

skimr

`skimr` provides a frictionless approach to summary statistics which conforms to the principle of least



surprise, displaying summary statistics the user can skim quickly to understand their data. It handles different data types and returns a `skim_df` object which can be included in a pipeline or displayed nicely for the human reader.

skimr

notes and vignettes carefully.

Installation

The current released version of `skimr` can be installed from CRAN. If you wish to install the current build of the next release you can do so using the following:

```
# install.packages("devtools")
devtools::install_github("ropensci/skimr")
```

The APIs for this branch should be considered reasonably stable but still subject to change if an issue is discovered.

To install the version with the most recent changes that have not yet been incorporated in the main branch (and may not be):

```
devtools::install_github("ropensci/skimr", ref = "develop")
```

Do not rely on APIs from the develop branch, as they are likely to change.

Skim statistics in the console

`skimr`:

- Provides a larger set of statistics than `summary()`, including missing, complete, n, and sd.
- reports each data types separately
- handles dates, logicals, and a variety of other types
- supports spark-bar and spark-line based on the [pillar package](#).

Separates variables by class:

```
skim(chickwts)
```

```
## — Data Summary —————
##                               Values
## Name                        chickwts
## Number of rows                71
```

skimr

```
## factor 1
## numeric 1
##
## Group variables None
##
## — Variable type: factor —
## skim_variable n_missing complete_rate ordered n_unique top_co
## 1 feed 0 1 FALSE 6 soy: 1
##
## — Variable type: numeric —
## skim_variable n_missing complete_rate mean sd p0 p25 p50
## 1 weight 0 1 261. 78.1 108 204. 258
```

Presentation is in a compact horizontal format:

```
skim(iris)

## — Data Summary —
##
## Name iris
## Number of rows 150
## Number of columns 5
##
## Column type frequency:
## factor 1
## numeric 4
##
## Group variables None
##
## — Variable type: factor —
## skim_variable n_missing complete_rate ordered n_unique top_co
## 1 Species 0 1 FALSE 3 set: 1
##
## — Variable type: numeric —
## skim_variable n_missing complete_rate mean sd p0 p25 p50
## 1 Sepal.Length 0 1 5.84 0.828 4.3 5.1 5.8
## 2 Sepal.Width 0 1 3.06 0.436 2 2.8 3
## 3 Petal.Length 0 1 3.76 1.77 1 1.6 4.3
## 4 Petal.Width 0 1 1.20 0.762 0.1 0.3 1.3
```

Built in support for strings, lists and other column

skimr

```
skim(dplyr::starwars)

## — Data Summary —————
##                               Values
## Name                         dplyr::starwars
## Number of rows                87
## Number of columns             14
##
## Column type frequency:
##   character      8
##   list           3
##   numeric        3
##
## Group variables              None
##
## — Variable type: character —————
##   skim_variable n_missing complete_rate min max empty n_unique
## 1 name          0          1      3 21    0      87
## 2 hair_color     5        0.943    4 13    0      11
## 3 skin_color     0          1      3 19    0      31
## 4 eye_color      0          1      3 13    0      15
## 5 sex            4        0.954    4 14    0       4
## 6 gender         4        0.954    8  9    0       2
## 7 homeworld     10        0.885    4 14    0      48
## 8 species        4        0.954    3 14    0      37
##
## — Variable type: list —————
##   skim_variable n_missing complete_rate n_unique min_length max
## 1 films          0          1      24      1
## 2 vehicles       0          1      11      0
## 3 starships      0          1      16      0
##
## — Variable type: numeric —————
##   skim_variable n_missing complete_rate mean    sd p0    p25 p
## 1 height        6          0.931 175.    34.8 66 167 18
## 2 mass          28          0.678  97.3 169.  15 55.6 2
## 3 birth_year    44          0.494  87.6 155.   8 35   5
```

Has a useful summary function

```
skim(iris) |>
  summary()
```

skimr

```
##                               Values
## Name                         iris
## Number of rows               150
## Number of columns            5
## _____
## Column type frequency:
##   factor                     1
##   numeric                    4
## _____
## Group variables              None
```

