Preliminaries in R

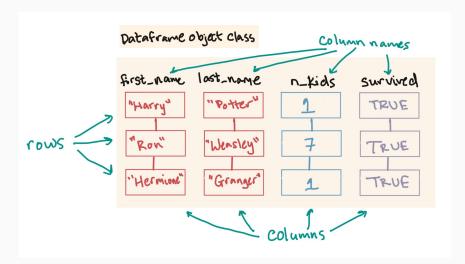
Dataframes

Structure of dataframe objects



A **dataframe** combines one or more vectors of the same length stuck together side-by-side.

Structure of dataframe objects



A dataframe contains **rows** and **columns**, and each column has a **column** name.

We'll be working with a specific class of dataframe called a tibble.

You can create tibble dataframe using the tibble function from the tibble package.

However, most often you will create a dataframe by reading in data from a file—most datasets will not be short enough that you want to enter them by hand in R.

We'll look at both methods of creating dataframes.

The format for creating a tibble dataframe using the tibble function is:

```
library(package = "tibble")
hp_data <- tibble(first_name = c("Harry", "Ron", "Hermione"),
                last name = c("Potter", "Weasley", "Granger"),
                n kids = c(1, 7, 1),
                survived = c(TRUE, TRUE, TRUE))
hp data
## # A tibble: 3 x 4
##
    first_name last_name n_kids survived
## <chr> <chr> <dbl> <lgl>
## 1 Harry Potter 1 TRUE
## 2 Ron Weasley 7 TRUE
## 3 Hermione Granger 1 TRUE
```

You can also create dataframes by joining together vector objects you previously created, as long as they have the same length and line up:

Usually, instead of creating a dataframe from vectors, you'll read one in from data on an outside file, for example using read_csv from the readr package.

For example, to read in a dataset from a csv file called "daily_show_guests.csv":

This has read data in from the external file into a dataframe object in my R session:

```
ls()
                                             "main_characters" "n_
## [1] "daily_show"
                          "hp_data"
```

Dataframes

You can use the functions dim, nrow, and ncol to figure out the dimensions (number of rows and columns) of a dataframe:

```
dim(x = daily_show)

## [1] 2693     5

nrow(x = daily_show)

## [1] 2693

ncol(x = daily_show)

## [1] 5
```

Dataframes

Base R also has some useful functions for quickly exploring dataframes:

- str: Show the structure of an R object, including a dataframe
- summary: Give summaries of each column of a dataframe.

The dplyr package has two functions for extracting data from dataframes by position: slice to extract rows based on their row position and select to extract columns based on their column position.

For example, if you wanted to get the first two rows of the hp_data dataframe, you could run:

If you wanted to get the first and fourth columns, you could run:

```
select(.data = hp_data, c(1, 4))

## # A tibble: 3 x 2

## first_name survived

## <chr> <lgl>
## 1 Harry TRUE

## 2 Ron TRUE

## 3 Hermione TRUE
```

You can compose calls from both functions. For example, you could extract the values in the first and fourth columns of the first two rows with: