Supplement 5

Experiment 1 – Descriptive plots of the SPARS stimulus-response relationship

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Contents

This script is part 1 of our analysis of the stimulus-response characteristics of the SPARS. This script generates exploratory plots of the relationship between stimulus intensity and SPARS rating.

Source URL: https://github.com/kamermanpr/SPARS/tree/supplementary_pdfs

Modelling of the stimulus-response relationship is described in "outputs/supplement_6.pdf", the diagnostics on the final linear mixed model are described in "outputs/supplement_7.pdf", the stability of the model is described in "outputs/supplement_8.pdf", the sensitivity of the scale to changes in stimulus intensity are described in "outputs/supplement_9.pdf", and the variance in ratings at each stimulus intensity is described in "outputs/supplement_10.pdf".

Import and inspect data

```
## $ trial number
           <dbl> 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 1...
## $ intensity
           <dbl> 3.75, 1.50, 3.25, 1.50, 3.00, 2.75, 1.00, 2....
           <chr> "3.75", "1.50", "3.25", "1.50", "3.00", "2.7...
## $ intensity char
           <dbl> -10, -40, -10, -25, -20, -25, -40, 2, -40, -...
## $ rating
## $ rating_positive
           <dbl> 40, 10, 40, 25, 30, 25, 10, 52, 10, 40, 54, ...
## $ EDA
           <dbl> 18315.239, 13904.177, 11543.449, 20542.834, ...
## $ age
           ## $ sex
           ## $ panas_positive
           ## $ panas negative
## $ dass42 anxiety
           ## $ dass42 stress
## $ pcs_rumination
           ## $ pcs_helplessness
```

Clean and transform data

We performed a basic clean-up of the data, and then calculated *Tukey trimean* at each stimulus intensity for each participant (participant average), and finally the *median* of the trimeans at each stimulus intensity across participants (group average).

```
#
                                           #
#
                    Clean
                                           #
data %<>%
 # Select required columns
 select(PID, block, block order, trial number, intensity, intensity char, rating)
#
                                           #
#
            Calculate 'Tukey trimean'
                                           #
#
                                           #
# Define tri.mean function
tri.mean <- function(x) {</pre>
 # Calculate quantiles
 q1 <- quantile(x, probs = 0.25, na.rm = TRUE)[[1]]
 q2 <- median(x, na.rm = TRUE)
 q3 \leftarrow quantile(x, probs = 0.75, na.rm = TRUE)[[1]]
 # Calculate trimean
 tm \leftarrow (q2 + ((q1 + q3) / 2)) / 2
 # Convert to integer
 tm <- as.integer(round(tm))</pre>
 return(tm)
}
```

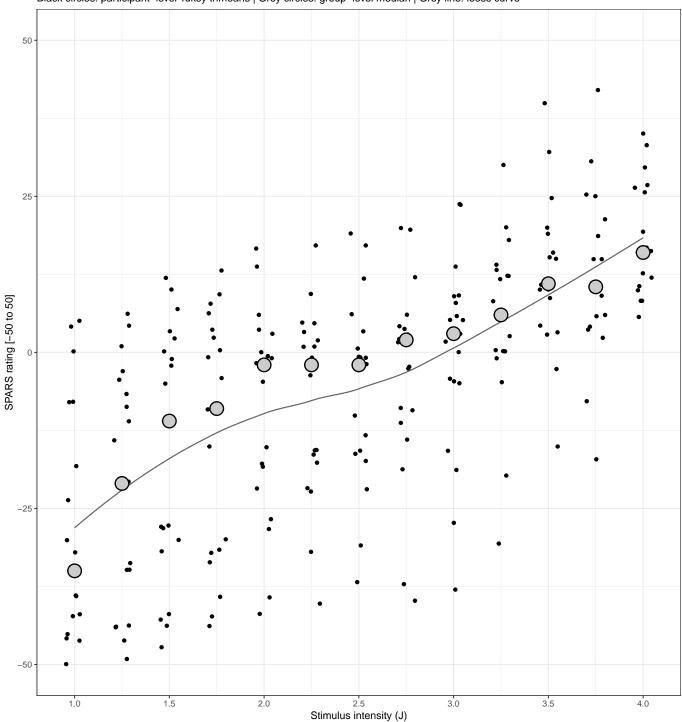
```
#
                                           #
#
                                           #
               Generate core data
# Calculate the participant average
data tm <- data %>%
 group_by(PID, intensity) %>%
 summarise(tri mean = tri.mean(rating)) %>%
 ungroup()
# Calculate the group average
data_group <- data_tm %>%
 group_by(intensity) %>%
 summarise(median = median(tri mean)) %>%
 ungroup()
```