Individual columns can be selected using tidyverse-style selectors

```
skim(iris, Sepal.Length, Petal.Length)

## — Data Summary —
##                               Values
## Name                         iris
## Number of rows               150
## Number of columns            5
## _____
## Column type frequency:
##   numeric                    2
## _____
## Group variables              None
##
## — Variable type: numeric —
##   skim_variable n_missing complete_rate mean    sd  p0 p25  p50
## 1 Sepal.Length      0           1 5.84 0.828 4.3 5.1 5.8
## 2 Petal.Length      0           1 3.76 1.77  1  1.6 4.35
```

Handles grouped data

`skim()` can handle data that has been grouped using `dplyr::group_by()`.

```
iris |>
  dplyr::group_by(Species) |>
  skim()
```

```
## — Data Summary —
```

skimr

```
## Number of rows      150
## Number of columns    5
## _____
## Column type frequency:
##   numeric            4
## _____
## Group variables      Species
##
## — Variable type: numeric —
##   skim_variable Species   n_missing complete_rate  mean  sd
## 1 Sepal.Length  setosa      0           1 5.01 0.352
## 2 Sepal.Length  versicolor 0           1 5.94 0.516
## 3 Sepal.Length  virginica   0           1 6.59 0.636
## 4 Sepal.Width   setosa      0           1 3.43 0.379
## 5 Sepal.Width   versicolor 0           1 2.77 0.314
## 6 Sepal.Width   virginica   0           1 2.97 0.322
## 7 Petal.Length  setosa      0           1 1.46 0.174
## 8 Petal.Length  versicolor 0           1 4.26 0.476
## 9 Petal.Length  virginica   0           1 5.55 0.552
## 10 Petal.Width  setosa      0           1 0.246 0.105
## 11 Petal.Width  versicolor 0           1 1.33 0.198
## 12 Petal.Width  virginica   0           1 2.03 0.275
```

Behaves nicely in pipelines

```
iris |>
  skim() |>
  dplyr::filter(numeric.sd > 1)

## — Data Summary —
##                                     Values
## Name                               iris
## Number of rows                      150
## Number of columns                    5
## _____
## Column type frequency:
##   numeric                           1
## _____
## Group variables                      None
##
## — Variable type: numeric —
##   skim_variable n_missing complete_rate mean  sd p0 p25  p50 p
```

skimr

Knitted results

Simply skimming a data frame will produce the horizontal print layout shown above. We provide a `knit_print` method for the types of objects in this package so that similar results are produced in documents. To use this, make sure the `skimmed` object is the last item in your code chunk.

```
faithful |>
  skim()
```

| Name | faithful |
|------------------------|----------|
| Number of rows | 272 |
| Number of columns | 2 |
| Column type frequency: | |
| numeric | 2 |
| Group variables | None |

Data summary

Data summary

Variable type: numeric

| skim_variable | n_missing | complete_rate | mean | sd | p0 | p1 |
|---------------|-----------|---------------|-------|-------|------|------|
| eruptions | 0 | 1 | 3.49 | 1.14 | 1.6 | 2.0 |
| waiting | 0 | 1 | 70.90 | 13.59 | 43.0 | 58.0 |

Customizing skimr

skimr

create statistics for new classes and develop skimmers for data structures that are not data frames.

Specify your own statistics and classes

Users can specify their own statistics using a list combined with the `skim_with()` function factory. `skim_with()` returns a new `skim` function that can be called on your data. You can use this factory to produce summaries for any type of column within your data.

Assignment within a call to `skim_with()` relies on a helper function, `sfl` or `skimr` function list. By default, functions in the `sfl` call are appended to the default skimmers, and names are automatically generated as well.

```
my_skim <- skim_with(numeric = sfl(mad))
my_skim(iris, Sepal.Length)
```

But you can also helpers from the `tidyverse` to create new anonymous functions that set particular function arguments. The behavior is the same as in `purrr` or `dplyr`, with both `.` and `.x` as acceptable pronouns. Setting the `append = FALSE` argument uses only those functions that you've provided.

```
my_skim <- skim_with(
  numeric = sfl(
    iqr = IQR,
    p01 = ~ quantile(.x, probs = .01)
    p99 = ~ quantile(., probs = .99)
  ),
  append = FALSE
)
my_skim(iris, Sepal.Length)
```

