

Data Cleaning with Tidyverse

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Resources

- [Tidyverse Documentation](#)
- [O'Reilly Learning Platform](#)
 - R for Data Science, 2nd Edition
 - R Programming for Statistics and Data Science

```
setwd("path_to_folder")
```

Part 1: About Tidyverse

```
install.packages("tidyverse")
```

```
library(tidyverse)
```

Tidyverse is a collection of packages focused on data analysis and data visualizations that share an underlying design philosophy, grammar, and data structures.

Packages Included in Tidyverse

tibble	lighter and more user-friendly version of data frames
tidyr	create tidy and meaningfully arranged data
readr	better importation of data into R
ggplot	data visualization functions
dplyr	data manipulation tools
lubridate	clean dates and times
purrr	better functional programming
forcats	handle, clean, and manipulate categorical variables

Part 2: Loading Data with Tidyverse

Readr function: read_csv

allows you to read a csv file into a tibble data frame

```
sw_df <- read_csv("starwars.csv")
```

Loading Proprietary Data

- [readxl](#) - this package allows you to read Excel files in a tibble data frame
- [haven](#) - this package allows you to read and export non-proprietary files for SPSS, SAS, and STATA

The Pipe Operator

The pipe operator allows you to run commands or operation on a single object based on an order of operations

- let's say you want to see the **name**, **height**, **mass**, and **species** of characters who were born on **Tatooine**

```
sw_df |> # object we are working on
  filter(homeworld == "Tatooine") |> # first operation
  select(name, height, mass, species) # second operation

# order of operations matter

sw_df |> # object we are working on
  select(name, height, mass, species) |> # first operation
  filter(homeworld == "Tatooine") # second operation

# why did this not work?
```

Tibble function: view

view the contents of a data frame in a separate viewer window or in the RStudio viewer pane.

```
view(sw_df)
```

Tibble function: glimpse

like the `str()` function in base `r`, this allow you see the structure of your data but in a more compact manner

```
glimpse(sw_df)
```

Part 3: Cleaning Data

Main Tidyverse Functions

filter	retains or filters out observations based on variable criteria
select	retains or filters out variables
arrange	sorts variables
mutate	change variable's observations OR create a new variable and observations using observations from another variable
group_by	group observations
summarise	get descriptive statistics about a variable
relocate	change the position of variables in the data frame
rename	change the name of an individual variable
drop_na	remove ALL missing values from a data frame or variable
replace_na	replace missing values with a specified

Dplyr function: filter

the **filter** function allows you to select rows in your data frame that meet specific conditions or criteria in a variable

```
sw_df

# let's filter the data frame so we are seeing characters who have blue eyes

sw_eye <- sw_df |>
  filter(eye_color == "blue")

sw_eye
```

Boolean operators

boolean operators allows you to build criteria in your code

Boolean operators

&	AND
	OR
==	EQUAL
!=	NOT EQUAL
<	LESS THAN
>	GREATER THAN
<=	LESS THAN OR EQUAL
>=	GREATER THAN OR EQUAL

```
# let's filter the data frame for characters who
# do have blue eyes
# and were born after 50 BBY

sw_eye50 <- sw_df |>
  filter(eye_color == "blue" & birth_year < 50)

sw_eye50
```

Dplyr function: select

the **select** function allows you to *keep* or *discard* variables

```
# keep variables

sw_select <- sw_df |>
  select(name, height, mass)

sw_select

# remove variables

sw_not_select <- sw_df |>
  select(-height, -mass)

sw_not_select
```

Dplyr function: mutate

the **mutate** function *creates* new variables in your data or *change* existing variables by performing calculations or transformations.

