

# Final Project Data

AUTHOR  
Tan Vo

```
life_expectancy <- readr::read_csv('https://raw.githubusercontent.com/rfordatascience/ti
life_expectancy_different_ages <- readr::read_csv('https://raw.githubusercontent.com/rfo
life_expectancy_female_male <- readr::read_csv('https://raw.githubusercontent.com/rforda
skim(life_expectancy)
```

Data summary

Name	life_expectancy
Number of rows	20755
Number of columns	4

Column type frequency:	
character	2
numeric	2

Group variables	None
-----------------	------

Variable type: character

skim_variable	n_missing	complete_rate	min	max	empty	n_unique	whitespace
Entity	0	1.00	4	59	0	261	0
Code	1694	0.92	3	8	0	238	0

Variable type: numeric

skim_variable	n_missing	complete_rate	mean	sd	p0	p25	p50	p75	p100	hist
Year	0	1	1975.73	38.42	1543	1961.00	1981.00	2001.00	2021.00	
LifeExpectancy	0	1	61.62	12.92	12	52.19	64.06	71.82	86.54	

```
skim(life_expectancy_different_ages)
```

Data summary

Name	life_expectancy_different...
Number of rows	20755
Number of columns	9

Column type frequency:	
character	2
numeric	7

Group variables	None
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Variable type: character

skim_variable	n_missing	complete_rate	min	max	empty	n_unique	whitespace
Entity	0	1.00	4	59	0	261	0
Code	1694	0.92	3	8	0	238	0

Variable type: numeric

skim_variable	n_missing	complete_rate	mean	sd	p0	p25	p50	p75	p100	hist
Year	0	1.00	1975.73	38.42	1543.00	1961.00	1981.00	2001.00	2021.00	
LifeExpectancy0	0	1.00	61.62	12.92	12.00	52.19	64.06	71.82	86.54	
LifeExpectancy10	833	0.96	68.82	7.08	22.11	63.63	69.49	73.95	86.96	
LifeExpectancy25	833	0.96	70.46	5.89	31.16	66.07	70.71	74.67	87.07	
LifeExpectancy45	833	0.96	73.54	4.07	50.03	70.51	73.22	76.21	87.31	
LifeExpectancy65	833	0.96	78.75	2.51	68.74	76.91	78.31	80.19	88.35	
LifeExpectancy80	833	0.96	86.09	1.30	82.10	85.20	85.80	86.72	91.57	

```
skim(life_expectancy_female_male)
```

Data summary

Name	life_expectancy_female_ma...
Number of rows	19922
Number of columns	4

Column type frequency:	
character	2
numeric	2

Group variables	None
-----------------	------

Variable type: character

skim_variable	n_missing	complete_rate	min	max	empty	n_unique	whitespace
Entity	0	1.00	4	59	0	259	0
Code	1600	0.92	3	8	0	237	0

Variable type: numeric

skim_variable	n_missing	complete_rate	mean	sd	p0	p25	p50	p75	p100	his
Year	0	1	1978.82	32.81	1751.00	1963.00	1983.0	2002.00	2021.00	
LifeExpectancyDiffFM	0	1	4.82	2.38	-9.88	3.23	4.6	6.22	29.55	

- **DATA EVALUATION**

Is the data in the right format for those mappings? - Yes.

Determine if the variables you have are the appropriate variable type. - Yes, it is appropriate.

Does something you want to use as a numeric variable look like a character string? - No.

See if there are any missing values, and if there are, if they are coded correctly. - No missing value.

Are the variable names usable? - Yes.

Do they have spaces in them that will make it hard to refer to them in R? - No.

For categorical variables, consider the categories. Are they readable? - Yes.

Are they understandable phrases or cryptic codes? - They are understandable phrases.

Are there the right number of categories, or will you want to reduce them? - They are correct but might consider to group them into different groups.

- **DATA WRANGLING**

1. Average Life Expectancy from 1950 to 2021

```
avg_life_expectancy <- life_expectancy |>
  filter(Year >= 1950 & Year <= 2021, !is.na(Code), Code != "OWID_WRL") |>
  group_by(Entity, Code) |>
  summarize(AvgLifeExpectancy = mean(LifeExpectancy, na.rm = TRUE)) |>
  ungroup()
```

`summarise()` has grouped output by 'Entity'. You can override using the  
`.groups` argument.

Export the filtered data as a CSV file

```
write_csv(avg_life_expectancy, "avg_life_expectancy.csv")
```

2. Average Life Expectancy Difference between Female & Male from 1950 to 2021

```
avg_life_expectancy_diff_female_male <- life_expectancy_female_male |>
  filter(Year >= 1950 & Year <= 2021, !is.na(Code), Code != "OWID_WRL") |>
  group_by(Entity, Code) |>
  summarize(AvgLifeExpectancyDiffFM = mean(LifeExpectancyDiffFM, na.rm = TRUE)) |>
  ungroup()
```

`summarise()` has grouped output by 'Entity'. You can override using the  
`.groups` argument.

Export the filtered data as a CSV file

```
write_csv(avg_life_expectancy_diff_female_male, "avg_life_expectancy_diff_female_male.csv")
```