

Making it **easier** to be **tidy**

Data Wrangling: Session 8

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Statistical Horizons, October 2023

Load the packages, as always














```
library(here)      # manage file paths
library(socviz)    # data and some useful functions
library(tidyverse) # your friend and mine
library(haven)     # for Stata, SAS, and SPSS files

library(broom)     # tidy model summaries
```

Moving ahead

Some helpful things

The RStudio Community

Topic		Replies	Views	Activity
  <input type="checkbox"/> Welcome to the RStudio Community! meta Welcome to community.rstudio.com — we're glad to have you! This welcome page will give you some advice on how to get the most out of the site if you're getting or giving help. We want this to be a friendly, inclusive com... read more		0	8.8k	2018-07-22
<input type="checkbox"/> what does that mean %in% in this code? • General	  	2	72	19m
<input type="checkbox"/> "William" in the Northeast ggplot Help • tidyverse ggplot2	  	5	69	28m
<input type="checkbox"/> Error while running RSelenium • shiny	 	4	27	28m
<input type="checkbox"/> Too slow to repsond •		0	16	40m
<input type="checkbox"/> Relative risk (RR) using categorical variables •		0	18	1h

The **reprex** package



Reference Articles ▾ News ▾ 

Overview

Prepare reprexes for posting to [GitHub issues](#), [StackOverflow](#), in Slack [messages](#) or [snippets](#), or even to paste into PowerPoint or Keynote slides. What is a `reprex`? It's a **re**producible **ex**ample, as coined by [Romain Francois](#).

Given R code on the clipboard, selected in RStudio, as an expression (quoted or not), or in a file ...

- run it via `rmarkdown::render()`,
- with deliberate choices re: `render()` arguments, knitr options, and Pandoc options.

Get resulting runnable code + output as

- Markdown, suitable for GitHub or Stack Overflow or Slack, or as
- R code, augmented with commented output, or as
- Plain HTML or (experimental) Rich Text

The result is returned invisibly, written to a file and, if possible, placed on the clipboard. Preview an HTML version in RStudio viewer or default browser.

Installation



Links

Download from CRAN at
<https://cloud.r-project.org/package=reprex>

Browse source code at
<https://github.com/tidyverse/reprex/>

Report a bug at
<https://github.com/tidyverse/reprex/issues>

License

[Full license](#)

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Community

[Contributing guide](#)

[Code of conduct](#)

Developers

Best demonstrated live

When asking for help, make a **reproducible example**

```
library(reprex)
```

```
library(tidyverse)
```

```
starwars %>%  
  count(homeworld, species) %>%  
  mutate(pct = n / sum(n) * 100) %>%  
  arrange(desc(pct))
```

```
## # A tibble: 58 × 4  
##   homeworld species      n    pct  
##   <chr>      <chr>   <int> <dbl>  
## 1 Tatooine   Human         8  9.20  
## 2 Naboo     Human         5  5.75  
## 3 <NA>      Human         5  5.75  
## 4 Alderaan  Human         3  3.45  
## 5 Naboo     Gungan        3  3.45  
## 6 <NA>      Droid         3  3.45  
## 7 Corellia  Human         2  2.30  
## 8 Coruscant Human         2  2.30  
## 9 Kamino    Kaminoan       2  2.30  
## 10 Kashyyyk Wookiee       2  2.30  
## # i 48 more rows
```

The **usethis** package

usethis **2.0.1.9000**



Setup

Reference

Articles ▾

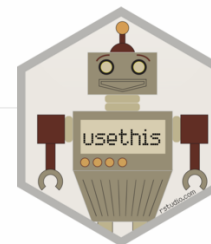
News ▾

Search...



usethis

usethis is a workflow package: it automates repetitive tasks that arise during project setup and development, both for R packages and non-package projects.



