

Supplement 4

Experiment 1 – Effect of stimulus order and blocking on SPARS rating

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The experimental protocol called for participants to be exposed to 13 stimuli, evenly spaced at 0.25J intervals over the range 1.00J to 4.00J. Each stimulus intensity was applied 8 times, giving a total of 104 exposures (trials). To prevent learning effects, the 104 trials were randomised across 4 experimental blocks (26 trials per block).

Despite using a randomized block approach, we still wanted to assess whether there were any:

- Trial order effects (does the intensity of the preceding stimulus affect the rating of a stimulus?)
 - Block order effects (does the rating of a given stimulus intensity change across experimental blocks?)
-

Import and inspect data

```
# Import
data <- read_rds('./data-cleaned/SPARS_A.rds')

# Inspect
glimpse(data)
```

```
## Observations: 1,927
## Variables: 19
## $ PID          <chr> "ID01", "ID01", "ID01", "ID01", "ID01", "ID0...
## $ block        <chr> "A", "A", "A", "A", "A", "A", "A", "A", "A", ...
## $ block_order  <dbl> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, ...
## $ 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....
## $ intensity_char <chr> "3.75", "1.50", "3.25", "1.50", "3.00", "2.7...
## $ rating       <dbl> -10, -40, -10, -25, -20, -25, -40, 2, -40, -...
```

```
## $ 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 <dbl> 21, 21, 21, 21, 21, 21, 21, 21, 21, 21, 21, ...
## $ sex <dbl> 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, ...
## $ panas_positive <dbl> 36, 36, 36, 36, 36, 36, 36, 36, 36, 36, 36, ...
## $ panas_negative <dbl> 10, 10, 10, 10, 10, 10, 10, 10, 10, 10, 10, ...
## $ dass42_depression <dbl> 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, ...
## $ dass42_anxiety <dbl> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, ...
## $ dass42_stress <dbl> 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, ...
## $ pcs_magnification <dbl> 6, 6, 6, 6, 6, 6, 6, 6, 6, 6, 6, ...
## $ pcs_rumination <dbl> 11, 11, 11, 11, 11, 11, 11, 11, 11, 11, 11, ...
## $ pcs_helplessness <dbl> 10, 10, 10, 10, 10, 10, 10, 10, 10, 10, 10, ...
```

Clean data and process

```
#####
#                                                                    #
#                               Clean                                #
#                                                                    #
#####
data %<>%
  # Select required columns
  select(PID, block, block_order, trial_number,
          intensity, rating, rating_positive) %>%
  # Format block and trial number type
  mutate(block_order = as.integer(block_order),
          trial_number = as.integer(trial_number)) %>%
  # 'Back-up' intensity data as a continuous variable before factorizing
  mutate(intensity_cont = intensity) %>%
  # Format intensity data as an ordered factor
  mutate(intensity = sprintf('%s%.2f%s', 'Stimulus: ', intensity, 'J'),
          intensity = factor(intensity,
                             levels = paste0('Stimulus: ',
                                                as.character(format(seq(from = 1,
                                                                           to = 4,
                                                                           by = 0.25),
                                                                           digits = 3))),
                             'J'),
          ordered = TRUE))

#####
#                                                                    #
#                               Lagged stimulus intensity data        #
#                               (lag 1: preceding stimulus)           #
#                                                                    #
#####
data_lag <- data %>%
  # Drop unneeded columns (block_order and positive_intensity)
  select(-rating_positive) %>%
```

