

## ds4biomed

## Exercise 2

## Pre-Workshop Exercise

Exercise 1

Exercise 2

Exercise 3

Summative

Start Over

This is a different version of the `ebola` dataset.

date <chr>	da <dbl>	Cases_Guinea <dbl>	Cases_Liberia <dbl>
1/5/2015	289	2776	NA
1/4/2015	288	2775	NA
1/3/2015	287	2769	8166
1/2/2015	286	NA	8157
12/31/2014	284	2730	8115
12/28/2014	281	2706	8018
12/27/2014	280	2695	NA
12/24/2014	277	2630	7977
12/21/2014	273	2597	NA
12/20/2014	272	2571	7862
1... Previous	1	2	3

- Tidy the dataset such that you get the dataset below
- Remember to drop missing values as the last step

date <chr>	da <dbl>	case_death <chr>	country <chr>	val... <dbl>
1/5/2015	289	Cases	Guinea	2776
1/5/2015	289	Cases	SierraLeone	10030
1/5/2015	289	Deaths	Guinea	1786
1/5/2015	289	Deaths	SierraLeone	2977
1/4/2015	288	Cases	Guinea	2775
1/4/2015	288	Cases	SierraLeone	9780
1/4/2015	288	Deaths	Guinea	1781
1/4/2015	288	Deaths	SierraLeone	2943
1/3/2015	287	Cases	Guinea	2769
1/3/2015	287	Cases	Liberia	8166

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Summative

Start Over

1... Previous **1** 2 3 4 5 6 ... 74 Next

R Code

 Start Over

 Run Code

```
1 ebola %>%  
2   ____ (____) %>%  
3   ____ (____, into = ____, sep = ____ ) %>%  
4   ____
```

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