



CHARLOTTE WICKHAM

HAPPY R USERS PURRRR: USING FUNCTIONAL PROGRAMMING TO SOLVE ITERATION PROBLEMS

GETTING SETUP

1. Download .zip of slides and code @ bit.ly/purrr-rstudioconf
2. Extract and open purrr_workshop.Rproj
3. Check you have packages:

```
library(purrr)
library(tidyverse)
```

SOLVE ITERATION PROBLEMS

FOR EACH _____ DO _____

You are already solving them:

copy & paste, for loops, (1/s)apply()

I'll show you an alternative `purrr::map()` & friends

Download .zip of slides and code @ bit.ly/purrr-rstudioconf

FUNCTIONAL PROGRAMMING

a programming paradigm

has some central concepts

you don't need to know them to use purrr, but I'll point them out



Download .zip of slides and code @ bit.ly/purrr-rstudioconf

Star Wars API - <http://swapi.co/>

Data extracted using <https://github.com/Ironholds/rwars/>

```
# loads objects: films, people, vehicles, starships,  
# planets & species
```

```
load("data/swapi.rda")
```

1. How many elements are in people?
2. Who is the first person listed in people? What information is given for this person?
3. What is the difference between people[1] and people[[1]]?

BEWARE!
ANSWERS ON FOLLOWING SLIDE

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```
length(people)
## [1] 87

people[[1]]
## $name
## [1] "Luke Skywalker"
##
## $height
## [1] "172"
##
## $mass
## [1] "77"
##
## $hair_color
## [1] "blond"
##
## $skin_color
## [1] "fair"
##
## $eye_color
## [1] "blue"
##
## $birth_year
## [1] "19BBY"
##
## $gender
## [1] "male"
##
## $homeworld
## [1] "http://swapi.co/api/planets/1/"
##
```

```
## $films
## [1] "http://swapi.co/api/films/6/"
## [2] "http://swapi.co/api/films/3/"
## [3] "http://swapi.co/api/films/2/"
## [4] "http://swapi.co/api/films/1/"
## [5] "http://swapi.co/api/films/7/"
##
## $species
## [1] "http://swapi.co/api/species/1/"
##
## $vehicles
## [1] "http://swapi.co/api/vehicles/14/"
## [2] "http://swapi.co/api/vehicles/30/"
##
## $starships
## [1] "http://swapi.co/api/starships/12/"
## [2] "http://swapi.co/api/starships/22/"
##
## $created
## [1] "2014-12-09T13:50:51.644000Z"
##
## $edited
## [1] "2014-12-20T21:17:56.891000Z"
##
## $url
## [1] "http://swapi.co/api/people/1/"
```




map()


```
map( .x, .f, ... )
```

for each element of .x do .f

.x

- ▶ a vector
- ▶ a list
- ▶ a data frame (for each column)

.f

We'll get to that...

HOW MANY FILMS HAS EACH CHARACTER BEEN IN?

for each person in `people`, count the number of films

```
map(people, _____)
```

STRATEGY

1. Do it for one element
2. Turn it into a recipe
3. Use `map()` to do it for all elements

```
luke <- people[[1]]
```

HOW MANY STARSHIPS HAS LUKE BEEN IN?

Write a line of code to find out.

Bored? Find the names of those starships...

DO IT FOR ONE

Solve the problem for one element

```
luke <- people[[1]]
```

```
length(luke$starships)
```

DO IT FOR ONE

Solve the problem for one element

```
luke <- people[[1]]
```

```
length(luke$starships)
```

DO IT FOR ONE

Solve the problem for one element

```
leia <- people[[5]]
```

```
length(leia$starships)
```


DO IT FOR ONE

Solve the problem for one element

```
_____ <- people[[?]]
```

```
length(_____ $starships)
```

TURN IT INTO A RECIPE

Make it a formula

Use .x as a placeholder

~ length(**.x**\$starships)

A formula

purrr's placeholder for
one element of our vector

DO IT FOR ALL!

Your recipe is the second argument to map

```
map( people ,  
  ~ length( .x$starships) )
```

A formula

purrr's placeholder for
one element of our vector


