

name	label	data_type	n_missing	complete_rate	n_unique	empty	min	median	max	mean	sd	whitespace	hist
<input type="text" value="All"/>	<input type="text" value="All"/>	<input type="text" value="All"/>	<input type="text" value="All"/>	<input type="text" value="All"/>	<input type="text" value="All"/>	<input type="text" value="All"/>	<input type="text" value="A"/>	<input type="text" value="All"/>	<input type="text" value="All"/>	<input type="text" value="All"/>	<input type="text" value="All"/>	<input type="text" value="All"/>	<input type="text" value="All"/>
<a href="#">subj</a>	de-identified subject id	character	0	1	102	0	1		6			0	
<a href="#">experiment.orig</a>	original name of experiment	character	0	1	6	0	3		5			0	
<a href="#">experiment.paper</a>	name of expeirment used in paper	character	0	1	6	0	4		11			0	
<a href="#">experiment.new</a>	specific name of sample to distinguish between older and younger infants	character	0	1	6	0	12		19			0	
<a href="#">trial</a>	which test trial (test1-test4)	character	0	1	4	0	5		5			0	
<a href="#">orig.look</a>	looking time generated by original offline coding (used in analysis)	numeric	16	0.96078431372549			0.95	14.6	60	20.6620714285714	16.2516161248682		
<a href="#">trialn</a>	index of trial	numeric	0	1			1.00	2.5	4	2.5	1.11940665223647		
<a href="#">secondary.look</a>	looking time generated by second coder	numeric	12	0.970588235294118			0.95	14.6	60	20.7583737373737	16.38607524009		
<a href="#">coder</a>	who did the secondary coding	character	0	1	3	0	5		7			0	