Exploratory plots

Group-level stimulus response curve

```
# Plot
data tm %>%
  ggplot(data = .) +
  aes(x = intensity,
      y = tri_mean) +
 geom_point(position = position_jitter(width = 0.05)) +
  geom_smooth(method = 'loess',
              se = FALSE,
              colour = '#656565',
              size = 0.6) +
  geom_point(data = data_group,
             aes(y = median),
             shape = 21,
             size = 6,
             stroke = 1,
             fill = '#CCCCCC') +
  labs(title = 'Group-level stimulus-response plot',
       subtitle = 'Black circles: participant-level Tukey trimeans | Grey circles: group-l
       x = 'Stimulus intensity (J)',
       y = 'SPARS rating [-50 to 50]') +
  scale_y_continuous(limits = c(-50, 50)) +
  scale_x_continuous(breaks = seq(from = 1, to = 4, by = 0.5))
```

Group-level stimulus-response plot
Black circles: participant-level Tukey trimeans | Grey circles: group-level median | Grey line: loess curve



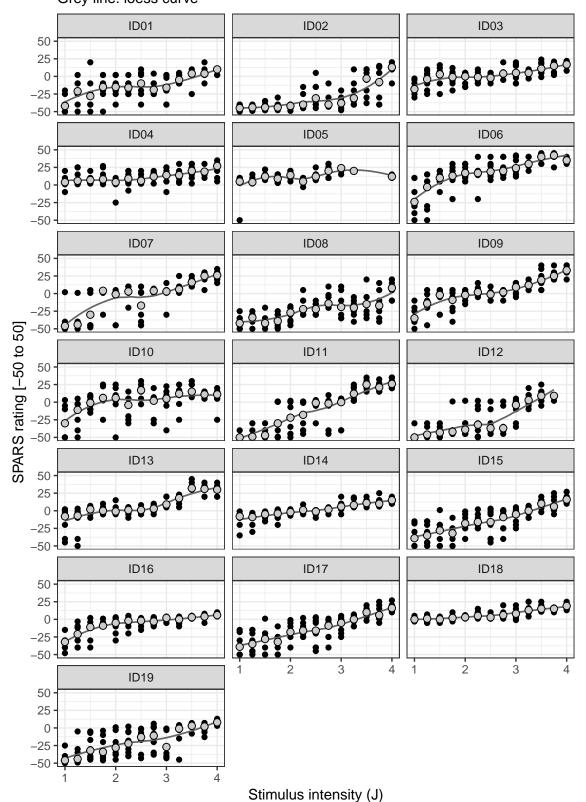
Participant-level stimulus response curves

All trials

```
y = rating) +
geom_point() +
geom_smooth(method = 'loess',
            se = FALSE,
            colour = '#656565',
            size = 0.6) +
geom_point(data = data_tm,
           aes(y = tri_mean),
           shape = 21,
           size = 2.25,
           fill = '#CCCCCC') +
labs(title = 'Participant-level stimulus-response plot',
     subtitle = 'Black circles: individual experimental blocks | Grey circles: Tukey tri
     x = 'Stimulus intensity (J)',
     y = 'SPARS rating [-50 to 50]') +
scale_y_continuous(limits = c(-50, 50)) +
facet_wrap(~ PID, ncol = 3) +
theme_bw()
```

Participant-level stimulus-response plot

Black circles: individual experimental blocks | Grey circles: Tukey trimean | Grey line: loess curve