```
map(people, ~ length(.x$starships))
```

Copy and paste ME.

Load then look at planet_lookup:

```
load("data/planet_lookup.rda")
```

```
planet_lookup
```

FIND THE NAME OF EACH CHARACTERS HOME WORLD.

Bored? Find the body mass index (BMI) of all characters.

$$\text{bmi} = (\text{mass in kg}) / ((\text{height in m})^2)$$

```
luke$homeworld
## [1] "http://swapi.co/api/planets/1/"

planet_lookup[luke$homeworld]
## http://swapi.co/api/planets/1/
## "Tatooine"

map(people, ~ planet_lookup[.x$homeworld])
## [[1]]
## http://swapi.co/api/planets/1/
## "Tatooine"

## [[2]]
## http://swapi.co/api/planets/1/
## "Tatooine"

## [[3]]
## http://swapi.co/api/planets/8/
## "Naboo"

...
```


ROAD_{map()}

map_lgl(.x, .f, ...)

Other types of output

Other ways of specifying .f

Other iteration functions

ROADmap()

map(.x, length, ...)

Other types of output

Other ways of specifying .f

Other iteration functions

ROADmap()

map2(.x, .y, .f, ...)

Other types of output

Other ways of specifying .f

Other iteration functions

map() details

`map()` **always** returns a list

SIMPLER OUTPUT:

`map_lgl()` **logical** vector

`map_int()` **integer** vector

`map_dbl()` **double** vector

`map_chr()` **character** vector

`walk()` - when you want nothing at all,
use a function for its side effects

Result: **No surprises!**

vector same length as `.x` or an ERROR


```
# names can be useful
people <- people %>% set_names(map_chr(people, "name"))
```

REPLACE `map()` WITH THE APPROPRIATELY TYPED FUNCTION

```
# How many starships has each character been in?
map(people, ~ length(.x[["starships"]]))
```

```
# What color is each character's hair?
map(people, ~ .x[["hair_color"]])
```

```
# Is the character male?
map(people, ~ .x[["gender"]] == "male")
```

```
# How heavy is each character?
map(people, ~ .x[["mass"]])
```

```
# How many starships has each character been in?
```

```
map_int(people, ~ length(.x[["starships"]]))
```

```
##      Luke Skywalker  C-3PO  R2-D2  Darth Vader  
##              2      0      0              1  ...
```

```
# What color is each character's hair?
```

```
map_chr(people, ~ .x[["hair_color"]])
```

```
##      Luke Skywalker  C-3PO  R2-D2  Darth Vader  
##      "blond"      "n/a"  "n/a"      "none"  ...
```

```
# Is the character male?
```

```
map_lgl(people, ~ .x[["gender"]] == "male")
```

```
##      Luke Skywalker  C-3PO  R2-D2  Darth Vader  
##              TRUE  FALSE  FALSE              TRUE  ...
```

```
# How heavy is each character?
```

```
map_dbl(people, ~ .x[["mass"]])
```

```
## Error: Can't coerce element 1 from a character to a double
```

```
# Doesn't work...because we get a string back
```

```
map(people, ~ .x[["mass"]])
```

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

```
## [1] "77"
```

```
##
```

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

```
## [1] "75"
```

```
...
```

```
# A little risky
```

```
map_dbl(people, ~ as.numeric(.x[["mass"]]))
```

```
## [1]  77.0   75.0   32.0  136.0  49.0  120.0   75.0   32.0   84.0
```

```
## ...
```

```
## There were 29 warnings (use warnings() to see them)
```

```
# Probably want something like:
```

```
map_chr(people, ~ .x[["mass"]]) %>%
```

```
  readr::parse_number(na = "unknown")
```

```
## [1]  77.0   75.0   32.0  136.0  49.0  120.0   75.0   32.0   84.0
```

```
## ...
```

. f CAN BE A FORMULA

`map(.x, .f = ~ DO SOMETHING WITH .x)`

```
map_int(people, ~ length(.x[["starships"]]))
```

```
map_chr(people, ~ .x[["hair_color"]])
```

```
map_chr(people, ~ .x[["mass"]])
```


.f CAN BE A STRING OR INTEGER

For each element, extract the named/numbered element

