

## Appendix 1. Exclusions in the dataset

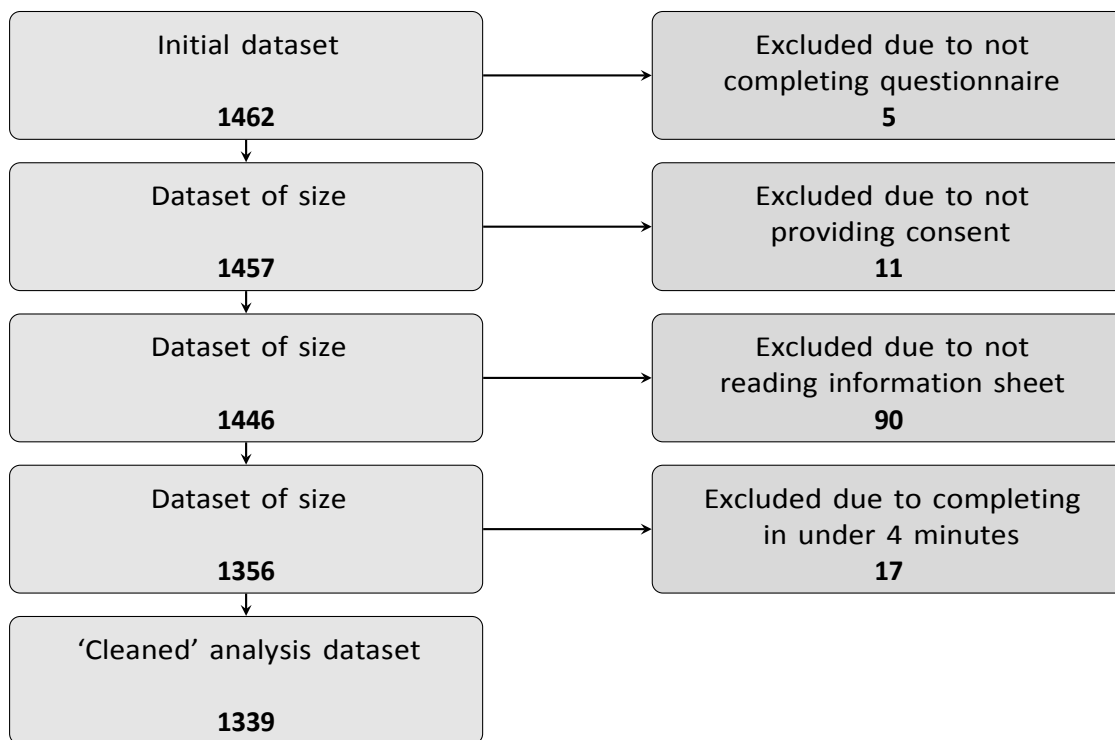


Figure A2 Initial exclusions made to dataset

## Appendix 2

The attitude items were summed using equation 1, and the attitude scale was then standardised using equation 2) to create a summary measure of attitudes towards length and quality of life which is referred to as  $ATLQL_{stan}$ , throughout the analyses. The ATLQL score is highest for those who indicated quality of life was most important and lowest for those who considered length more important.

$$ATLQL_{norm} = \frac{(A_1 + (6 - A_2) + (6 - A_3) + A_4) - 4}{16} \in [0, 1] \quad (1)$$

$$ATLQL_{stan} = \frac{ATLQL_{norm} - \bar{x}}{\sigma} \quad (2)$$

### **Appendix 3 –Pooling data across the different TTO variants**

We ran a regression to explore the extent to which we could pool responses from the four TTO variants to which respondents were randomised involving slightly different elicitation procedures.

The regression model that was constructed containing dummy variables representing the four TTO variants and all seven states, with a dummy for the severity of the health states to which the respondent was randomised (Mild and Moderate). Also included were interaction terms (cross products) between states and TTO variants. This regression was run to establish if any variants were significantly affecting elicited TTO values. A significant  $p$ -value for a given variant-term<sup>1</sup> would indicate the TTO values generated from this variant were significantly different to those obtained under other variants, and therefore could not be pooled together. In the event that particular variant-terms were found to be significant, it was determined whether the values obtained from these variants could be pooled (and which if any must be dropped) by means of an  $F$ -test. The regression was run on the cleaned dataset of 1339, and the results are shown in Table A1).

Here we label these variants IS: Iterative Sequential; NIS: Non-iterative Sequential; IC: Iterative Concurrent; NIC: Non-iterative Concurrent.

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<sup>1</sup> We will use 'variant-term' to refer to any term included in the regression models which contains one of *IC*, *NIC* or *NIS* either as a cross product or alone.