Installation

Install the released version of usethis from CRAN:

```
install.packages("usethis")
```

Or install the development version from GitHub with:

```
# install.packages("devtools")
devtools::install_github("r-lib/usethis")
```

Usage

Most `use_*()` functions operate on the *active project*: literally, a directory on your computer. If you've just used usethis to create a new package or project, that will be the active project. Otherwise, usethis verifies that current working directory is or is below a valid project directory and that becomes the active project. Use `proj_get()` or `proj_sitrep()` to manually query the project and [read more in the docs](#).

A few usethis functions have no strong connections to projects and will expect you to provide a path.

Links

Download from CRAN at
<https://cloud.r-project.org/package=usethis>

Browse source code at
<https://github.com/r-lib/usethis/>

Report a bug at
<https://github.com/r-lib/usethis/issues>

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Community

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[Code of conduct](#)

Developers

[Hadley Wickham](#)

Author

[Jennifer Bryan](#)

Author, maintainer

[Malcolm Barrett](#)

Author

The packages that made these slides



Never **paste** tables into a slide again!

```
---  
  
# Never .kjh-orange[copy and paste] code to a slide again!  
  
.pull-left[  
  
![:scale 100%](img/xaringan-sample.png)  
  
]  
  
.pull-right[  
  
```${r}  
Oh no, its the GSS
gss_sm %>%
 count(bigregion, religion)
```,  
  
]
```

religion	Northeast	Midwest	South	West
Protestant	158	325	650	238
Catholic	162	172	160	155
Jewish	27	3	11	10
None	112	157	170	180
Other	28	33	50	48
NA	1	5	11	1

Tables, tables, tables

The **gtsummary** package is very powerful. There are a number of other very good tidy table-making options too.

```
library(gtsummary)
```

```
trial
```

```
## # A tibble: 200 × 8
##   trt      age marker stage grade response death ttdeath
##   <chr>  <dbl>  <dbl> <fct> <fct>    <int> <int>    <dbl>
## 1 Drug A    23  0.16  T1    II         0     0      24
## 2 Drug B     9  1.11  T2     I         1     0      24
## 3 Drug A    31  0.277 T1    II         0     0      24
## 4 Drug A    NA  2.07  T3   III         1     1     17.6
## 5 Drug A    51  2.77  T4   III         1     1     16.4
## 6 Drug B    39  0.613 T4     I         0     1     15.6
## 7 Drug A    37  0.354 T1    II         0     0      24
## 8 Drug A    32  1.74  T1     I         0     1     18.4
## 9 Drug A    31  0.144 T1    II         0     0      24
## 10 Drug B   34  0.205 T3     I         0     1     10.5
## # i 190 more rows
```

Tables, tables, tables

The **gtsummary** package is very powerful. There are a number of other very good tidy table-making options too.

```
trial >
  tbl_summary(
    include = c(age, grade, response),
    by = trt, # split table by group
    missing = "no" # don't list missing data separately
  ) >
add_n() > # add column with total number of non-missing observations
add_p()
```

Tables, tables, tables

Characteristic	N	¹ ">Drug A, N = 98 ¹	¹ ">Drug B, N = 102 ¹	² ">p-value ²
Age	189	46 (37, 59)	48 (39, 56)	0.7
Grade	200			0.9
I		35 (36%)	33 (32%)	
II		32 (33%)	36 (35%)	
III		31 (32%)	33 (32%)	
Tumor Response	193	28 (29%)	33 (34%)	0.5
¹ Median (IQR); n (%)				
² Wilcoxon rank sum test; Pearson's Chi-squared test				

Tables, tables, tables

gtsummary() straight out of the box:

```
gss_sm >
  select(race, degree, marital) >
  drop_na() >
  tbl_summary(
    by = race, # split table by group
    missing = "no" # don't list missing data separately
  ) >
  bold_labels()
```

