

Homework 06: Instructions

Summarizing Data

1. Using either the `gng_long.Rds` or the `gng_long_cleaned.Rds` data files (your own or one I shared with you), determine how many go and no-go trials there are for each individual at each wave. Your data frame would look something like the one below.

	<code>id_subject</code>	<code>wave</code>	<code>target</code>	<code>total_trials</code>
	<code><chr></code>	<code><dbl></code>	<code><chr></code>	<code><dbl></code>
1	FAQAAW	1	Go	?
2	FAQAAW	1	No-Go	?
3	FAQAAW	2	Go	?
4	FAQAAW	2	No-Go	?
5	FAQAAW	3	Go	?
6	FAQAAW	3	No-Go	?

2. Take the following `list()` and edit it so that your summary table will include the mean, median, and standard deviation. Make sure your functions can handle NA values.

```
summarize_across <- list(  
  mean = ~mean(.x, na.rm = TRUE) # or mean(na.omit(.x))  
)
```

2. Using the `gng_long_cleaned.Rds` data, aggregate your data by participant, wave, and trial type (e.g., go vs. no-go trials). Then generate a summary data frame containing the mean, median, and standard deviation across `accuracy` and `rt` so that you have the summary statistics for each participant, at each wave, and for both go and no-go trials. Remove the grouping structure so that if you assign it to an object or write the data frame, it is not grouped.
3. Using the `sspan_long_cleaned.Rds` data, generate a summary data frame containing the mean, median, and standard deviation across `sspan` and `totalcorrectsquares` by wave.