```
map(.x, .f = "some_name")
```

equivalent to

```
map(.x, ~ .x[[some_name]])
```

.f CAN BE A STRING OR INTEGER

For each element, extract the named/numbered element

```
map(.x, .f = some_number )
```

equivalent to

```
map(.x, ~ .x[[some_number]])
```

```
map_chr(people, ~ .x[["hair_color"]])
```

becomes

```
map_chr(people, "hair_color")
```

.f CAN BE A FUNCTION

`map(.x, .f = some_function, ...)`

equivalent to

`map(.x, ~ some_function(.x, ...))`

gets passed on to .f

```
char_starships <- map(people, "starships")  
map_int(char_starships, length)
```

```
# In one go  
map(people, "starships") %>% map_int(length)
```

```
# equivalent to  
map_int(people, ~ length(.x[["starships"]])
```

don't be afraid to do things in
little steps and pipe them
together

WHAT ABOUT `sapply()` & `lapply()`?

What type of object does `sapply()` return? It depends.

Motivation for `purrr`:

- consistent return type,
- useful shortcuts,
- consistent syntax for more complicated iteration

STAR WARS CHALLENGES

Which film (see `films`) has the most characters?

Create the `planet_lookup` vector from earlier.

Which species has the most possible eye colors?

```
# Which film (see films) has the most characters?
```

```
map(films, "characters") %>%
```

```
  map_int(length) %>%
```

```
  set_names(map_chr(films, "title")) %>%
```

```
  sort()
```

```
# Create the planet_lookup vector from earlier.
```

```
planet_lookup <- map_chr(planets, "name") %>%
```

```
  set_names(map(planets, "url"))
```



```
# Which species has the most possible eye colors?
```

```
species[[1]]$eye_colors
```

```
map_chr(species, "eye_colors") %>%
```

```
  strsplit(",", "") %>%
```

```
  map_int(length)
```

```
# this is lazy, what about n/a and unknown?
```

FUNCTIONS CAN BE ARGUMENTS

Functions are **first class** citizens in R, they can occur anywhere a number could:
as arguments,
as return values,
assigning them to variables,
storing in data structures.

Higher order function: a function that takes a function as input or returns a function

`map()` is a higher order function.

**purrr and
list columns**

PURRR AND LIST COLUMNS

Data should be in a data frame as soon as it makes sense!

Data frame: **cases** in rows, **variables** in columns

YOUR TURN:

What are the **cases** and **variables** in the people data?

```
# A tibble: 87 × 4
```

	name	films	height	species
	<chr>	<list>	<dbl>	<chr>
1	Luke Skywalker	<chr [5]>	172	http://swapi.co/api/species/1/
2	C-3P0	<chr [6]>	167	http://swapi.co/api/species/2/
3	R2-D2	<chr [7]>	96	http://swapi.co/api/species/2/
4	Darth Vader	<chr [4]>	202	http://swapi.co/api/species/1/
5	Leia Organa	<chr [5]>	150	http://swapi.co/api/species/1/

```
# ... with 82 more rows
```

PURRR CAN HELP TURN LISTS INTO TIBBLES

```
people_tbl <- tibble(  
  name      = c\("Han Solo", "Leia Organa"\),  
  films     = c\("The Force Awakens", "The Last Jedi"\),  
  height    = c\(180, 150\),  
  species   = c\("Human", "Human"\),  
)
```

Full code in [code/star_wars-tbl.R](#)

PURRR CAN HELP TURN LISTS INTO TIBBLES

```
people_tbl <- tibble(  
  name      = people %>% map_chr("name"),  
  films     = people %>% map("films"),      will result in list column  
  height    = people %>% map_chr("height") %>%  
    readr::parse_number(na = "unknown"),  needs some parsing  
  species   = people %>% map_chr("species", .null = NA_character_)  
)  
                                           isn't in every element
```

Full code in `code/star_wars-tbl.R`

COMBINE PURRR WITH DPLYR TO WORK WITH LIST COLUMNS

```
people_tbl$films
```

```
people_tbl %>%
```

```
  mutate(
```

```
    film_numbers = map(films, ~ film_number_lookup[.x]),
```

```
    n_films = map_int(films, length)
```

```
)
```

Code to create tibble in 04-purrr-list-columns.R

Create a new character column that collapses the film numbers into a single string,

e.g. for Luke: " 6, 3, 2, 1, 7"

```
people_tbl <- people_tbl %>%  
  mutate(  
    films_squashed = map_chr(film_numbers, paste,  
                             collapse = ", ")  
  )  
people_tbl %>% select(name, n_films, films_squashed)
```

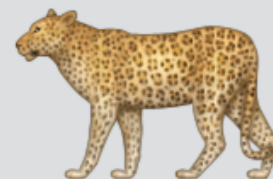
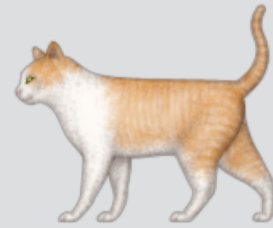
**More iteration
functions**

to each element of .x apply .f

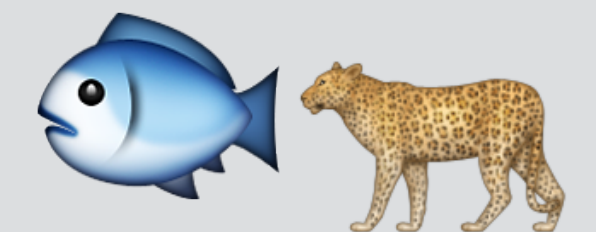
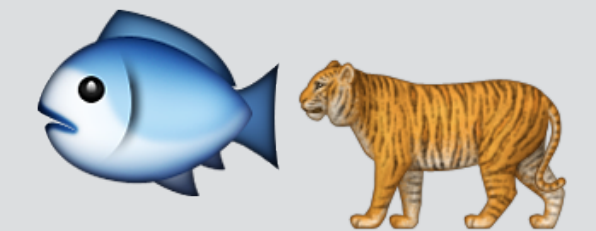
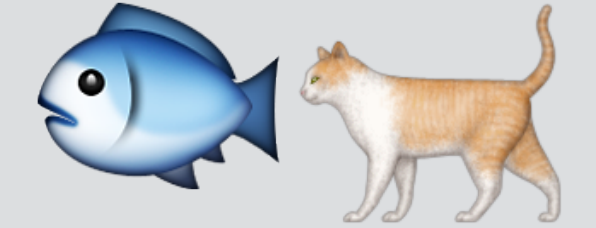
map(.x , .f)

to each `cat` apply `give_fish`

`map(`



`, give_fish)`




to each element of .x apply .f

`walk(.x , .f)`

Expect nothing in return

You actually get .x invisibly back,
good for piping

to each **cat** apply **love**

walk( , love)

Expect nothing in return

You actually get .x invisibly back,
good for piping

For functions called for their side effects:

- ▶ printing to screen
- ▶ plotting to graphics device
- ▶ file manipulation (saving, writing, moving etc.)
- ▶ system calls

to each element of `.x` and corresponding element of `.y` apply `.f`

