

Reshaping Data with tidyr in R

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> Content

Definitions

- The majority of data analysis in R is performed in **data frames**. These are rectangular datasets consisting of rows and columns.
- An **observation** contains all the values or variables related to a single instance of the objects being analyzed. For example, in a dataset of movies, each movie would be an observation.
- A **variable** is an attribute for the object, across all the observations. For example, the release dates for all the movies.
- Tidy data** provides a standard way to organize data. Having a consistent shape for datasets enables you to worry less about data structures and more on getting useful results. The principles of tidy data are:

- Every column is a variable.
- Every row is an observation.
- Every cell is a single value.

> Helpful syntax before getting started

Installing and loading tidyr

```
# Install tidyr through tidyverse
install.packages("tidyverse")
```

```
# Install it directly
install.packages("tidyr")
```

```
# Load tidyr into R
library(tidyr)
```

The %>% Operator

%>% is a special operator in R found in the magrittr and tidyr packages. %>% lets you pass objects to functions elegantly, and helps you make your code more readable. The following two lines of code are equivalent.

```
# Without the %>% operator
second_function(first_function(dataset, arg1, arg2), arg3)
```

```
# With the %>% operator
dataset %>% some_function(arg1, arg2) %>% second_function(arg3)
```

> Datasets used throughout this cheat sheet

Throughout this cheat sheet we will use a dataset of the top grossing movies of all time, stored as movies.

title	release_year	release_month	release_day	directors	box_office_busd
Avatar	2009	12	18	James Cameron	2.922
Avengers: Endgame	2019	4	22	Anthony Russo, Joe Russo	2.798
Titanic	1997	11	01	James Cameron	2.202
Star Wars Ep. VII: The Force Awakens	2015	12	14	J.J Abrams	2.068
Avengers: Infinity War	2018	4	23	Anthony Russo, Joe Russo	2.048

The second dataset involves an experiment with the number of unpopped kernels in bags of popcorn, adapted from the Popcorn dataset in the Stat2Data package.

brand	trial_1	trial_2	trial_3	trial_4	trial_5	trial_6
Orville	26	35	18	14	8	6
Seaway	47	47	14	34	21	37

The third dataset is JSON data about music containing nested elements. The JSON is parsed into nested lists using parse_json() from the jsonlite package.

artist	singles	
	Title	Tracks
Bad Bunny	Gato de Noche	Gato de Noche, Ñengo Flow
	La Jumpa	La Jumpa, Arcángel
	Scary Hours 2	What's Next, Wants and Needs, Lemon Pepper Freestyle, NA, Lil Baby, Rick Ross

The fourth dataset is a synthetic dataset containing attributes of people. sex is a character vector, and hair_color is a factor.

sex	hair_color	height_cm	weight_kg
female	brown	166	72
male	blonde	184	
female	black	153	
male	black	192	93

> Uniting and separating columns

```
# Combine several columns into a single vector column with unite()
movies %>%
  unite(release_date, c(release_year, release_month, release_day), sep = "-")
```

```
# Split a single vector column into several columns with separate()
movies %>%
  separate(directors, into = c("director1", "director2"), sep=",", fill = "right")
```

```
# Split a single column into several rows with separate_rows()
movies %>%
  separate_rows(directors, sep=",")
```

> Packing and unpacking columns

```
# Combine several columns into a data frame column with pack()
movies_packed <- movies %>%
  pack(release_date = c(release_year, release_month, release_day))
# The release_date column is a data frame with 5 rows, 3 columns
```

```
# Split a single data frame column into several columns with unpack()
movies_packed %>%
  unpack(release_date)
# release_date column replaced with release_year/release_month/release_day columns
```

> Pivoting

```
# Move side-by-side columns to consecutive rows with pivot_longer()
popcorn_long <- popcorn %>%
  pivot_longer(trial_1:trial_6, names_to = "trial", values_to = "n_unpopped")
# "brand" columns contains "Orville" and "Seaway"
# "trial" column contains "trial_1" to "trial_6"
# "n_unpopped" column contains the numbers
```

```
# Move values in different rows to columns with pivot_wider()
popcorn_long %>%
  pivot_wider(brand, names_from = "trial", values_from = "n_unpopped")
# Same contents and shape as popcorn dataset
```

