



23 OCTOBER 2018

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01.72 - Kaat Tilley

```
> library(cowsay)
> say("Welcome to the coding club!", "turkey")
```

```
-----
Welcome to the coding club!
-----
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       `~)-) /          /` }
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```



LOOPS

Install the package suite:

```
install.packages("tidyverse")  
install.packages("rgbif")
```

Load the package suite:

```
library(tidyverse)  
library(rgbif)
```

Old skool...

For Loop

```
for (variable in sequence){  
  Do something  
}
```

Example

```
for (i in 1:4){  
  j <- i + 10  
  print(j)  
}
```

New kids:

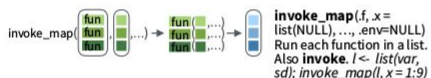
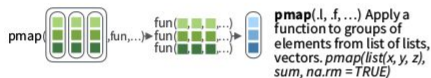
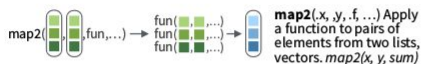
[purrr](#)

Apply functions with purrr : : CHEAT SHEET



Apply Functions

Map functions apply a function iteratively to each element of a list or vector.



lmap(x, f, ...) Apply function to each list-element of a list or vector.
imap(x, f, ...) Apply .f to each element of a list or vector and its index.

OUTPUT

map(), **map2()**, **pmap()**, **imap** and **invoke_map** each return a list. Use a suffixed version to return the results as a specific type of flat vector, e.g. **map2_chr**, **map2_dbl**, **map2_dfc**, **map2_dfr**, **map2_int**, **map2_lgl**, etc.

Use **walk**, **walk2**, and **pwalk** to trigger side effects. Each return its input invisibly.

function	returns
map	list
map_chr	character vector
map_dbl	double (numeric) vector
map_dfc	data frame (column bind)
map_dfr	data frame (row bind)
map_int	integer vector
map_lgl	logical vector
walk	triggers side effects, returns the input invisibly

SHORTCUTS - within a purrr function:

"name" becomes **function(x) x[["name"]]**, e.g. **map(l, "a")** extracts a from each element of l

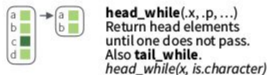
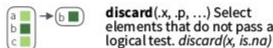
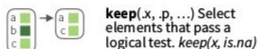
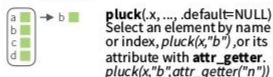
~ x becomes **function(x) x**, e.g. **map(l, ~ 2 * x)** becomes **map(l, function(x) 2 * x)**

~ .x .y becomes **function(x, y) x.y**, e.g. **map2(l, p, ~ x + y)** becomes **map2(l, p, function(l, p) l + p)**

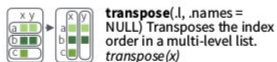
~ .1 ..2 etc becomes **function(.1, ..2, etc) ..1 ..2 etc**, e.g. **pmap(list(a, b, c), ~ .3 + ..1 - ..2)** becomes **pmap(list(a, b, c), function(a, b, c) c + a - b)**

Work with Lists

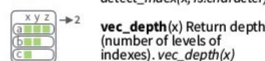
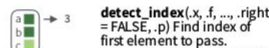
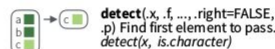
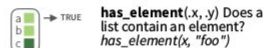
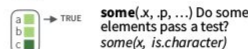
FILTER LISTS



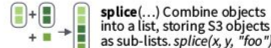
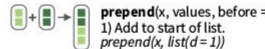
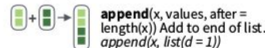
RESHAPE LISTS



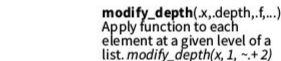
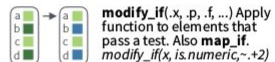
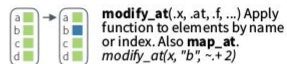
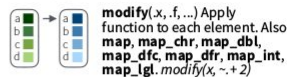
SUMMARISE LISTS



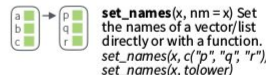
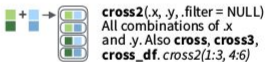
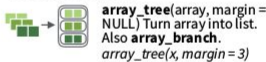
JOIN (TO) LISTS



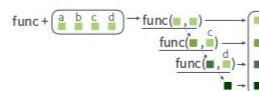
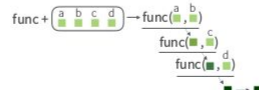
TRANSFORM LISTS



WORK WITH LISTS



Reduce Lists



Modify function behavior

compose() Compose multiple functions.

lift() Change the type of input a function takes. Also **lift_dbl**, **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 errors).


Share your snippets and solutions during the coding session:

Go to <https://hackmd.io/jwSucdiFQDCcIFsbHgLCCg> and post your code in between backticks:

For example:

```
```\n\nlibrary(tidyverse)\n\nmy_data <- ... \n\n```
```

# The concept

We defined a number of challenges. If you were able to achieve a challenge, add a  to your laptop screen.

The objective is that **everyone** achieves !

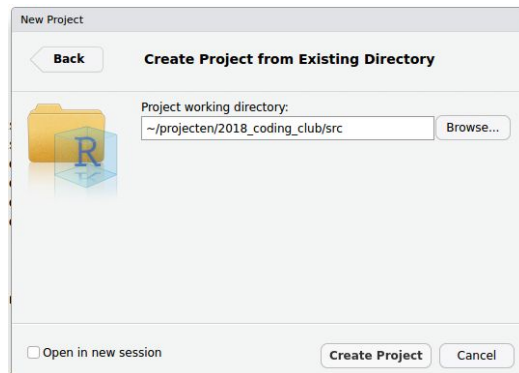
- Someone has more  than you? **Ask for help!**
- Someone has less  than you? **Provide help!**



- Download coding club material and work locally, **not in sync** with the Google drive



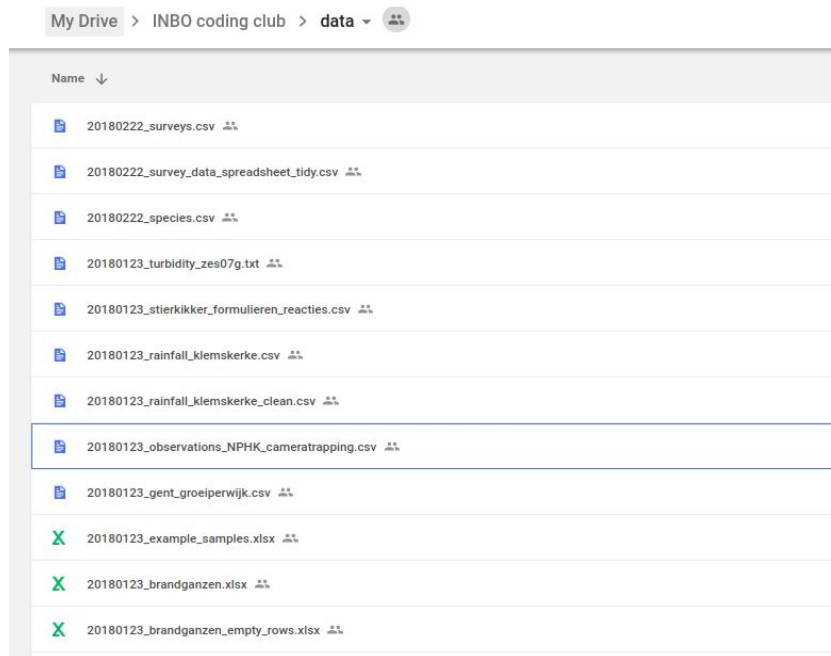
- Create new Rstudio project in the **/src** folder



- Download coding club material and work locally, not in sync with the Google drive
- Create new Rstudio project in the **src** folder...
- Use relative paths to data files!

```
> library(readr)
```

```
> read_csv2("../data/20180123_gent_groeiperwijk.csv")
```



For this coding club:

`20180522_gent_groeiperwijk_tidy.csv`

`20180222_species.csv`



```
for (variable in sequence){
 Do something
}
```

This code makes and saves a plot of the demographic evolution of Ghent's districts for year 2000.

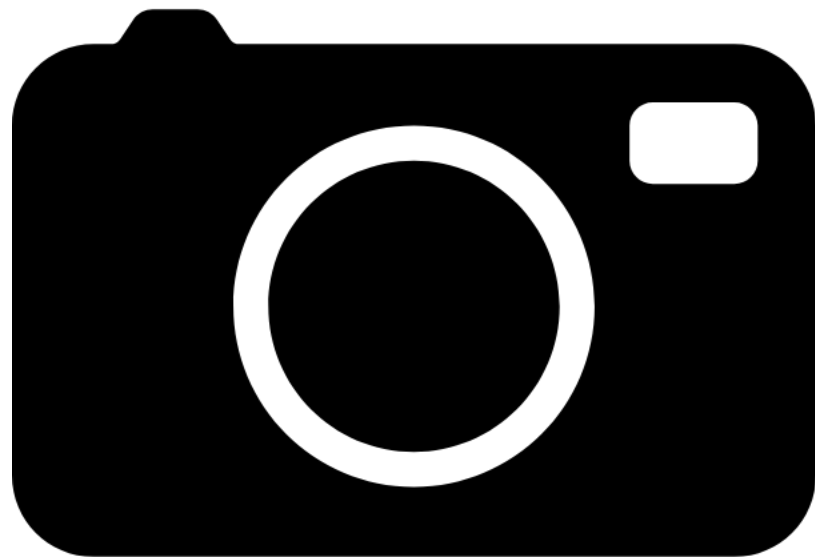
```
library(readr)

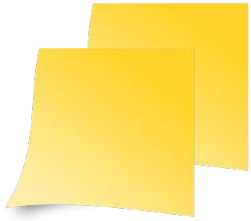
groei_gent_df <- read_csv("../data/20180522_gent_groeiperwijk_tidy.csv")

year_plot <- groei_gent_df %>% filter(year == 2000) %>%
 ggplot(aes(x = wijk, y = growth)) + geom_bar(stat = "identity") +
 coord_flip()

ggsave(file.path("../", "images", "district_evol_2000.png"), year_plot)
```

How to do the same for all years from 2000 to 2003 using a `for` loop?





Apply function `name_backbone()` to a vector/list of taxa (animals in the example below).

`name_backbone()` : lookup names in the GBIF backbone taxonomy.

```
library(rgbif)
```

```
name_backbone("Branta", rank = "GENUS")
```

```
name_backbone("Sus", rank = "GENUS")
```


```
animals <- c("Branta", "Sus")
```

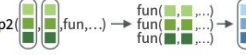
```
...
```


Can you get the result as a `data.frame` using `purrr` package as well?


## Apply Functions

Map functions apply a function iteratively to each element of a list or vector.

`map()`  **map(x, f, ...)** Apply a function to each element of a list or vector. *map(x, is.logical)*

`map2()`  **map2(x, y, f, ...)** Apply a function to pairs of elements from two lists, vectors. *map2(x, y, sum)*

`pmap()`  **pmap(l, f, ...)** Apply a function to groups of elements from list of lists, vectors. *pmap(list(x, y, z), sum, na.rm = TRUE)*

`invoke_map()`  **invoke\_map(f, x = list(NULL), ..., env=NULL)** Run each function in a list. Also **invoke**. *l <- list(var, sd); invoke\_map(l, x=1:9)*

**imap(x, f, ...)** Apply function to each list-element of a list or vector.  
**imap(x, f, ...)** Apply .f to each element of a list or vector and its index.