```
map2( .x , .y , .f )
```

to each **cat** and corresponding **times** apply **rep**

map2(



,

3

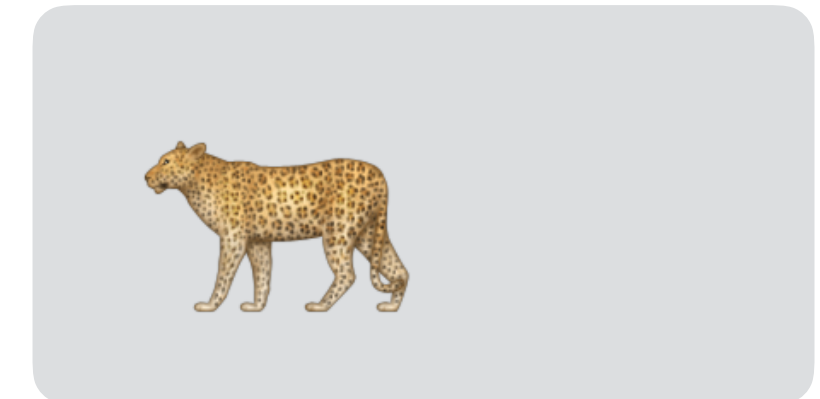
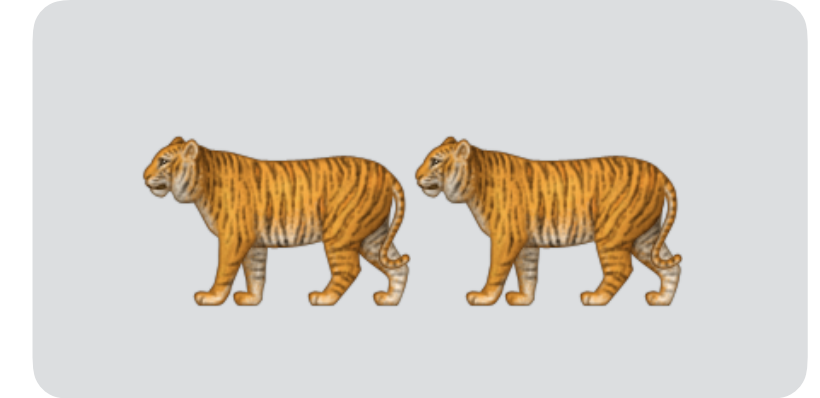
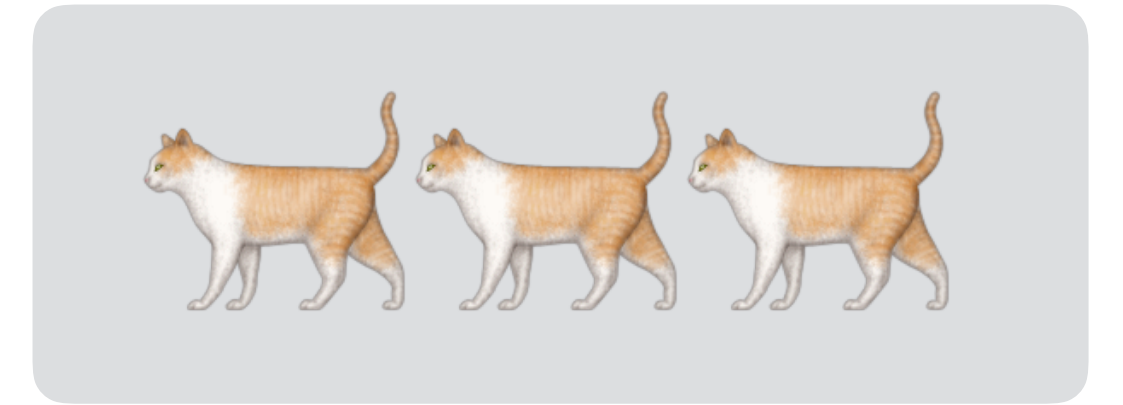
, rep)



2



1



Always get a list back, or use:

walk2(), map2_lgl(), map2_int(), map2_dbl(), map2_chr()

DISCUSS WITH YOUR NEIGHBOR

1. For each function, which two arguments might be useful to iterate over?

`download.file()`

`rnorm()`

`lm()`

`predict.lm()`

`write.csv()`

2. Which functions should we use `walk2()` or a typed version of `map2()`?

`download.file()` for each url download to destfile `walk2()`, `map2_int()`

`rnorm()` for each n generate a Normal sample with mean mean (or sd)

(See `purrr::rerun()` for repeating a function many times)

`lm()` for each data fit a model (formula)

`predict.lm()` for each model (object), generate predictions at data (newdata)

`readr::write_csv()` for each data frame (x) save to path `walk2()`

Similar for `ggplot::ggsave()` for each plot save to filename

NATIONAL ELECTRONIC INJURY SURVEILLANCE SYSTEM (NEISS)

From <https://github.com/hadley/neiss>