Tables, tables, tables

Characteristic	¹ n > White , N = 2,096 ¹	¹ n > Black , N = 487 ¹	¹ n > Other , N = 276 ¹
degree			
Lt High School	197 (9.4%)	60 (12%)	71 (26%)
High School	1,057 (50%)	292 (60%)	112 (41%)
Junior College	166 (7.9%)	33 (6.8%)	17 (6.2%)
Bachelor	426 (20%)	71 (15%)	39 (14%)
Graduate	250 (12%)	31 (6.4%)	37 (13%)
marital			
Married	979 (47%)	121 (25%)	110 (40%)
Widowed	196 (9.4%)	35 (7.2%)	18 (6.5%)
Divorced	363 (17%)	93 (19%)	39 (14%)
Separated	55 (2.6%)	27 (5.5%)	20 (7.2%)
Never Married	503 (24%)	211 (43%)	89 (32%)
¹ n (%)			

Tables, tables, tables

With a bit more work ...

```
trial ▷  
  select(trt, age, marker) ▷  
  tbl_summary(  
    by = trt,  
    type = all_continuous() ~ "continuous2",  
    statistic = all_continuous() ~ c("{N_nonmiss}",  
                                     "{mean} ({sd})",  
                                     "{median} ({p25}, {p75})",  
                                     "{min}, {max}"),  
    missing = "no"  
  ) ▷  
  italicize_levels()
```


Tables, tables, tables

Characteristic	Drug A, N = 98	Drug B, N = 102
Age		
<i>N</i>	91	98
<i>Mean (SD)</i>	47 (15)	47 (14)
<i>Median (IQR)</i>	46 (37, 59)	48 (39, 56)
<i>Range</i>	6, 78	9, 83
Marker Level (ng/mL)		
<i>N</i>	92	98
<i>Mean (SD)</i>	1.02 (0.89)	0.82 (0.83)
<i>Median (IQR)</i>	0.84 (0.24, 1.57)	0.52 (0.19, 1.20)
<i>Range</i>	0.00, 3.87	0.01, 3.64

The power of lists

Similar to earlier, but simpler:

```
library(gapminder)

## Fit as a function, for clarity
fit_ols <- function(df) {
  lm(lifeExp ~ log(gdpPercap), data = df)
}

out_le <- gapminder >
  filter(continent %in% "Oceania") >
  group_by(continent) >
  nest() >
  mutate(model = map(data, fit_ols),
         mod_sum = map(model, glance),
         mod_terms = map(model, tidy, conf.int = TRUE),
         ) >
  unnest(cols = c(mod_terms))
```

The power of lists

```
out_le
```

```
## # A tibble: 8 × 11
## # Groups:   continent [4]
##   continent data      model mod_sum term estimate std.error statistic p.value
##   <fct>      <list>  <lis> <list> <chr>    <dbl>    <dbl>    <dbl>    <dbl>
## 1 Asia      <tibble> <lm>  <tibble> (Int...  9.58     2.71      3.54 4.46e- 4
## 2 Asia      <tibble> <lm>  <tibble> log(...  6.25     0.331    18.9 3.73e- 57
## 3 Europe    <tibble> <lm>  <tibble> (Int... 13.0      1.92      6.76 5.52e- 11
## 4 Europe    <tibble> <lm>  <tibble> log(...  6.31     0.205    30.8 8.06e-103
## 5 Africa    <tibble> <lm>  <tibble> (Int...  7.60     2.63      2.89 4.03e- 3
## 6 Africa    <tibble> <lm>  <tibble> log(...  5.69     0.361    15.8 1.86e- 47
## 7 Americas  <tibble> <lm>  <tibble> (Int... -19.1     4.82     -3.95 9.65e- 5
## 8 Americas  <tibble> <lm>  <tibble> log(...  9.72     0.558    17.4 2.51e- 47
## # i 2 more variables: conf.low <dbl>, conf.high <dbl>
```

The power of lists

```
## Nice formatting of the numbers
## There are many convenience packages
## like this; it's not too hard to write your own, either
# remotes::install_github("tjmahr/printy")

text_ready <- out_le >
  mutate(
    across(c(estimate, conf.low, conf.high),
           printy::fmt_fix_digits, 2),
    se = printy::fmt_fix_digits(std.error, 3),
    ci = glue::glue("[{conf.low}, {conf.high}]")
  ) >
  select(continent, term, estimate, se, ci)
```