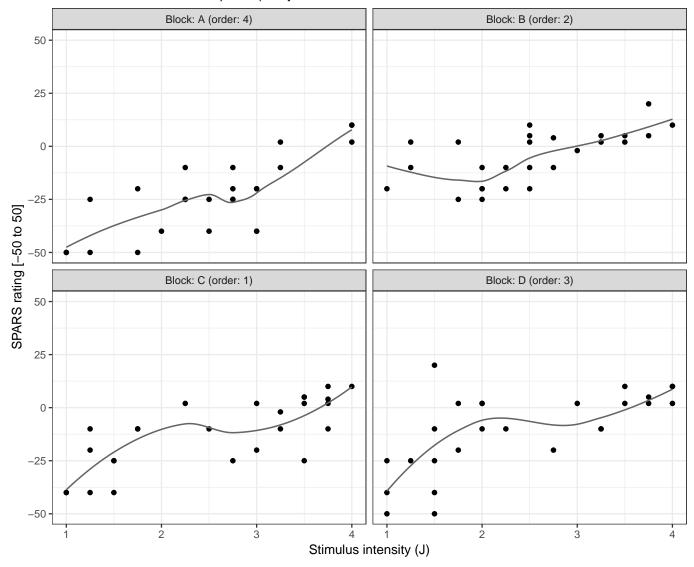


Trials by experimental block

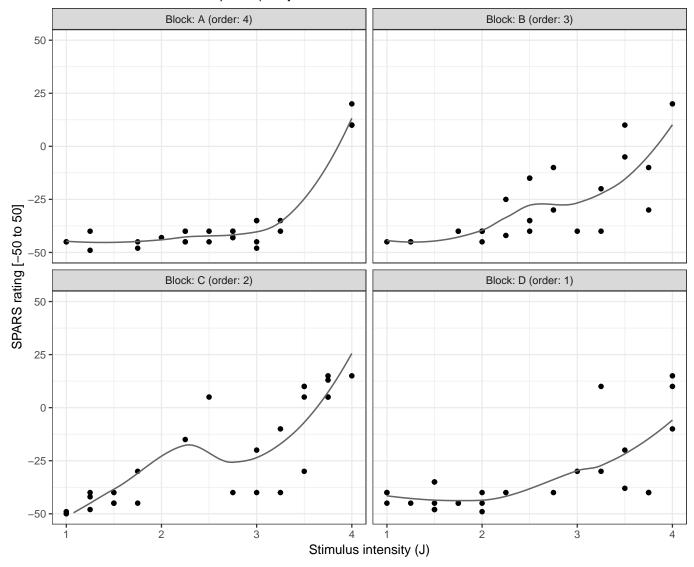
```
# Process data
data_block <- data %>%
    # Rename blocks
mutate(block = sprintf('Block: %s (order: %i)', block, block_order)) %>%
    # Nest by PID
```

```
group_by(PID) %>%
 nest() %>%
  # Generate plots
 mutate(plots = map2(.x = data,
                      .y = unique(PID),
                      ~ ggplot(data = .x) +
                        aes(x = intensity,
                            y = rating) +
                        geom_point() +
                        geom_smooth(method = 'loess',
                                    se = FALSE,
                                    colour = '#656565',
                                    size = 0.6) +
                        labs(title = paste(.y, ': Participant-level stimulus-response plot
                             subtitle = 'Black circles: individual data points | Grey line
                             x = 'Stimulus intensity (J)',
                             y = 'SPARS rating [-50 to 50]') +
                        scale_y_continuous(limits = c(-50, 50)) +
                        facet_wrap(~ block, ncol = 2)))
# Print plots
walk(.x = data_block$plots, ~ print(.x))
```

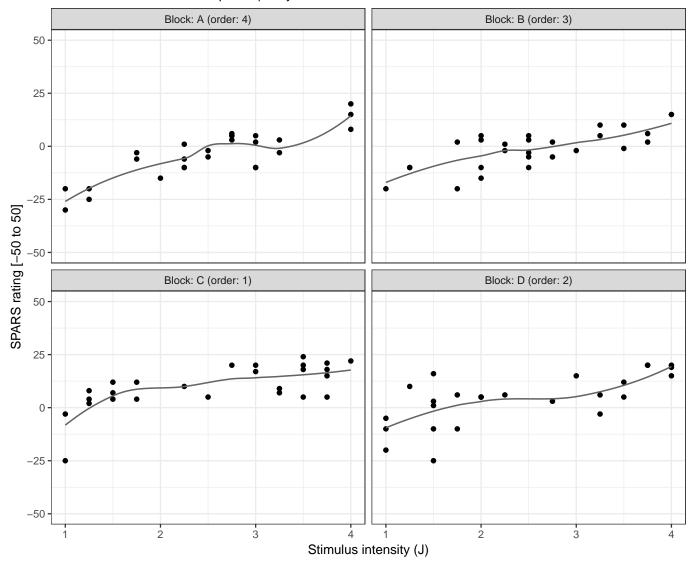
ID01 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



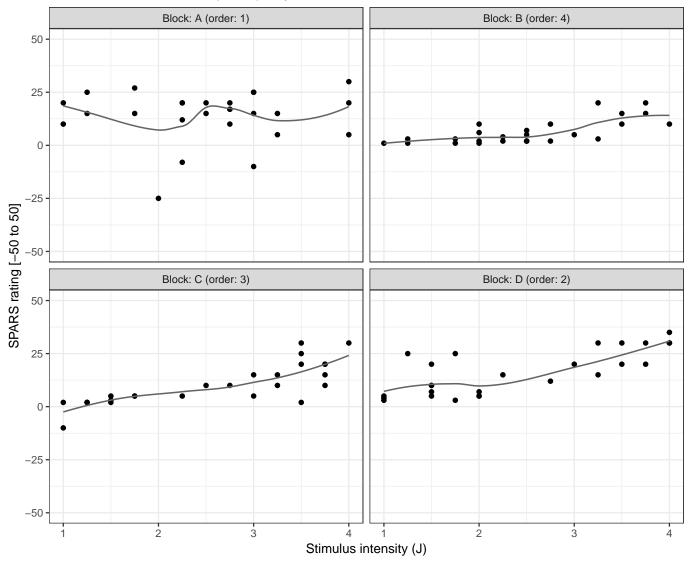
ID02 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



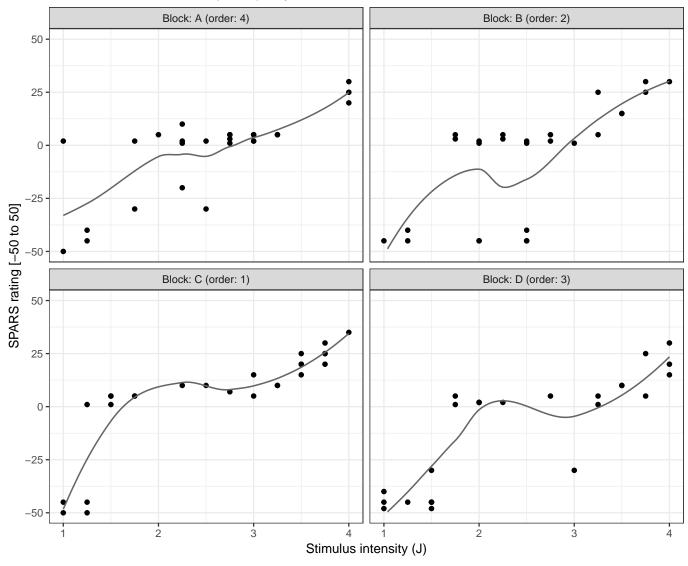
ID03 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



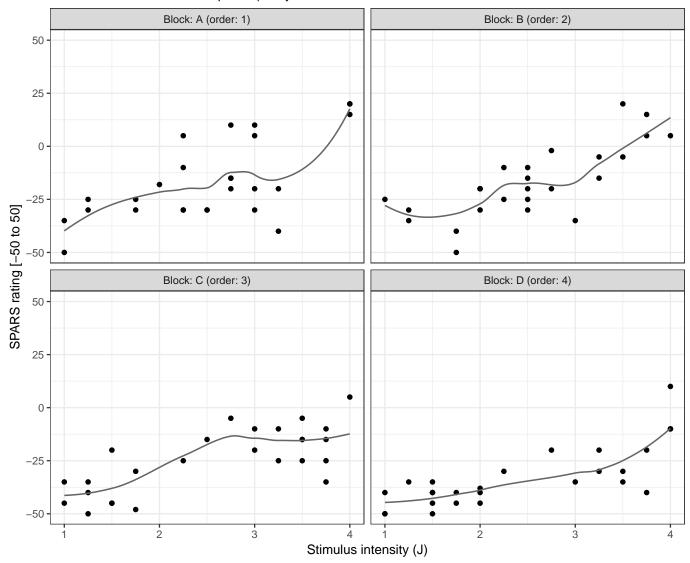
ID04 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



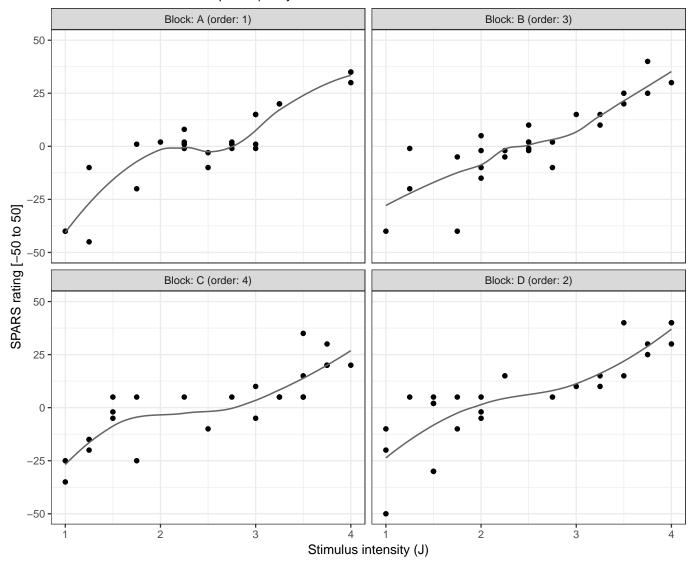
ID07 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