```
load("data/neiss_by_day.rda")
```

common_prods: 11 product codes with at least 50,000 injuries 2009-2014

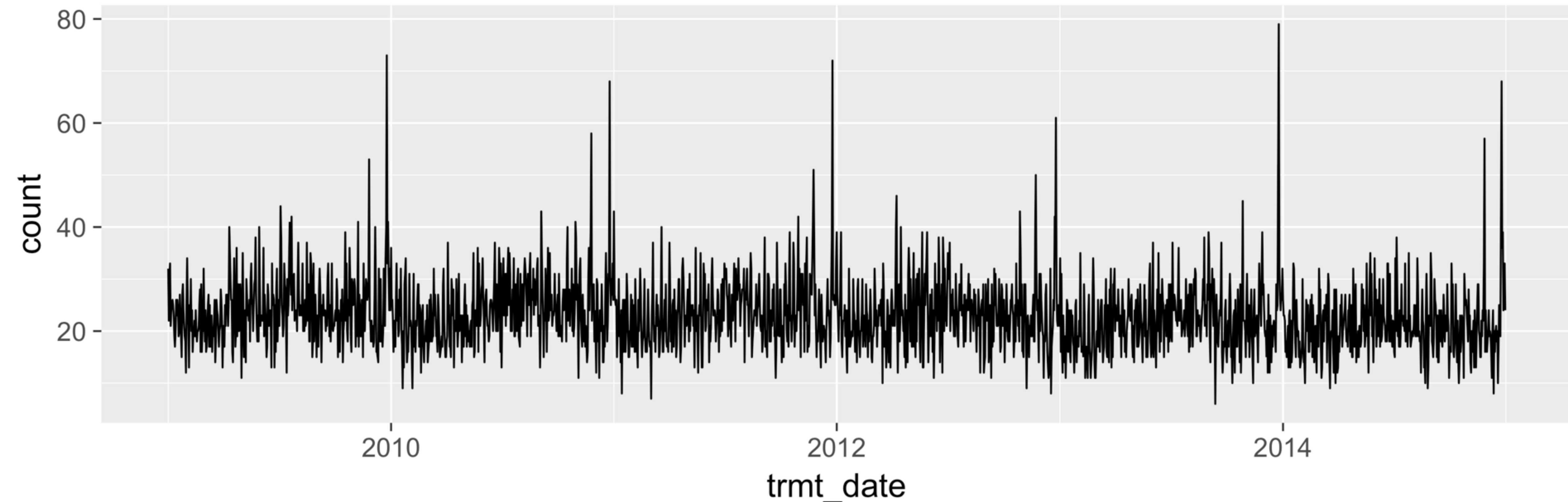
common_names: corresponding product description

per_day: a list with 11 elements, one for each product, injuries summarized to daily counts

ARE THERE PRODUCTS WITH PERIODIC PATTERNS IN INCIDENCE?

TAKING A LOOK

```
plots <- map(per_day, ~ ggplot(.x, aes(trmt_date, count)) + geom_line())  
plots[[1]] # try: walk(plots, print)
```



CAN WE SAVE THIS PLOT FOR ALL PRODUCTS?

DO IT FOR ONE

Solve the problem for one pair of elements

```
one_plot <- plots[[1]]
```

```
one_code <- common_codes[[1]]
```

```
ggsave(paste0(one_code, ".png"), one_plot)
```

TURN IT INTO A RECIPE

Make it a formula

Use .x and .y as
placeholders

```
~ ggsave(paste0(  .x  , ".png"),  .y  )
```

code plot

DO IT FOR ALL!

Your recipe is the .f argument to map2

```
walk2( common_codes, plots,  
  ~ ggsave(paste0( .x , ".png"), .y ))  
                        code      plot
```

WHEN THE SHORTCUT, ISN'T A SHORTCUT

```
walk2(paste0(common_codes, ".png"),  
      plots, ggsave)
```

WHEN THE SHORTCUT, ISN'T A SHORTCUT

```
walk2(paste0(common_codes, ".png"),  
      plots, ggsave,  
      width = 10, height = 3)
```

This fits a **naive** model with effects for month and day of the week to the first product:

```
lm(count ~ month + wday, data = per_day[[1]])
```

1. Fit the model to all products
2. Use `modelr::rsquare` to find the R-squared for each model (you'll probably want to look at `?modelr::rsquare`)
3. Bored? Repeat the plots but title them with the product names in `common_names`

```
models <- map(per_day, ~ lm(count ~ month + wday, data = .x))
```

```
map2_db1(models, per_day, modelr::rsquare)
```

```
plots[[3]]
```

```
common_names[[3]]
```

SHOULD REALLY BE USING LIST COLUMNS...