**Table 1** Model (1) Output for Effect of TTO Variant on TTO Values

Util	Coefficient	Std. Err.	z	$p >  z $
11121	0.1632	0.0209	7.83	0.000*
21211	0.1387	0.0202	6.87	0.000*
12212	0.0788	0.0206	3.83	0.000*
13224	-0.2540	0.0213	-11.92	0.000*
23242	-0.2972	0.0216	-13.78	0.000*
23314	-0.2454	0.0215	-11.43	0.000*
Moderate	0.0149	0.0363	0.41	0.681
IS	0.0220	0.0346	0.64	0.524
NIS	0.0230	0.0352	0.65	0.514
IC	-0.0219	0.0345	-0.63	0.526
11121_IS	-0.0212	0.0283	-0.75	0.454
21211_IS	-0.0235	0.0278	-0.84	0.399
12212_IS	-0.0942	0.0281	-3.35	0.001*
13224_IS	0.0229	0.0289	0.79	0.428
23242_IS	0.0054	0.0291	0.19	0.853
23314_IS	-0.0055	0.0290	-0.19	0.849
Even_IS	0.0209	0.0498	0.42	0.675
11121_NIS	0.0243	0.0294	0.83	0.409
21211_NIS	0.0047	0.0285	0.17	0.868
12212_NIS	-0.0281	0.0292	-0.96	0.336
13224_NIS	-0.0393	0.0296	-1.32	0.185
23242_NIS	-0.0575	0.0299	-1.92	0.055
23314_NIS	-0.0657	0.0299	-2.20	0.028*
Even_NIS	0.0262	0.0506	0.52	0.604
11121_IC	0.0095	0.0283	0.34	0.736
21211_IC	-0.0052	0.0278	-0.19	0.852
12212_IC	-0.0469	0.0282	-1.66	0.096
13224_IC	0.0206	0.0287	0.72	0.474
23242_IC	-0.0105	0.0290	-0.36	0.718
23314_IC	-0.0055	0.0289	-0.19	0.849
Even_IC	-0.0206	0.0496	-0.42	0.677
Constant	0.6524	0.0250	26.12	0.000*

\*Significant at the 5% level.

This regression used variant NIC and state 13122 as the base case and therefore included no dummies for this variant, state or their interactions.

The dummies for the state being valued are significant as expected. The only other terms which are significantly contributing to the TTO values are  $12212 \times IS$  ( $p$ -value = 0.001) and  $23314 \times NIS$  ( $p$ -value = 0.028). We interpret that - relative to the base-case variant, IS does cause respondents to give systematically different values for health state 12212, and similarly for 23314 under variant NIS. These results suggest that we cannot pool data collected from the four variants in their entirety.

We set up an  $F$ -test with the null and alternative hypotheses were set up as follows:

- $H_0$ : The full model does not provide a significantly better fit than the restricted.
- $H_1$ : The full model provides a significantly better fit than the restricted.

For a  $p$ -value  $\geq \alpha = 0.05$  we would fail to reject  $H_0$ , and pooling would not be appropriate.

We test whether dropping the 12212×*IS* and 23314×*NIS* terms would allow us to pool the remaining values across variants and health states. We conduct an  $F$ -test with Model (1) reported in table A1 as the unrestricted, and Model (2) as the restricted model.

$$util = \alpha + \beta_1 11121 + \beta_2 21211\epsilon + \beta_3 12212 + \beta_4 13224 + \beta_5 23242 + \beta_6 23314 + \beta_7 \text{EVEN} + \beta_8 12212 \times IS + \beta_9 23314 \times NIS + \epsilon \quad \text{Model (2)}$$

Model (2) contains only two variant-terms, 12212×*IS* and 23314×*NIS*, which were significant in Model (1). The test returns a  $p$ -value of 0.1191 which is not significant at the 5% level and therefore there is no significant difference between the way the two models fit the data. That is, 12212×*IS* and 23314×*NIS* jointly influence respondent's values to the same extent as 12212×*IS* + 23314×*NIS* + {*All other variant-terms*}. A corollary of this is that if these variables 12212×*IS* and 23314×*NIS* were to be dropped - along with the observations to which they apply<sup>2</sup>, the remaining values elicited across all four variants could be pooled. Model 1 was rerun on the dataset with the two variant terms omitted, and confirms that no variant-term is now contributing to the values ( $\chi^2(23) = 31.15$  Prob >  $\chi^2 = 0.1191$ ).

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<sup>2</sup> These numbered 167 and 169 respectively - note however that no respondents were dropped, only one value for each of the IS and NIS respondents



## Appendix 4

We ran the choice regression with three-way interactions and used a chi-test to determine whether the dummies and their interactions were simultaneously zero. This is similar to testing for significant differences between a model with these variant variables added and a model without them, i.e. the difference between full model and reduced models.

**Table 1** Choices

<i>Latent propensity to choose Life</i>	Base Model				Base model with attitudes			
	Coefficient	Std. Err.	z	$p> z $	Coefficient	Std. Err.	z	$p> z $
<i>A</i>								
Constant	1.1295	0.1354	8.34	0.000*	1.078	0.1376	7.68	0.000*
Years	0.0935	0.0167	5.59	0.000*	0.0871	0.0170	4.92	0.000*
Male	-0.0558	0.0738	-0.76	0.449	-0.0550	0.0748	-0.74	0.462
Age	-0.0017	0.0027	-0.62	0.534	-0.0005	0.0027	-0.19	0.853
Moderate	-0.4787	0.0741	-6.46	0.000*	-0.5080	0.0754	-6.74	0.000*
Male× years	0.0213	0.0096	2.20	0.028*	0.0104	0.0099	1.05	0.293
Age× years	0.0008	0.0004	2.27	0.023*	0.0011	0.0003	3.41	0.001*
Moderate_×years	-0.0917	0.0107	-8.55	0.000*	-0.0947	0.0109	-8.71	0.000*
<i>ATLQL<sub>stan</sub></i>					0.0476	0.0386	1.24	0.215
<i>ATLQL<sub>stan</sub>×Y ears</i>					-0.0236	0.0179	-1.32	0.187
<i>ATLQL<sub>stan</sub>×Y ears×male</i>					0.0334	0.0102	3.33	-0.001*
<i>ATLQL<sub>stan</sub>×Y ears×age</i>					-0.0027	0.0004	-0.73	-0.465
<i>ATLQL<sub>stan</sub>×Y ears×Moderate</i>					-0.0335	0.0107	-3.14	0.002*

\*Significant at the 5% level.

We find that the chi-squared tests of differences for a model including the dummies and their interactions were statistically significant (e.g. non-iterative sequential  $\chi^2(5) = 39.07$  Prob >  $\chi^2 = 0.000$ ). In this model we found that attitudes continue to affect choices when three-way interactions are included.