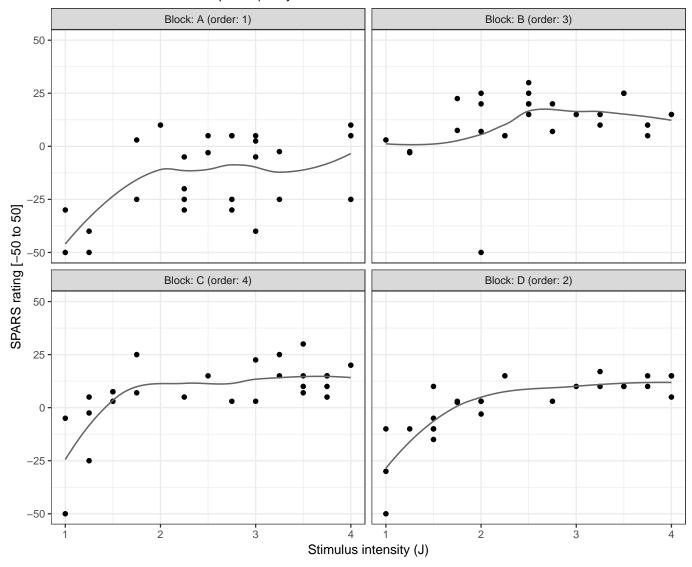
ID08 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



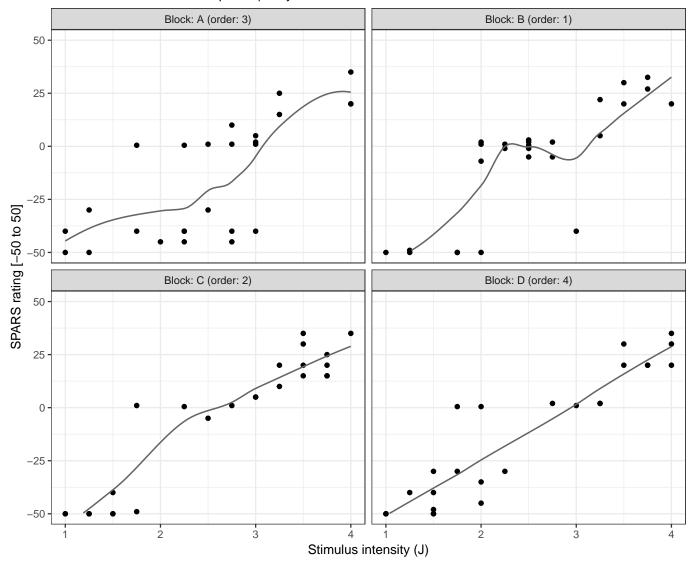
ID09 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



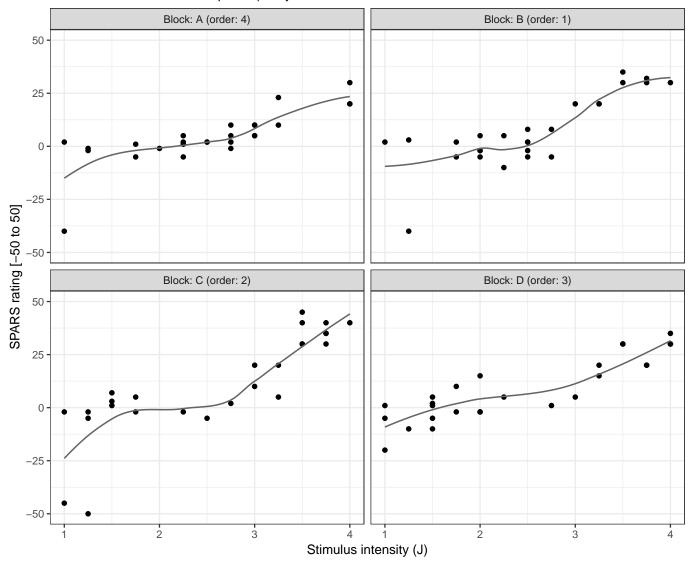
ID10 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



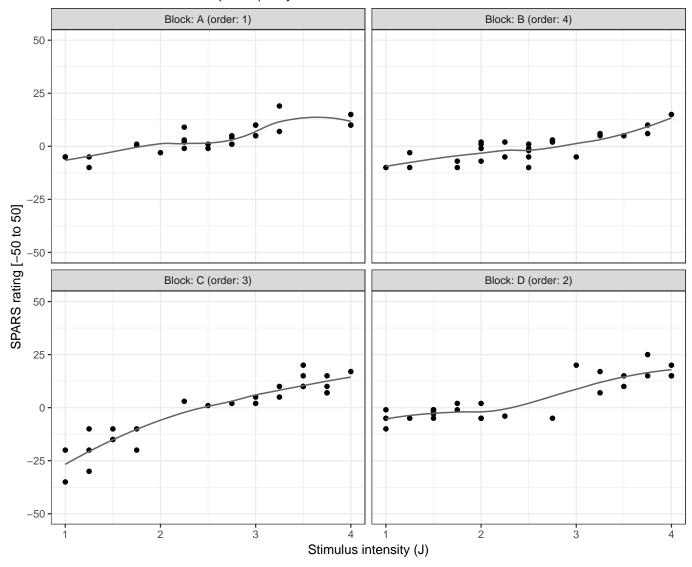
ID11 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



