### Cleaning dirty data with {janitor}

Workshop: I2DS Tools for Data Science

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## Agenda

- 1. Overview
- 2. Data cleaning
- 3. Data exploring
- 4. Resources
- 5. Summary

#### The New York Times

# For Big-Data Scientists, 'Janitor Work' Is Key Hurdle to Insights

The New York Times



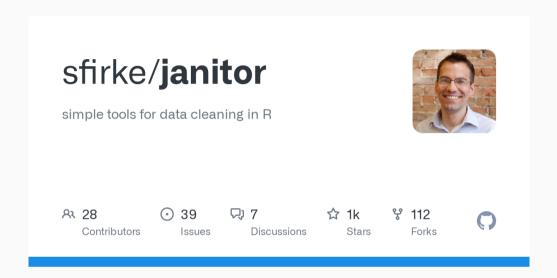
### Overview

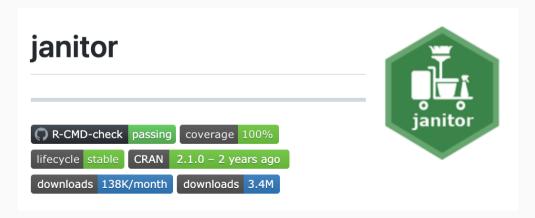
#### **(janitor)** is a package built by **Sam Firke**.

- Simple functions for cleaning and examining data
- Optimized for user-friendliness
- For beginning and intermediate R users
- Advanced R users: with {janitor} they can do data wrangling faster

#### Let's get started

- --> install.packages("janitor")
- --> library(janitor)





Clean data frame names with clean\_names()

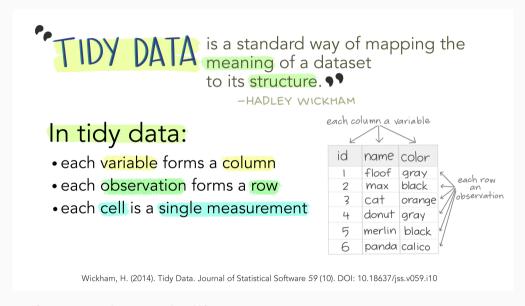
### tidyverse grammar

Consistent variable names are important

#### snake\_case

"Variable and function names should use only lowercase letters, numbers, and underscores. Use underscores (so called snake\_case) to separate words within a name."

The tidyverse style guide



Julie Lowndes and Allison Horst

#### Clean data frame names with clean\_names()

```
> # Create a test data frame with dirty variable names
> dirty_df ← as.data.frame(matrix(ncol = 7))
> names(dirty_df) ← c("firstName", "a@b!c'ß??",
                 "success.rate.in.%.(2022)",
+
                 "IDENTICAL", "IDENTICAL",
                 "", "#")
> clean_df ← dirty_df %>%
    clean_names()
> colnames(clean df)
## [1] "first name"
                                      "a_b_css"
## [3] "success_rate_in_percent_2022" "identical"
## [5] "identical 2"
## [7] "number"
```

#### Clean data frame names with clean\_names()

```
> # Create a test data frame with dirty variable names
> dirty df \leftarrow as.data.frame(matrix(ncol = 7))
> names(dirty_df) \leftarrow c("firstName", "a@b!c'\(\beta\)??",
                   "success.rate.in.%.(2022)".
 +
                   "IDENTICAL", "IDENTICAL",
> clean df ← dirty df %>%
     clean names()
 >
> colnames(clean df)
## [1] "first name"
                                         "a b css"
  [3] "success_rate_in_percent_2022" "identical"
```

```
[5] "identical 2"
                                       " x "
[7] "number"
```

- **Consistent format** for letter cases and separators
  - snake\_case is default, but other cases like camelCase are available
- Handles **special characters** and **spaces**, including transliterating characters like ß to ss
- Converts "%" to "percent" and "#" to "number"
- Adds numbers to duplicated names
- **Spacing** (or lack thereof) around numbers is preserved
- Works with the **%>% operator**
- Can be used for data frames and objects
- make names clean also works for character vectors

