before =
1) Add to start of list.
prepend(x, list(d = 1))



splice(...) Combine objects into a list, storing S3 objects as sub-lists. *splice(x, y, "foo")*

TRANSFORM LISTS



modify(.x, .f, ...) Apply function to each element. Also map, map_chr, map_dbl, map_dfc, map_int, map_lgl. $modify(x, \sim +2)$



modify_at(.x, .at, .f, ...) Apply function to elements by name or index. Also **map_at**. *modify_at(x, "b", ~.+ 2)*



modify_if(.x, .p, .f, ...) Apply function to elements that pass a test. Also **map_if**. *modify_if*(x, is.numeric, ~.+2)

modify_depth(.x,.depth,.f,...) Apply function to each element at a given level of a list. *modify_depth(x, 1, ~.+ 2)*

WORK WITH LISTS



array_tree(array, margin = NULL) Turn array into list. Also array_branch. array_tree(x, margin = 3)

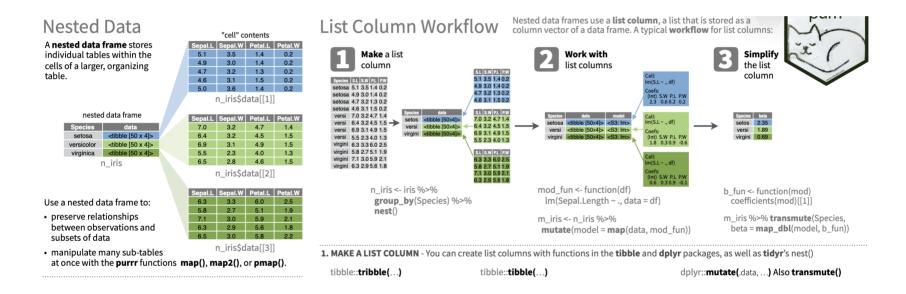


cross2(.x, .y, .filter = NULL) All combinations of .x and .y. Also cross, cross3, cross df. cross2(1:3, 4:6)



set_names(x, nm = x) Set the names of a vector/list directly or with a function. set_names(x, c("p", "q", "r")) set_names(x, tolower)

Working with lists and nested data



Adverbs: Modify function behavior

Modify function behavior

compose() Compose multiple functions.

lift() Change the type
of input a function
takes. Also lift_dl,
lift_dv, lift_ld, lift_lv,
lift_vd, lift_vl.

rerun() Rerun expression n times.

negate() Negate a predicate function (a pipe friendly!)

partial() Create a version of a function that has some args preset to values.

safely() Modify func to return list of results and errors. quietly() Modify function to return list of results, output, messages, warnings.

possibly() Modify function to return default value whenever an error occurs (instead of error).

Learn more!

Jenny Bryan's purrr tutorial: A detailed introduction to purrr. Free online.

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

Free online.