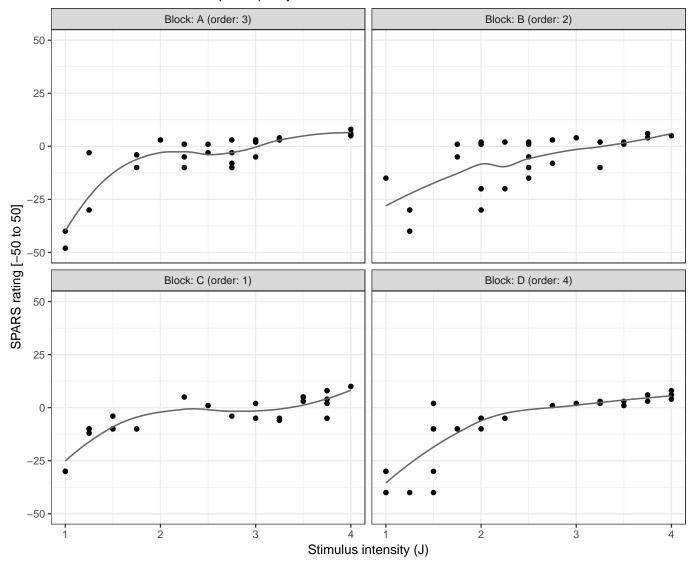
ID13 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



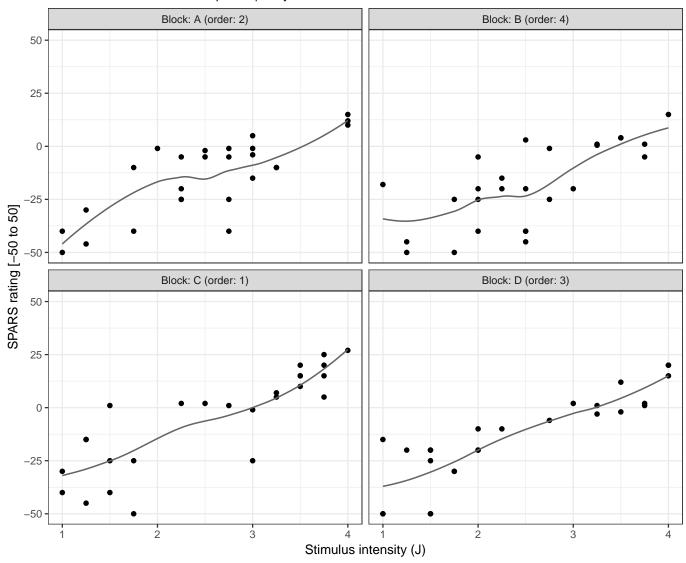
ID14 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



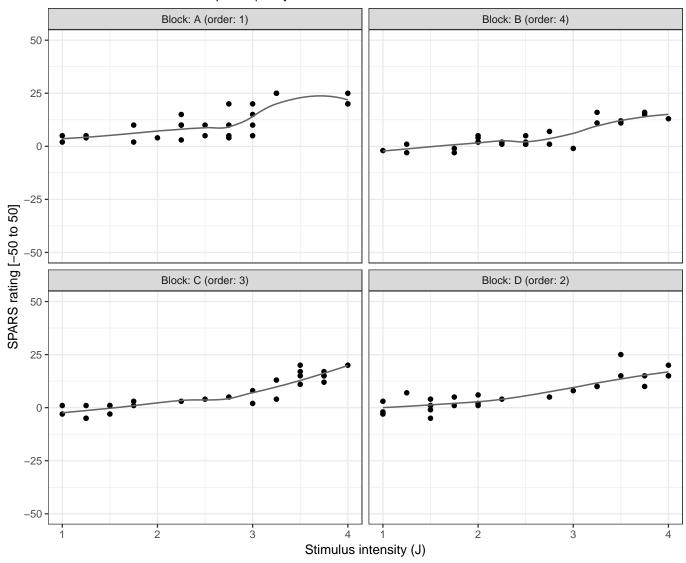
ID16 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



