A tutorial on setting up a reproducible workflow in R and R Studio with Quarto

Supplementary Materials

# Appendix A: Supplementary Materials

## Model formulas

The regression formula for the full model (model b2):

*Note:* rt = reaction time (ms); stimulus = blah; compatibility = blah; pid = subject/participant identifier.

## A supplementary table

Here, I just reproduce the table from the main manuscript to save time making anything else. But of course in a real paper, you wouldn’t do this, you’d just show whatever you needed. It is labelled differently to reflect that it is a supplementary table ([Table 1](#tbl-s1)).

|  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Table 1: Supplementary table created with tinytable::tt()   | term | value | .lower | .upper | | --- | --- | --- | --- | | intercept | 613.9 | 593.62 | 635 | | stimulus | 6.4 | 0.77 | 12 | | Note. This is a footnote. | | | | |

## A supplementary figure

Let’s take a look at the mixing of chains in the model. This is a useful model diagnostic check to see that the model built ok ([Figure 1](#fig-s1)).

|  |
| --- |
| Figure 1: Caterpillar plots showing the mixing of chains across parameters in model b2. |