[10] "Culmen Length (mm)"

[16] "Delta 13 C (o/oo)"

[13] "Body Mass (g)"

```
> # install.packages("palmerpenguins")
> library(palmerpenguins)
> dirty_penguins_df ← penguins_raw
> colnames(dirty_penguins_df)

## [1] "studyName" "Sample Number" "Species"
## [4] "Region" "Island" "Stage"
## [7] "Individual ID" "Clutch Completion" "Date Egg"
```

"Flipper Length (mm)"

"Delta 15 N (o/oo)"

"Culmen Depth (mm)"

"Sex"

"Comments"

[16] "delta 13 c o oo"

"comments"

```
> # install.packages("palmerpenguins")
> library(palmerpenguins)
> dirty penguins df ← penguins raw
> colnames(dirty penguins df)
   [1] "studyName"
                              "Sample Number"
                                                     "Species"
   [4] "Region"
                              "Island"
                                                     "Stage"
   [7] "Individual ID"
                              "Clutch Completion"
                                                     "Date Egg"
  [10] "Culmen Length (mm)"
                             "Culmen Depth (mm)"
                                                     "Flipper Length (mm)"
  [13] "Body Mass (g)"
                                                     "Delta 15 N (o/oo)"
                              "Sex"
  [16] "Delta 13 C (o/oo)"
                              "Comments"
> clean penguins df ← penguins raw %>%
    clean names()
> colnames(clean penguins df)
    [1] "study name"
                            "sample number"
                                                 "species"
   [4] "region"
                            "island"
                                                 "stage"
##
   [7] "individual id"
                            "clutch completion" "date egg"
  [10] "culmen length mm"
                            "culmen_depth_mm"
                                                 "flipper_length_mm"
  [13] "body mass g"
                            "sex"
                                                 "delta 15 n o oo"
```

## Columns: 2

## \$ v1 <dbl> 1, 3

#### Remove content with remove\_empty()

- remove\_empty() rows and columns
  - Cleans Excel files that contain empty rows and columns after being read into R
  - Adding quiet = FALSE let's you know how many rows or columns were removed.

[7] "individual id"

[13] "body\_mass\_g"

[10] "culmen length mm"

[16] "delta 13 c o oo"

#### Drop constant columns with remove\_constant()

```
remove constant() columns
                                                                     remove constant()
Drops columns from a data frame that contain only a
                                                                 >
single constant value
                                                                 > adelie clean df %>%
                                                                     names()
> adelie df ← clean penguins df %>%
    filter(species = "Adelie Penguin (Pygoscelis adeli
                                                                    [1] "study_name"
                                                                    [4] "individual_id"
                                                                ##
> adelie df %>%
                                                                    [7] "culmen_length_mm"
    names()
                                                                ## [10] "body mass g"
                                                                ## [13] "delta 13 c o oo"
    [1] "study name"
                            "sample number"
                                                 "species"
    [4] "region"
                            "island"
                                                 "stage"
##
```

"clutch\_completion" "date\_egg"

"flipper length mm"

"delta 15 n o oo"

"culmen depth mm"

"sex"

"comments"

```
> adelie_clean_df ← adelie_df %>%
                            "sample number"
                                                "island"
                            "clutch_completion" "date_egg"
                            "culmen depth mm"
                                                "flipper le
                            "sex"
                                                "delta 15 n
                            "comments"
```

#### Check content of columns with compare\_df\_cols()

Imagine you have a set of data frames that you want to combine by binding the rows together but <code>rbind()</code> fails. You can check if the column classes match and see if they are **matching** by running <code>compare\_df\_cols()</code>.

- takes unquoted names of data frames, tibbles, or a list of data frames
- --> Returns a summary of how they compare
  - What column types are there?
  - How do column types differ?
  - Which are missing or present in the different inputs?

```
> chinstrap df ← clean penguins df %>%
     filter(species = "Chinstrap penguin (Pygoscelis ar
>
> # rbind(adelie. chinstrap)
>
> compare df cols(adelie clean df, chinstrap df) %>%
     tail()
##
        column name adelie clean df chinstrap df
             region
                               <NA>
                                       character
## 12
## 13 sample number
                            numeric
                                         numeric
## 14
                          character
                                       character
                sex
```

<NA>

<NA>

character

character

character

character

## 15

## 16

## 17

species

study name

stage

#### tabyl() – a tidy, fully-featured approach to counting things

Why not use table()?