```
accidents <- tibble(  
  name = common_names,  
  code = common_codes,  
  data = per_day)
```

```
accidents %>%  
  mutate(  
    model = map(data, ~ lm(count ~ month + wday, data = .x)),  
    rsquare = map2_dbl(model, data, modelr::rsquare)) %>%  
  arrange(rsquare) %>%  
  select(name, rsquare)
```


CHALLENGES:

challenges/01-mtcars.R - Fit and summarise many regression models

challenges/02-word_count.R - Count the number of words of all files in a directory

challenges/03-starwars.R - Print who used which vehicles in the films

challenges/04-weather.R - Download, tidy, plot and save daily temperatures

challenges/05-swapi.R - Download all Star Wars data using rwars package

Next up: a few remaining iteration functions, a comment about other functions in purrr, wrap up.

to each element of each vector in `.l`, apply `.f`

`pmap(.l , .f , ...)`

```
.f(.l[[1]][[1]], .l[[2]][[1]], .l[[3]][[1]], ...)
```

```
.f(.l[[1]][[2]], .l[[2]][[2]], .l[[3]][[1]], ...)
```




```
.f(.l[[1]][[3]], .l[[2]][[3]], .l[[3]][[1]], ...)
```




and so on




or by name if supplied




to each element in animal, reaction, and animal2, apply c




pmap (data.frame (



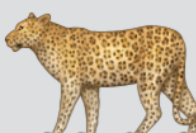












no more formula shortcut

for each function in `.f`, apply it to `.x`

`invoke_map(.f, .x, ...)`

`.f[[1]](.x, ...)`

`.f[[2]](.x, ...)`

`.f[[3]](.x, ...)`

and so on

for each function in .f, apply it to .x

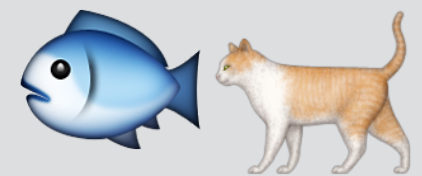
`invoke_map(`

`give_fish`

, )

`double`

`count_legs`



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OTHER FEATURES OF PURRR

06-other-features.R

LISTS AND FUNCTIONS

Key objects in purrr

`purrr` provides a pile of functions to make working with them easier

WITH YOUR NEIGHBOUR

Look at the help for `safely()` and `transpose()`

What kind of objects do they expect as input?

What kind of objects are returned as output?

SAFELY() TO HANDLE ERRORS

```
urls <- list(  
  example = "http://example.org",  
  asdf = "http://asdfasdasdkfjlda"  
)
```

```
map(urls, read_lines)
```

```
safe_readLines <- safely(readLines)  
safe_readlines
```

```
# Use the safe_readLines() function with map(): html  
html <- map(urls, safe_readLines)
```

TRANSPOSE() TO HANDLE RESULTS

```
# Easier to handle transposed  
str(html)  
str(transpose(html))  
  
# Extract the results: res  
res <- transpose(html)[["result"]]  
  
# Extract the errors: errs  
errs <- transpose(html)[["error"]]
```

WRAP UP

purrr provides:

- ▶ functions that write for loops for you
- ▶ with consistent syntax, and
- ▶ convenient shortcuts for specifying functions to iterate

Choosing the right function depends on:

- type of iteration
- type of output

Check out "Bonus" cheatsheet
in your conference packet

LEARNING MORE

R for Data Science:

- ▶ <http://r4ds.had.co.nz/iteration.html>
- ▶ <http://r4ds.had.co.nz/many-models.html>

DataCamp Writing functions in R

<https://www.datacamp.com/courses/writing-functions-in-r>

Jenny Bryan's purrr tutorial

<https://github.com/jennybc/purrr-tutorial>

THANK YOU

Slides and code @ bit.ly/purrr-rstudioconf

 @cvwickham

<http://cwick.co.nz>

cwickham@gmail.com