NOTE: if you name your variable as an *existing variable*, it will *overwrite* the existing variable. If you give it a *new name*, it will create a *new variable*

```
# create a new variable

sw_df

sw_df <- sw_df |>
  mutate(bmi = height/mass) |> # run mutate operation
  relocate(bmi, .after = mass) # relocate variable in data frame

# let's overwrite the old variable

sw_overwrite <- sw_df |>
  mutate(height = height/12) # overwrite variable

sw_overwrite
```

Dplyr function: arrange

the **arrange** function allows you to *sort* variables

```
# oldest characters

sw_df |>
  arrange(desc(birth_year))

# characters with the same skin color than the same hair color

sw_df |>
  arrange(desc(skin_color), hair_color)
```

Dplyr function: group_by & summarise

the **group_by** function allows you to *group* common observations in a variable and **summarise** function allows you to get descriptive statistics about the groupings

```
sw_group <- sw_df |>
  group_by(sex) |>
  summarise(avg_height = mean(height)) # get the mean of height for each group

sw_group

# you can add as many summaries as you want

sw_group <- sw_df |>
  group_by(sex) |>
  summarise(avg_height = mean(height), # get mean
            n = n()) # get number

sw_group

write_csv(sw_group, "sw_sex_dv.csv") # export
```

Base Function: as.character

The **as.** function along with **mutate** will allow you to change the data type of a variable. For this example we are going to recode the *character_id* variable to interpret the data type as a *character* instead of a *double*

```
sw_df <- sw_df |>
  mutate(character_id = as.character(character_id))
```

```
sw_df
```

Dyplr Function: recode

we can rename the values of observations within a variable using the **mutate** function in combination with the **recode** or **recode_factor** functions

```
# let's create a new variable with to give numeric levels instead of value labels

sw_df <- sw_df |>
  mutate(gender_num = recode(gender,
                             "masculine" = 0,
                             "feminine" = 1)) |>
  relocate(gender_num, .after = gender)

sw_df
```

Dplyr function: rename

the **rename** function allows you rename variables in your data frame

```
glimpse(sw_df)

sw_df <- sw_df |>
  rename("gender_label" = gender)

glimpse(sw_df)
```

Tidyr function: drop_na

we can remove **all** missing data from data frames or variables using the **drop_na** function

```
# we can see if are data frame has missing NA values using the is.na function.

which(is.na(sw_df$bmi))

# because there are missing values we cannot calculate some descriptive statistics
```

```
mean(sw_df$bmi)

# we can drop all NA values from the data frame

sw_dropNA <- sw_df |>
  drop_na()

mean(sw_dropNA$bmi)

# we can also just drop NAs from a variable

sw_dropNA_var <- sw_df |>
  drop_na(bmi)

mean(sw_dropNA_var$bmi)
```

Tidyr function: replace_na

you can also recode the NA values for observations with the **replace_na** function

```
# let's replace the NAs the gender_label variable with "unknown"

sw_df <- sw_df |>
  mutate(gender_label = replace_na(gender_label, "unknown")) |>
  mutate(gender_label = as_factor(gender_label))

levels(sw_df$gender_label)
```

Readr function: write_csv

the **write_csv** function allows us to export data frames to a csv file once we are done cleaning it up or when we have done some analysis that we want to export

```
# now that we have this data frame cleaned let's save it

# let's export the file

write_csv(sw_df, "starwars_clean.csv")
```

Part 4: Explore Your Data

- [Psych Package](#) - built-in functions for factor analysis, reliability analysis, descriptive statistics and data visualization.

```
install.packages("psych")  
library(psych)
```

```
sw_ds <- describe(sw_df)  
  
write_csv(sw_ds, "starwars_ds.csv")
```

- [SummaryTools Package](#) - simplifies data exploration and descriptive statistics generation for data frames and vectors.

```
install.packages("summarytools")  
library(summarytools)
```

```
descr(sw_df)  
  
freq(sw_df$sex)  
  
ctable(sw_df$sex, sw_df$gender)
```

- [DataExplorer package](#) - automates and streamlines the process of exploring and visualizing datasets.

```
install.packages("DataExplorer")  
library(DataExplorer)
```

```
create_report(reviews_df)
```