The printy package is by [T.J. Mahr](#)

The power of lists

```
text_ready
```

```
## # A tibble: 8 × 5
## # Groups:   continent [4]
##   continent term      estimate se      ci
##   <fct>      <chr>      <chr>    <chr> <glue>
## 1 Asia      (Intercept)    9.58    2.706 [4.26, 14.90]
## 2 Asia      log(gdpPercap) 6.25    0.331 [5.60, 6.90]
## 3 Europe    (Intercept)   12.97    1.917 [9.19, 16.74]
## 4 Europe    log(gdpPercap) 6.31    0.205 [5.91, 6.71]
## 5 Africa    (Intercept)    7.60    2.632 [2.43, 12.77]
## 6 Africa    log(gdpPercap) 5.69    0.361 [4.98, 6.40]
## 7 Americas  (Intercept)  -19.07    4.824 [-28.56, -9.58]
## 8 Americas  log(gdpPercap) 9.72    0.558 [8.62, 10.82]
```

The power of **lists**

Now ...

```
stats <- text_ready ▷  
  mutate(term = janitor::make_clean_names(term)) ▷  
  printy::super_split(continent, term) # Thanks again, TJ Mahr
```

The power of lists

Why are we doing this?

```
stats
```

```
## $Africa
## $Africa$intercept
## # A tibble: 1 × 5
## # Groups:   continent [1]
##   continent term      estimate se      ci
##   <fct>      <chr>      <chr>   <chr> <glue>
## 1 Africa    intercept  7.60    2.632 [2.43, 12.77]
##
## $Africa$log_gdp_percap
## # A tibble: 1 × 5
## # Groups:   continent [1]
##   continent term      estimate se      ci
##   <fct>      <chr>      <chr>   <chr> <glue>
## 1 Africa    log_gdp_percap 5.69    0.361 [4.98, 6.40]
##
##
## $Americas
## $Americas$intercept
## # A tibble: 1 × 5
## # Groups:   continent [1]
##   continent term      estimate se      ci
##   <fct>      <chr>      <chr>   <chr> <glue>
## 1 Americas  intercept -19.07   4.824 [-28.56, -9.58]
##
## $Americas$log_gdp_percap
## # A tibble: 1 × 5
```

The power of **lists**

The Intercept term for Africa was 'r stats\$Africa\$intercept\$estimate' 'r stats\$Africa\$intercept\$ci'.

For Europe it was 'r stats\$Europe\$intercept\$estimate' 'r stats\$Europe\$intercept\$ci'

The Intercept term for Africa was 7.60 [2.43, 12.77].

For Europe it was 12.97 [9.19, 16.74].

For more, see [this post by TJ Mahr](#):

<https://www.tjmahr.com/lists-knitr-secret-weapon/>

Test the output of your functions

```
countries ← read_csv(here("data", "countries.csv"))
```

```
countries
```

```
## # A tibble: 213 × 4
##   cname      iso3 iso2 continent
##   <chr>      <chr> <chr> <chr>
## 1 Afghanistan AFG  AF   Asia
## 2 Algeria      DZA  DZ   Africa
## 3 Armenia      ARM  AM   Asia
## 4 Australia    AUS  AU   Oceania
## 5 Austria      AUT  AT   Europe
## 6 Azerbaijan   AZE  AZ   Asia
## 7 Bahrain      BHR  BH   Asia
## 8 Belarus      BLR  BY   Europe
## 9 Belgium      BEL  BE   Europe
## 10 Brazil      BRA  BR   South America
## # i 203 more rows
```