- It doesn't work with the %>% operator
- It doesn't output data frames
- Its results are hard to format

#### One variable

```
> table(clean_penguins_df$sex)
```

```
## FEMALE MALE ## 165 168
```

#### tabyl() - a tidy, fully-featured approach to counting things

#### Why not use table()?

- It doesn't work with the %>% operator
- It doesn't output data frames
- Its results are hard to format

#### Instead better use tabyl()

- Tidyverse-aligned primarily built upon the {dplyr} and {tidyr} packages
- Compatible with the {knitr} package
- Useful for data exploration
- Generate frequencies along with the percent of total
- Counts combinations of one, two, or three variables

#### One variable

```
> table(clean penguins df$sex)
##
## FEMALE
           MALE
##
      165
            168
> tabyl(clean_penguins_df$sex)
   clean penguins df$sex n percent valid percent
                                             0.4954955
##
                   FFMALF 165 0.47965116
###
                     MALE 168 0.48837209
                                             0.5045045
                     <NA> 11 0.03197674
                                                    NA
##
> clean penguins no na df ← clean penguins df %>%
    drop na(sex)
```

Torgersen

24 23

#### tabyl() - a tidy, fully-featured approach to counting things

**Two variables** Two-way tabyl / "crosstab" or "contingency" table

```
> clean_penguins_no_na_df %>%
+ tabyl(island, sex)

## island FEMALE MALE
## Biscoe 80 83
## Dream 61 62
```

#### tabyl() - a tidy, fully-featured approach to counting things

**Two variables** Two-way tabyl / "crosstab" or "contingency" table

```
> penguins_crosstab_table 		 clean_penguins_no_na_df %>%
+ tabyl(island, sex) %>%
+ adorn_totals("col") %>%  # total in each row
+ adorn_percentages("row") %>%  # percentage value per row
+ adorn_pct_formatting(digits = 2) %>%  # rounded percentage value
+ adorn_ns() %>%  # adds the absolute numbers
+ adorn_title()  # adds the variable name
> penguins_crosstab_table
```

```
## sex

## island FEMALE MALE Total

## Biscoe 49.08% (80) 50.92% (83) 100.00% (163)

## Dream 49.59% (61) 50.41% (62) 100.00% (123)

## Torgersen 51.06% (24) 48.94% (23) 100.00% (47)
```

#### adorn\_\*() options

- tabyl() can be formatted with a suite of adorn\_\*
   functions to add information and for pretty formatting:
- adorn\_totals(): Add totals row, column, or both.
- **adorn\_percentages()**: Calculate percentages along either axis or the entire tabyl
- adorn\_pct\_formatting(): Format percentage columns, controlling the number of digits to display and whether to append the % symbol
- adorn\_rounding(): Round a data frame of numbers

- adorn\_ns(): Add Ns to a tabyl drawn from the tabyl's underlying counts or they can be supplied by the user
- adorn\_title(): Add a title to a tabyl pptions include putting the column title in a new row on top of the data frame or combining the row and column titles in the data frame's first name slot.

#### tabyl() Three variables Three-way tabyl

```
> clean_penguins_no_na_df %>% tabyl(island, species, sex) %>%
+ adorn_percentages("all") %>%
+ adorn_pct_formatting(digits = 1) %>%
+ adorn_title() %>%
+ kable()
```

	species				species		
island	Adelie Penguin (Pygoscelis adeliae)	Chinstrap penguin (Pygoscelis antarctica)	Gentoo penguin (Pygoscelis papua)	island	Adelie Penguin (Pygoscelis adeliae)	Chinstrap penguin (Pygoscelis antarctica)	Gentoo penguin (Pygoscelis papua)
Biscoe	13.3%	0.0%	35.2%	Biscoe	13.1%	0.0%	36.3%
Dream	16.4%	20.6%	0.0%	Dream	16.7%	20.2%	0.0%
Torgersen	14.5%	0.0%	0.0%	Torgersen	13.7%	0.0%	0.0%