And you can remove default skimmers by setting them to `NULL`.

```
my_skim <- skim_with(numeric = sfl(hist = NULL))
my_skim(iris, Sepal.Length)
```

Skimming other objects

`skimr` has summary functions for the following types of data by default:

skimr

- character
- factor
- logical
- complex
- Date
- POSIXct
- ts
- AsIs


`skimr` also provides a small API for writing packages that provide their own default summary functions for data types not covered above. It relies on R S3 methods for the `get_skimmers` function. This function should return a `sf1`, similar to customization within `skim_with()`, but you should also provide a value for the `class` argument. Here's an example.

```
get_skimmers.my_data_type <- function(column) {
  sf1(
    .class = "my_data_type",
    p99 = quantile(., probs = .99)
  )
}
```

Limitations of current version

We are aware that there are issues with rendering the inline histograms and line charts in various contexts, some of which are described below.

Support for spark histograms

With versions of R before 4.2.1, there are known issues with printing the spark-histogram characters when printing a data frame. For example, " " is printed as "<U+2582><U+2585><U+2587>". This longstanding problem [originates in the low-level code](#) for printing dataframes. While some cases have been addressed, there are, for example, reports of this issue in Emacs ESS. While this is a deep issue, there is [ongoing work to address it in base R](#). We recommend upgrading to at least R 4.2.1 to address this issue.

This means that while `skimr` can render the histograms to the console

skimr

- converting a `skimr` data frame to a vanilla R data frame, but tibbles render correctly
- in the context of rendering to a pdf using an engine that does not support utf-8.

One workaround for showing these characters in Windows is to set the CTYPE part of your locale to Chinese/Japanese/Korean with `Sys.setlocale("LC_CTYPE", "Chinese")`. The helper function `fix_windows_histograms()` does this for you.

And last but not least, we provide `skim_without_charts()` as a fallback. This makes it easy to still get summaries of your data, even if unicode issues continue.

Printing spark histograms and line graphs in knitted documents

Spark-bar and spark-line work in the console, but may not work when you knit them to a specific document format. The same session that produces a correctly rendered HTML document may produce an incorrectly rendered PDF, for example. This issue can generally be addressed by changing fonts to one with good building block (for histograms) and Braille support (for line graphs). For example, the open font “DejaVu Sans” from the `extrafont` package supports these. You may also want to try wrapping your results in `knitr::kable()`. Please see the vignette on using fonts for details.

Displays in documents of different types will vary. For example, one user found that the font “Yu Gothic UI Semilight” produced consistent results for Microsoft Word and Libre Office Write.

Inspirations

- [TextPlots](#) for use of Braille characters
- [spark](#) for use of block characters.

The earliest use of unicode characters to generate sparklines appears to be [from 2009](#).

Exercising these ideas to their fullest requires a font with good support

Contributing

We welcome issue reports and pull requests, including potentially adding support for commonly used variable classes. However, in general, we encourage users to take advantage of skimr's flexibility to add their own customized classes. Please see the [contributing](#) and [conduct](#) documents.

Links

[View on CRAN](#)

[Browse source code](#)

[Report a bug](#)

License

[GPL-3](#)

Community

[Contributing guide](#)

Citation

[Citing skimr](#)

Developers

Elin Waring

Maintainer, author

Michael Quinn

Author

Amelia McNamara

Author

Eduardo Arino de la Rubia

Author

Hao Zhu

Author

Shannon Ellis

Author

[More about authors...](#)

skimr

Software Peer-Review

 Peer Reviewed



ABOUT

About
rOpenSci
Software
Review
Our Team
Jobs
Donate
Contact Us

COMMUNITY

Our
Community
Community
calls
Events
Join the
Discussion
Code of
conduct

RESOURCES

Packages
Use Cases
Talks &
Publications
Documentation
Newsletter
Cite rOpenSci

rOpenSci is a fiscally sponsored project of
NumFOCUS.