### OUTPUT

function	returns
<b>map()</b> , <b>map2()</b> , <b>pmap()</b> , <b>imap</b> and <b>invoke_map</b>	list
<b>map_chr</b>	character vector
<b>map_dbl</b>	double (numeric) vector
<b>map_dfc</b>	data frame (column bind)
<b>map_dfr</b>	data frame (row bind)
<b>map_int</b>	integer vector
<b>map_lgl</b>	logical vector
<b>walk</b>	triggers side effects, returns the input invisibly

Use **walk**, **walk2**, and **pwalk** to trigger side effects. Each return its input invisibly.

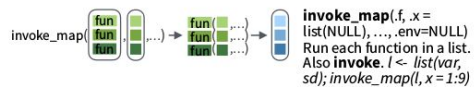
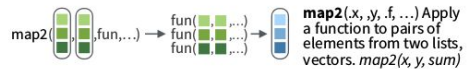


Apply function `name_backbone()` to all scientific names of the species data (genus + species) and add the results (multiple columns) to original data frame `df_species`

```
df_species <- read_csv("../data/20180222_species.csv")
df_species %>%
 mutate(scientific_name = str_c(genus, species,
 sep = " ")) %>%
 ...
```

## Apply Functions

Map functions apply a function iteratively to each element of a list or vector.



**lmap(x, f, ...)** Apply function to each list-element of a list or vector.  
**imap(x, f, ...)** Apply *f* to each element of a list or vector and its index.

### OUTPUT

function	returns
<b>map()</b> , <b>map2()</b> , <b>pmap()</b> , <b>imap</b> and <b>invoke_map</b>	list
<b>map_chr</b>	character vector
<b>map_dbl</b>	double (numeric) vector
<b>map_dfc</b>	data frame (column bind)
<b>map_dfr</b>	data frame (row bind)
<b>map_int</b>	integer vector
<b>map_lgl</b>	logical vector
<b>walk</b>	triggers side effects, returns the input invisibly

Use **walk**, **walk2**, and **pwalk** to trigger side effects. Each return its input invisibly.

```
A tibble: 55 x 27
 usageKey scientificName canonicalName rank status confidence matchType kingdom phylum order family genus species
 <int> <chr> <chr> <chr> <chr> <int> <chr> <chr> <chr> <chr> <chr> <chr>
1 2491757 Amphispizza bi... Amphispizza b... SPEC... ACCEP... 98 EXACT Animal... Chord... Pass... Ember... Amph... Amphis...
2 2437568 Ammospermophi... Ammospermoph... SPEC... ACCEP... 96 FUZZY Animal... Chord... Rode... Sciur... Ammo... Ammosp...
3 2491123 Ammodramus sa... Ammodramus s... SPEC... ACCEP... 98 EXACT Animal... Chord... Pass... Ember... Ammo... Ammodr...
...
8 2444480 Crotalus scut... Crotalus scu... SPEC... ACCEP... 92 FUZZY Animal... Chord... Squa... Viper... Crot... Crotal...
9 8071886 Cnemidophorus... Cnemidophoru... SPEC... SYNON... 97 EXACT Animal... Chord... Squa... Teiid... Aspi... Aspido...
10 5227544 Cnemidophorus... Cnemidophoru... SPEC... SYNON... 98 EXACT Animal... Chord... Squa... Teiid... Aspi... Aspido...
.. with 45 more rows, and 14 more variables: kingdomKey<int>, phylumKey <int>, classKey <int>, orderKey <int>,
familyKey <int>, genusKey <int>, speciesKey <int>, synonym <lgl>, class <chr>, acceptedUsageKey <int>,
species_id <chr>, genus <chr>, species1 <chr>, taxa <chr>
```



Go to <https://hackmd.io/jwSucdiFQDCcIFsbHgLCCg...>



Zaal: Herman Teirlinck - 01.71 - Frans Breziers

Datum: 2018-11-29, van 10:00 tot 12:00

*(registration announced via [DG\\_useR@inbo.be](mailto:DG_useR@inbo.be))*