> Nesting and unnesting

```
# Expand nested data frame columns with unnest_longer()
# Vectors inside the nested data are given their own row
# The number of columns remains unchanged
music %>%
  unnest_longer(singles)
```

artist	single\$title	singles\$tracks
Bad Bunny	Gato de Noche	2 Variables
Bad Bunny	La Jumpa	2 Variables
Drake	Scary Hours 2	1 Variable

```
# Expand nested data frame columns with unnest_wider()
# Top-level elements inside the nested data are given their own column
# The number of rows remains unchanged
music %>%
  unnest_wider(singles)
```

artist	title	tracks
Bad Bunny	["Gato de Noche", "La Jumpa"]	[[{"title": "Gato de Noche", "collaborator": "Ñengo Flow"}, {"title": "..."}]]
Drake	["Scary Hours 2"]	[[{"title": "What's Next"}, {"title": "wants and needs", "collaborator": "..."}]]

```
# Expand selected nested data frame columns with hoist()
# Replacement for unnest_wider() %>% select()
music %>%
  hoist(singles, single_titles = "title")
```

artist	single_titles	singles
Bad Bunny	["Gato de Noche", "La Jumpa"]	[[{"tracks": [{"title": "Gato de Noche", "collaborator": "Ñengo Flow"}]]
Drake	["Scary Hours 2"]	[[{"tracks": [{"title": "What's Next"}, {"title": "Wants and Needs", "collaborator": "..."}]]

```
# Expand nested data frame columns with unnest_longer()
# Every top-level element of the nested data gets its own column in the result
# Vectors inside the nested data are given their own row
music_unnested <- music %>%
  unnest(singles)
# Roughly equivalent to music %>% unnest_longer(singles) %>% unnest_wider(singles)
```

artist	title	tracks
Bad Bunny	Gato de Noche	[[{"title": "Gato de Noche", "collaborator": "Ñengo Flow"}]]
Bad Bunny	La Jumpa	[[{"title": "La Jumpa", "collaborator": "Arcángel"}]]
Drake	Scary Hours 2	[[{"title": "What's Next"}, {"title": "Wants and Needs", "collaborator": "Lil Baby"}, {"title": "..."}]]

```
# Summarize parts of a data frame as a list of dataframes with nest()
music_unnested %>%
  nest(singles = c(title, tracks))
```

artist	singles
Bad Bunny	[[{"title": "Gato de Noche", "tracks": [{"title": "Gato de Noche", "collaborator": "Ñengo Flow"}]}, {"title": "La Jumpa", "tracks": [{"title": "La Jumpa", "collaborator": "Arcángel"}]}]]
Drake	[[{"title": "Scary Hours 2", "tracks": [{"title": "What's Next"}, {"title": "Wants and Needs", "collaborator": "Lil Baby"}, {"title": "..."}]}]]

> Dealing with missing data

```
# Drop rows containing any missing values in the specified columns with drop_na()
people %>%
  drop_na(weight_kg)
```

```
# Replace missing values with a default value with replace_na()
people %>%
  replace_na(list(weight_kg = 100))
```

> Creating grids

```
# Get all combinations of input values with expand_grid()
expand_grid(
  sex = c("male", "female", "female"),
  hair_color = c("red", "brown", "blonde", "black", "red")
)
# 2 column data frame with rows like "male", "red".
```

```
# Get all combinations of input values, deduplicating and sorting with crossing()
crossing(
  sex = c("male", "female", "female"),
  hair_color = c("red", "brown", "blonde", "black", "red")
)
# Same as expand_grid() but "red" rows only appear once and order is alphabetical
```

```
# Get all combinations of values in data frame columns with expand()
# All factor levels included, even if they don't appear in data
people %>%
  expand(sex, hair_color)
# Equivalent to expand_grid(unique(people$sex), levels(people$hair_color))
```

```
# Get all combinations of values that exist in data frame columns with expand() + nesting()
people %>%
  expand(nesting(sex, hair_color))
# As previous, but filtered to rows that exist in people dataset
```

```
# Expand the data frame, then full join to itself with complete()
people %>%
  complete(sex, hair_color)
# Same output as expand, with additional height_cm and weight_kg columns
```

```
# Fill in sequence of numeric or datetime columns with expand() + full_seq()
people %>%
  expand(height_cm_expanded = full_seq(height_cm, 1))
# 1 column data frame with height_cm_expanded values
# from min height_cm to max height_cm in steps of 1
```