Test the output of your functions

```
get_stmf <- function(url = "https://www.mortality.org/File/GetDocument/Public/STMF/Outputs",
  fname = "stmf",
  date = lubridate::today(),
  ext = "csv",
  dest = "data-raw/data",
  save_file = c("n", "y"),
  ... ) {
  save_file <- match.arg(save_file)
  target <- fs::path(url, fname, ext = ext)
  message("target: ", target)

  destination <- fs::path(here::here("data-raw/data"),
    paste0(fname, "_", date), ext = ext)

  tf <- tempfile(fileext = ext)
  curl::curl_download(target, tf)

  switch(save_file,
    y = fs::file_copy(tf, destination),
    n = NULL)

  janitor::clean_names(read_csv(tf, ... ))
}
```

Test the output of your functions

```
stmf_raw <- read_csv(here("data", "stmf.csv"), skip = 2) >
janitor::clean_names() >
rename(deaths_total = d_total, rate_total = r_total) >
select(country_code:sex, deaths_total, rate_total, split:forecast, everything()) >
pivot_longer(
  cols = d0_14:r85p,
  names_to = c("measure", "age_group"),
  names_pattern = "(r|d)(.*)"
) >
pivot_wider(names_from = measure,
            values_from = value) >
mutate(age_group = stringr::str_replace(age_group, "_", "-"),
       age_group = stringr::str_replace(age_group, "p", "+")) >
rename(death_count = d, death_rate = r) >
mutate(approx_date = paste0(year, "-", "W",
                           stringr::str_pad(week, width = 2, pad = "0"), "-", "7"),
       approx_date = ISOweek::ISOweek2date(approx_date)) >
select(country_code:sex, split:forecast, approx_date,
       age_group:death_rate, deaths_total, rate_total) >
mutate(country_code = replace(country_code, country_code = "AUS2", "AUS"),
       country_code = replace(country_code, country_code = "NZL_NP", "NZL"))
```

Test the output of your functions

```
stmf_raw
```

```
## # A tibble: 576,840 × 13
##   country_code year  week sex  split split_sex forecast approx_date age_group
##   <chr>         <dbl> <dbl> <chr> <dbl>      <dbl>      <dbl> <date>      <chr>
## 1 AUS          2015     1 m      1         0         0 2015-01-04 0-14
## 2 AUS          2015     1 m      1         0         0 2015-01-04 15-64
## 3 AUS          2015     1 m      1         0         0 2015-01-04 65-74
## 4 AUS          2015     1 m      1         0         0 2015-01-04 75-84
## 5 AUS          2015     1 m      1         0         0 2015-01-04 85+
## 6 AUS          2015     1 f      1         0         0 2015-01-04 0-14
## 7 AUS          2015     1 f      1         0         0 2015-01-04 15-64
## 8 AUS          2015     1 f      1         0         0 2015-01-04 65-74
## 9 AUS          2015     1 f      1         0         0 2015-01-04 75-84
## 10 AUS         2015     1 f      1         0         0 2015-01-04 85+
## # i 576,830 more rows
## # i 4 more variables: death_count <dbl>, death_rate <dbl>, deaths_total <dbl>,
## #   rate_total <dbl>
```

Test the output of your functions

```
md_ccodes <- tibble(country_code = unique(stmf_raw$country_code)) ▷
  left_join(countries, by = c("country_code" = "iso3")) ▷
  mutate(cname = replace(cname, country_code = "DEUTNP", "Germany"),
         iso2 = replace(iso2, country_code = "DEUTNP", "DE"),
         continent = replace(continent, country_code = "DEU", "Europe"),
         cname = replace(cname, country_code = "FRATNP", "France"),
         iso2 = replace(iso2, country_code = "FRATNP", "FR"),
         continent = replace(continent, country_code = "FRA", "Europe"),
         cname = replace(cname, country_code = "GBRTENW", "England and Wales"),
         cname = replace(cname, country_code = "GBR_SCO", "Scotland"),
         cname = replace(cname, country_code = "GBR_NIR", "Northern Ireland"),
         continent = replace(continent, country_code %in% c("GBRTENW", "GBR_SCO", "GBR_NIR"), "Europe")
  ) ▷
  left_join(countries)

stmf <- left_join(stmf_raw, md_ccodes) ▷
  select(country_code, cname:iso3, everything()) ▷
  mutate(iso3 = replace(iso3, iso2 = "DE", "DEU"),
         iso3 = replace(iso3, iso2 = "FR", "FRA"))
```

