## ds4biomed

## Pre-Workshop Exercise

Exercise 1

Exercise 2

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## Exercise 1

Take a look at the ebola dataset.

ebola

date <chr></chr>	country <chr></chr>	<b>289</b> <dbl></dbl>		<b>287</b> · <dbl></dbl>	<b>286</b> · <dbl></dbl>
1/5/2015	Guinea	2776	NA	NA	NA
1/5/2015	SierraLeone	10030	NA	NA	NA
1/4/2015	Guinea	NA	2775	NA	NA
1/4/2015	SierraLeone	NA	9780	NA	NA
1/3/2015	Guinea	NA	NA	2769	NA
1/3/2015	Liberia	NA	NA	8166	NA
1/3/2015	SierraLeone	NA	NA	9722	NA
1/2/2015	Liberia	NA	NA	NA	8157
12/31/2014	Guinea	NA	NA	NA	NA
12/31/2014	Liberia	NA	NA	NA	NA
1 Previous	1 2 3 4	4 5	6	37 I	Vext

- Tidy the dataset such that you get the dataset below.
- You can use the last\_col() to select the last column of the dataset.
- Remember to drop missing values as the last step.

date <chr></chr>	country <chr></chr>	day cases <chr> <dbl></dbl></chr>
1/5/2015	Guinea	289 2776
1/5/2015	SierraLeone	289 10030
1/4/2015	Guinea	288 2775
1/4/2015	SierraLeone	288 9780
1/3/2015	Guinea	287 2769
1/3/2015	Liberia	287 8166

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date <chr></chr>		cou <chi< th=""><th>ntry ^&gt;</th><th></th><th></th><th>day <chr></chr></th><th>cases <dbl></dbl></th></chi<>	ntry ^>			day <chr></chr>	cases <dbl></dbl>
1/3/2015		SierraLeone			287	9722	
1/2/2015		Liberia		286	8157		
12/31/2014		Guinea		284	2730		
12/31/2014		Liberia			284	8115	
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R Code	Start Over	▶ Run Code
1 ebol		to =, values_to =) %:

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