### Fixing dates

excel\_numeric\_to\_date()
 Fix dates stored as serial numbers
 Converts serial date numbers from Excel to class Date

```
> excel_numeric_to_date(44886)
## [1] "2022-11-21"
```

convert\_to\_date()
 Convert a mix of date and datetime formats to date

```
> convert_to_date(c("2020-02-29", "40000.1"))
## [1] "2020-02-29" "2009-07-06"
> convert_to_datetime(40000.1)
## [1] "2009-07-06 02:24:00 UTC"
```

### Rounding numbers

## [1] 4 3

Careful: In base R round() uses **"banker's rounding"**, i.e., halves are rounded to the nearest *even* number.

```
> numbers \( \sigma c(3.5, 2.5) \)
> round(numbers)

## [1] 4 2

Instead use: round_half_up()
directionally-consistent rounding behavior

> round_half_up(numbers)
```

```
round_to_fraction()
round decimals to precise fractions of a given denominator
> round_to_fraction(0.175, denominator = 4)
## [1] 0.25
> round_to_fraction(0.2, denominator = 4)
## [1] 0.25
> round_to_fraction(0.25000000001, denominator = 4)
## [1] 0.25
```

#### **Detect duplicated records with** get\_dupes()

Checking for duplicated records is important because they can interfere with your tabulations in the analysis. { janitor } makes this tedious task simple.

```
get_dupes()
```

- Detects duplicate records during data cleaning
- Returns the records (and inserts a count of duplicates)
   so you can examine the potentially problematic cases

```
> clean_penguins_df %>%
+ get_dupes(individual_id) %>%
+ head(6)
```

```
## # A tibble: 6 × 18
     individual id dupe c...¹ study...² sampl...³ species region :
##
     <chr>
                       <int> <chr>
                                        <dbl> <chr> <chr> <
                                            21 Adelie... Anvers R
## 1 N11A1
                            2 PAL0708
## 2 N11A1
                            2 PAL0809
                                            47 Gentoo... Anvers I
                            2 PAL0708
                                            22 Adelie... Anvers I
## 3 N11A2
## 4 N11A2
                            2 PAL0809
                                            48 Gentoo... Anvers I
## 5 N12A1
                            2 PAL0708
                                            23 Adelie... Anvers I
## 6 N12A1
                            2 PAL0809
                                            49 Gentoo... Anvers I
## # ... with 9 more variables: date egg <date>, culmen_length
       culmen_depth_mm <dbl>, flipper_length_mm <dbl>, body
## #
       sex <chr>, delta 15 n o oo <dbl>, delta 13 c o oo <dl
## #
       comments <chr>, and abbreviated variable names ¹dupe
## #
## #
       <sup>2</sup>study name, <sup>3</sup>sample number, <sup>4</sup>clutch completion
                                                        20 / 22
```

### Resources

#### Original material:

- Package janitor documentation
- GitHub Repository

#### Further resources:

- exploringdata.org How to Clean Data: {janitor}
   Package
- towardsdatascience.com Cleaning and Exploring Data with the "janitor" Package
- jenrichmond.rbind.io Cleaning penguins with the janitor package

More on tidying data:

tidyr cheatsheet

Unfortunately, there is no cheatsheet for { janitor }. Here are two alternative tips:

- ls("package:janitor") gives you a list of all functions
- or type janitor:: in your console and you get to scroll up and down the list of all functions in the package.

BTW, this works for all packages ;)

### Summary

- Integrate {janitor} into your data wrangling and cleaning pipeline
- It works faster than regular functions from the {tidyverse}
- It can help you to:
  - **quickly** clean variable names
  - remove empty rows and columns
  - remove constant columns
  - easily create better tables that work in the tidyverse, can be stored as data frames and are almost worth publishing!



### **Happy data cleaning!**