Test the output of your functions

```
stmf
```

```
## # A tibble: 576,840 × 17
##   country_code cname      iso2 continent iso3   year  week sex  split split_sex
##   <chr>         <chr>    <chr> <chr>    <chr> <dbl> <dbl> <chr> <dbl>    <dbl>
## 1 AUS          Austral... AU    Oceania AUS   2015     1 m     1      0
## 2 AUS          Austral... AU    Oceania AUS   2015     1 m     1      0
## 3 AUS          Austral... AU    Oceania AUS   2015     1 m     1      0
## 4 AUS          Austral... AU    Oceania AUS   2015     1 m     1      0
## 5 AUS          Austral... AU    Oceania AUS   2015     1 m     1      0
## 6 AUS          Austral... AU    Oceania AUS   2015     1 f     1      0
## 7 AUS          Austral... AU    Oceania AUS   2015     1 f     1      0
## 8 AUS          Austral... AU    Oceania AUS   2015     1 f     1      0
## 9 AUS          Austral... AU    Oceania AUS   2015     1 f     1      0
## 10 AUS         Austral... AU    Oceania AUS   2015     1 f     1      0
## # i 576,830 more rows
## # i 7 more variables: forecast <dbl>, approx_date <date>, age_group <chr>,
## #   death_count <dbl>, death_rate <dbl>, deaths_total <dbl>, rate_total <dbl>
```

For example, manually

```
## stmf
stmf_colnames ← c("country_code", "cname", "iso2", "continent", "iso3", "year",
                  "week", "sex", "split", "split_sex", "forecast", "approx_date",
                  "age_group", "death_count", "death_rate", "deaths_total", "rate_total")

all.equal(colnames(stmf), stmf_colnames)
```

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

For example, manually

```
## stmf
stmf_colnames ← c("country_code", "cname", "iso2", "continent", "iso3", "year",
                  "week", "sex", "split", "split_sex", "forecast", "approx_date",
                  "age_group", "death_count", "death_rate", "deaths_total", "rate_total")

all.equal(colnames(stmf), stmf_colnames)

## [1] TRUE
```

Imagine how you might build up a set of tests and checks

For example, manually

```
## stmf
stmf_colnames ← c("country_code", "cname", "iso2", "continent", "iso3", "year",
                  "week", "sex", "split", "split_sex", "forecast", "approx_date",
                  "age_group", "death_count", "death_rate", "deaths_total", "rate_total")

all.equal(colnames(stmf), stmf_colnames)

## [1] TRUE
```

Imagine how you might build up a set of tests and checks

But you don't have to manage this manually

Use **testthat** to check things

```
## countries
test_that("countries conforms to spec", {
  countries_colnames ← c("cname", "iso3", "iso2", "continent")
  expect_equal(colnames(countries), countries_colnames)
})

## stmf
test_that("stmf conforms to spec", {
  stmf_colnames ← c("country_code", "cname", "iso2", "continent", "iso3", "year",
    "week", "sex", "split", "split_sex", "forecast", "approx_date",
    "age_group", "death_count", "death_rate", "deaths_total", "rate_total")
  expect_equal(colnames(stmf), stmf_colnames)
})
```

Use **testthat** to check things

```
testthat::test_dir(here("tests", "testthat"))
```

```
## ✓ | OK F W S | Context
```

```
##
```

```
## - | 0 | stmf
```

```
## - | 0 | Validating package data objects
```

```
## ✓ | 2 | Validating package data objects
```

```
##
```

```
## == Results ==
```

```
## [ FAIL 0 | WARN 0 | SKIP 0 | PASS 2 ]
```

testthat in practice

Oriented towards package development

Consider packaging your datasets! Benefits to documentation/codebooks etc

One-table example: **uscenpops**

More extensive: **covdata**

How R packages work: **Wickham & Bryan**

Summarizing your wrangling with **skimr**

We might want to make a **codebook** of our data

```
library(skimr)
```

Summarizing your wrangling with **skimr**

We might want to make a **codebook** of our data

```
library(skimr)  
organdata ← read_csv(here("data", "organdonation.csv"))
```

