## ds4biomed

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## Pre-Workshop Exercise

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## Exercise 3

Using the same <code>ebola</code> dataset example from the previous exercise:

date <chr></chr>	da Cas	es_Guinea <dbl></dbl>	Cases_Liberia <dbl></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
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- Tidy the dataset such that it looks like the below example
- Remember to drop missing values as the last step

date <chr></chr>	day country <dbl×chr></dbl×chr>	<b>Ca</b> <dbl></dbl>	Deaths <dbl></dbl>
1/5/2015	289 Guinea	2776	1786
1/5/2015	289 SierraLeone	10030	2977
1/4/2015	288 Guinea	2775	1781
1/4/2015	288 SierraLeone	9780	2943
1/3/2015	287 Guinea	2769	1767
1/3/2015	287 Liberia	8166	3496
1/3/2015	287 SierraLeone	9722	2915
1/2/2015	286 Liberia	8157	3496
12/31/2014	284 Guinea	2730	1739

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ds4biomed	date <chr></chr>	day c <dbl×< th=""><th>country chr&gt;</th><th></th><th><b>Ca</b> <dbl></dbl></th><th>Deaths <dbl></dbl></th></dbl×<>	country chr>		<b>Ca</b> <dbl></dbl>	Deaths <dbl></dbl>
	12/31/2014	284 L	iberia		8115	3471
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