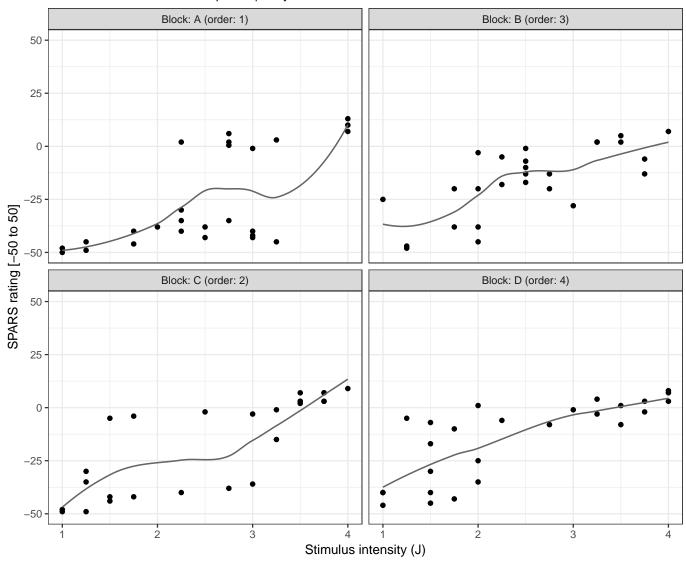
ID17 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



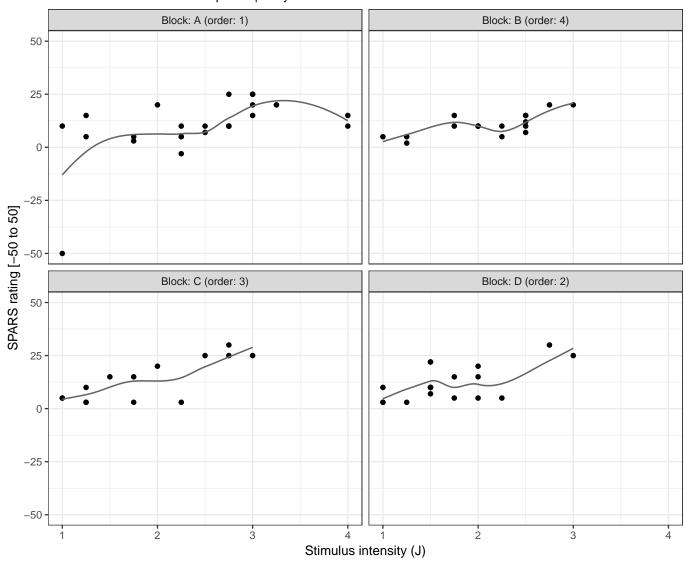
ID18 : Participant-level stimulus-response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



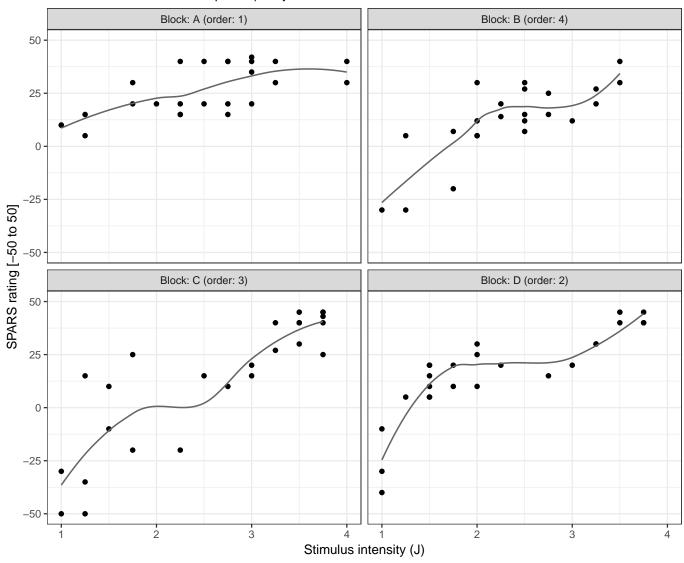
ID19 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



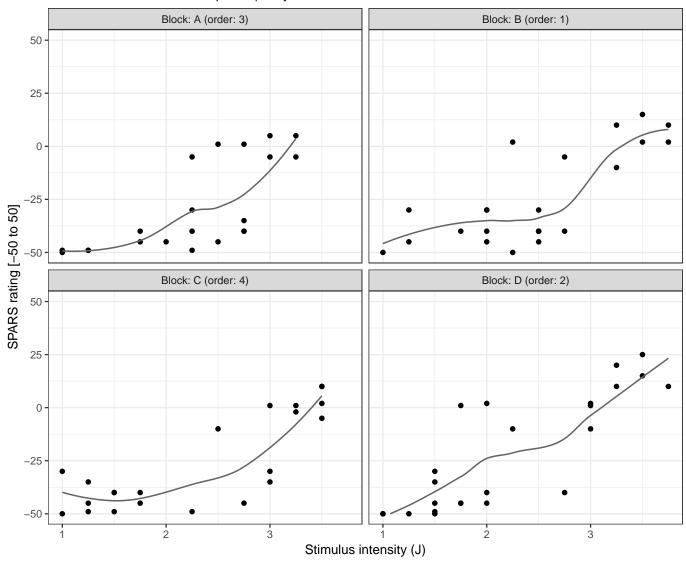
ID05 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