Summarizing your wrangling with **skimr**

```
organdata > skim(where(is.numeric)) > partition()
```

Variable type: numeric

skim_variable	n_missing	complete_rate	mean	sd	p0	p25	p50	p75	p100	hist
year	34	0.86	1996.50	3.46	1991.00	1993.75	1996.50	1999.25	2002.00	
donors	34	0.86	16.48	5.11	5.20	13.00	15.10	19.60	33.90	
pop	17	0.93	39921.29	62219.22	3514.00	6938.00	15531.00	57301.00	288369.00	
pop.dens	17	0.93	12.00	11.09	0.22	1.94	9.49	19.11	38.89	
gdp	17	0.93	22986.18	4665.92	12917.00	19546.00	22756.00	26180.00	36554.00	
gdp.lag	0	1.00	22574.92	4790.71	11434.00	19034.25	22158.00	25886.50	36554.00	
health	0	1.00	2073.75	733.59	791.00	1581.00	1956.00	2407.50	5665.00	
health.lag	0	1.00	1972.99	699.24	727.00	1542.00	1850.50	2290.25	5267.00	
pubhealth	21	0.91	6.19	0.92	4.30	5.50	6.00	6.90	8.80	
roads	17	0.93	113.04	36.33	58.21	83.46	111.22	139.57	232.48	
cerebvas	17	0.93	610.80	144.45	300.00	500.00	604.00	698.00	957.00	
assault	17	0.93	16.53	17.33	4.00	9.00	11.00	16.00	103.00	
external	17	0.93	450.06	118.19	258.00	367.00	421.00	534.00	853.00	
txp.pop	17	0.93	0.72	0.20	0.22	0.63	0.71	0.83	1.12	

Summarizing your wrangling with **skimr**

```
organdata > skim(!where(is.numeric)) > partition()
```

Variable type: character

skim_variable	n_missing	complete_rate	min	max	empty	n_unique	whitespace
country	0	1.00	5	14	0	17	0
world	14	0.94	6	11	0	3	0
opt	28	0.88	2	3	0	2	0
consent.law	0	1.00	8	8	0	2	0
consent.practice	0	1.00	8	8	0	2	0
consistent	0	1.00	2	3	0	2	0
ccode	0	1.00	2	4	0	17	0

Custom Summaries

```
stmf_country_years <- function(df = stmf) {  
  df ▷  
    dplyr::select(cname, year) ▷  
    dplyr::group_by(cname, year) ▷  
    dplyr::tally() ▷  
    dplyr::mutate(n = as.character(n),  
                 n = dplyr::recode(n, "0" = "-", .default = "Y")) ▷  
    dplyr::group_by(year, cname) ▷  
    dplyr::arrange(year) ▷  
    tidyr::pivot_wider(names_from = year, values_from = n) ▷  
    dplyr::mutate(dplyr::across(where(is.character), dplyr::recode, .missing = "-")) ▷  
    dplyr::arrange(cname)  
}
```

Custom Summaries

knitr::kable(stmf_country_years())

cname	1990	1991	1992	1993	1994	1995	1996	1997	1998	1999	2000	2001	2002	2003	2004	2005	2006	2007	2008	2009	2010	2011	2012	2013	2014	2015	2016	2017	2018	2019	2020
Australia	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y
Austria	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Belgium	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Bulgaria	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Canada	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Chile	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y
Croatia	-	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Czech Republic	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Denmark	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
England and Wales	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Estonia	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Finland	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
France	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Germany	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Greece	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y
Hungary	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Iceland	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y
Israel	-	-	-	-	-	-	-	-	-	-	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y

