

# 01: Clean Data

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## Setup

### Load Libraries

```
library(readxl)
library(dplyr)
```

### Set File Paths

```
root <- "/Users/pei-chin/Research/Behavioral Science and Marketing_data_task"

data_raw      <- file.path(root, "data_raw")
data_processed <- file.path(root, "data_processed")
data_clean    <- file.path(root, "data_clean")
scripts       <- file.path(root, "scripts")
output        <- file.path(root, "output")
figures       <- file.path(output, "figures")
```

### Load Processed Data

```
load(file.path(data_processed, "processed_data.RData"))
```

## Clean Data

Sanity checks have been completed and documented in the script 00\_Appendix: Data Quality Checks. This section focuses on dropping variables unnecessary for analysis.

## Remove Irrelevant Columns

These column indices correspond to metadata, system-generated fields, and other variables that are not relevant for analysis, so they are removed here.

```
df_data <- processed_data %>%  
  select(-c(1:6), -c(8:17), -c(19:26), -c(62, 64, 65, 68, 69, 70))
```

## Apply Inclusion Criteria and Final Variable Selection

```
clean_data <- df_data %>%  
  # Apply inclusion criteria  
  filter(Finished == "TRUE", consent == "AGREE", passedattn == "yes") %>%  
  # Remove variables not needed for analysis  
  # Some variables were not shown to participants in the survey interface,  
  # so they are removed here to avoid ambiguity in interpretation  
  select(  
    -real_imaginary, -initials_box, -describe, -real_imaginary, -describe,  
    -feelings_exp, -attention_1, -attention_1_TEXT, -attention_2, -attention_2_TEXT,  
    -attention_3, -attention_3_TEXT, -target_sex_3_TEXT, -sex_3_TEXT, -comments,  
    -consent, -Finished, -passedattn, -initials, -initiator_type  
  ) %>%  
  # Create participant ID  
  
  mutate(id = row_number()) %>%  
  # Create a high vs. low self-blame grouping variable using a median split at 50  
  # This step distinguishes participants by their tendency toward self-blame  
  # While dichotomizing a continuous variable has methodological limitations,  
  # a binary grouping (0/1) is used here for simplicity and time constraints  
  mutate(high_blame = if_else(blame_1 > 50, 1, 0)) %>%  
  # Move ID to the first column for easier inspection and reference  
  relocate(id)
```

## Save and Verify

### Save Cleaned Dataset

```
save(  
  clean_data,  
  file = file.path(data_clean, "clean_data.RData")  
)
```

### Quick Check

```
dim(clean_data)
```

```
## [1] 45 24
```