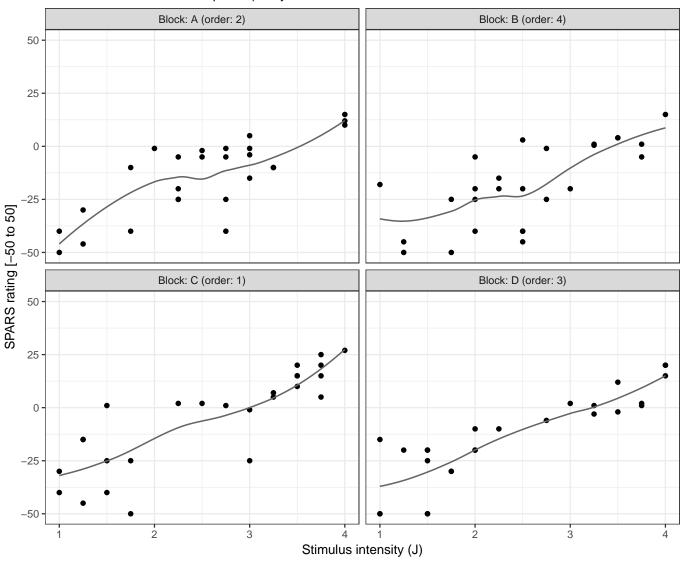
ID06 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



ID12 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



ID15 : Participant–level stimulus–response plots conditioned on experimental block Black circles: individual data points | Grey line: loess curve



Session information

attached base packages:

sessionInfo()

[1] stats

```
## R version 3.5.1 (2018-07-02)
## Platform: x86_64-apple-darwin15.6.0 (64-bit)
## Running under: macOS 10.14
##
## Matrix products: default
## BLAS: /Library/Frameworks/R.framework/Versions/3.5/Resources/lib/libRblas.0.dylib
## LAPACK: /Library/Frameworks/R.framework/Versions/3.5/Resources/lib/libRlapack.dylib
##
## locale:
## [1] en_GB.UTF-8/en_GB.UTF-8/en_GB.UTF-8/C/en_GB.UTF-8/en_GB.UTF-8
##
```

datasets methods

base

graphics grDevices utils

```
##
## other attached packages:
    [1] bindrcpp_0.2.2 patchwork_0.0.1 forcats_0.3.0
                                                         stringr_1.3.1
##
##
    [5] dplyr 0.7.6
                        purrr 0.2.5
                                         readr 1.1.1
                                                         tidyr 0.8.1
##
    [9] tibble_1.4.2
                        ggplot2_3.0.0
                                         tidyverse_1.2.1 magrittr_1.5
##
## loaded via a namespace (and not attached):
    [1] Rcpp_0.12.19
                         cellranger_1.1.0 pillar_1.3.0
                                                            compiler_3.5.1
##
##
    [5] plyr 1.8.4
                         bindr 0.1.1
                                           tools 3.5.1
                                                            digest 0.6.17
##
    [9] lubridate 1.7.4
                         jsonlite 1.5
                                           evaluate 0.11
                                                            nlme 3.1-137
## [13] gtable 0.2.0
                         lattice 0.20-35
                                           pkgconfig_2.0.2
                                                            rlang_0.2.2
##
   [17] cli_1.0.1
                         rstudioapi_0.8
                                           yam1_2.2.0
                                                            haven_1.1.2
## [21] withr 2.1.2
                                           httr 1.3.1
                                                            knitr 1.20
                         xml2 1.2.0
## [25] hms 0.4.2
                         rprojroot 1.3-2
                                           grid_3.5.1
                                                            tidyselect 0.2.4
## [29] glue_1.3.0
                         R6_2.2.2
                                           readxl_1.1.0
                                                            rmarkdown_1.10
## [33] modelr_0.1.2
                         backports_1.1.2
                                           scales_1.0.0
                                                            htmltools_0.3.6
## [37] rvest 0.3.2
                         assertthat 0.2.0 colorspace 1.3-2 labeling 0.3
## [41] stringi 1.2.4
                         lazyeval 0.2.1
                                           munsell 0.5.0
                                                            broom 0.5.0
## [45] crayon_1.3.4
```