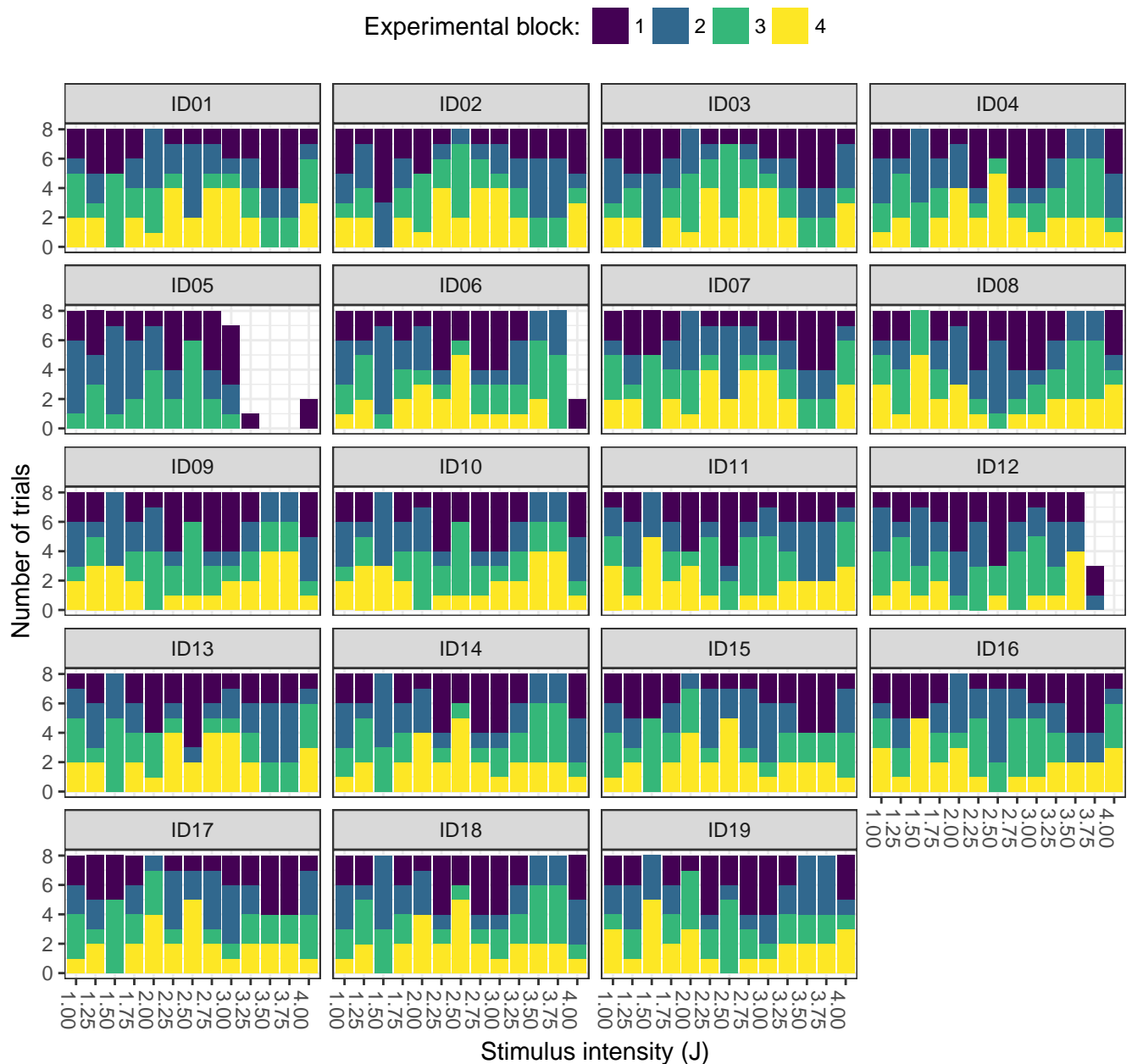
```
# Get lag-1 stimulus intensity by PID
group_by(PID) %>%
mutate(intensity_lag = lag(intensity_cont)) %>%
# Ungroup and remove incomplete cases created by lag function (i.e., first trial)
ungroup() %>%
filter(complete.cases(.))
```

Summary

The figure below summarises the exposure (number of trials) of each participant at each stimulus intensity.

```
data %>%
  ggplot(data = .) +
  aes(x = intensity,
      fill = factor(block_order)) +
  geom_bar() +
  scale_fill_viridis_d(name = 'Experimental block:') +
  facet_wrap(~PID, ncol = 4) +
  labs(title = 'Number of trials per participant at each stimulus intensity, stratified by
          y = 'Number of trials',
          x = 'Stimulus intensity (J)') +
  scale_x_discrete(labels = sprintf('%.2f', seq(from = 1, to = 4, by = 0.25))) +
  theme_bw() +
  theme(legend.position = 'top',
        axis.text.x = element_text(angle = -90))
```

Number of trials per participant at each stimulus intensity, stratified by experimental



Participants *ID05*, *ID06*, and *ID12* did not complete the trial per protocol.

Trial order effects

Question: Is the SPARS rating for a given stimulus intensity dependent on the intensity of the preceding stimulus?

The analysis includes data from experimental trials, ignoring experimental blocks. Because each stimulus intensity is repeated multiple times in each participant, a hierarchical analysis by study participant and stimulus intensity was conducted.

User-defined function

Function to calculate the *Tukey trimeans* for plots.

```

# Trimean
tri.mean <- function(x) {
  # Calculate quantiles
  q1 <- quantile(x, probs = 0.25, na.rm = TRUE)[[1]]
  q2 <- median(x, na.rm = TRUE)
  q3 <- quantile(x, probs = 0.75, na.rm = TRUE)[[1]]
  # Calculate trimean
  tm <- (q2 + ((q1 + q3) / 2)) / 2
  # Convert to integer
  tm <- as.integer(round(tm))
  return(tm)
}

```

Participant-level exploratory plots

Participant-level plots of SPARS rating against the intensity of the preceding stimulus, across the range of stimuli used (1.00J to 4.00J).

```

# Plot data
plot_data <- data_lag %>%
  # Nest by ID
  group_by(PID) %>%
  nest() %>%
  # Calculate trimean
  mutate(trimean = map(.x = data,
                        ~ group_by(.data = .x,
                                    intensity) %>%
                                    summarise(tm = tri.mean(rating))))

# Plot
plot_trial <- plot_data %>%
  mutate(plot = pmap(..1 = list(data, trimean, unique(PID)),
                     ~ ggplot(data = ..1) +
                       aes(x = intensity_lag,
                           y = rating) +
                       geom_point() +
                       geom_smooth(se = FALSE,
                                   colour = '#666666',
                                   size = 0.6,
                                   na.rm = TRUE) +
                       geom_hline(data = ..2,
                                   aes(yintercept = tm),
                                   colour = 'red',
                                   size = 0.6,
                                   linetype = 2) +
                       labs(title = paste0(..3, ': SPARS rating versus intensity of the pre
                                   subtitle = 'Red line (dashed): Tukey trimean of SPARS rating |
                                   y = 'SPARS rating [-50 to 50]',
                                   x = 'Intensity of previous stimulus (J)') +
                       scale_y_continuous(limits = c(-50, 50)) +
                       scale_x_continuous(limits = c(1, 4)) +

```

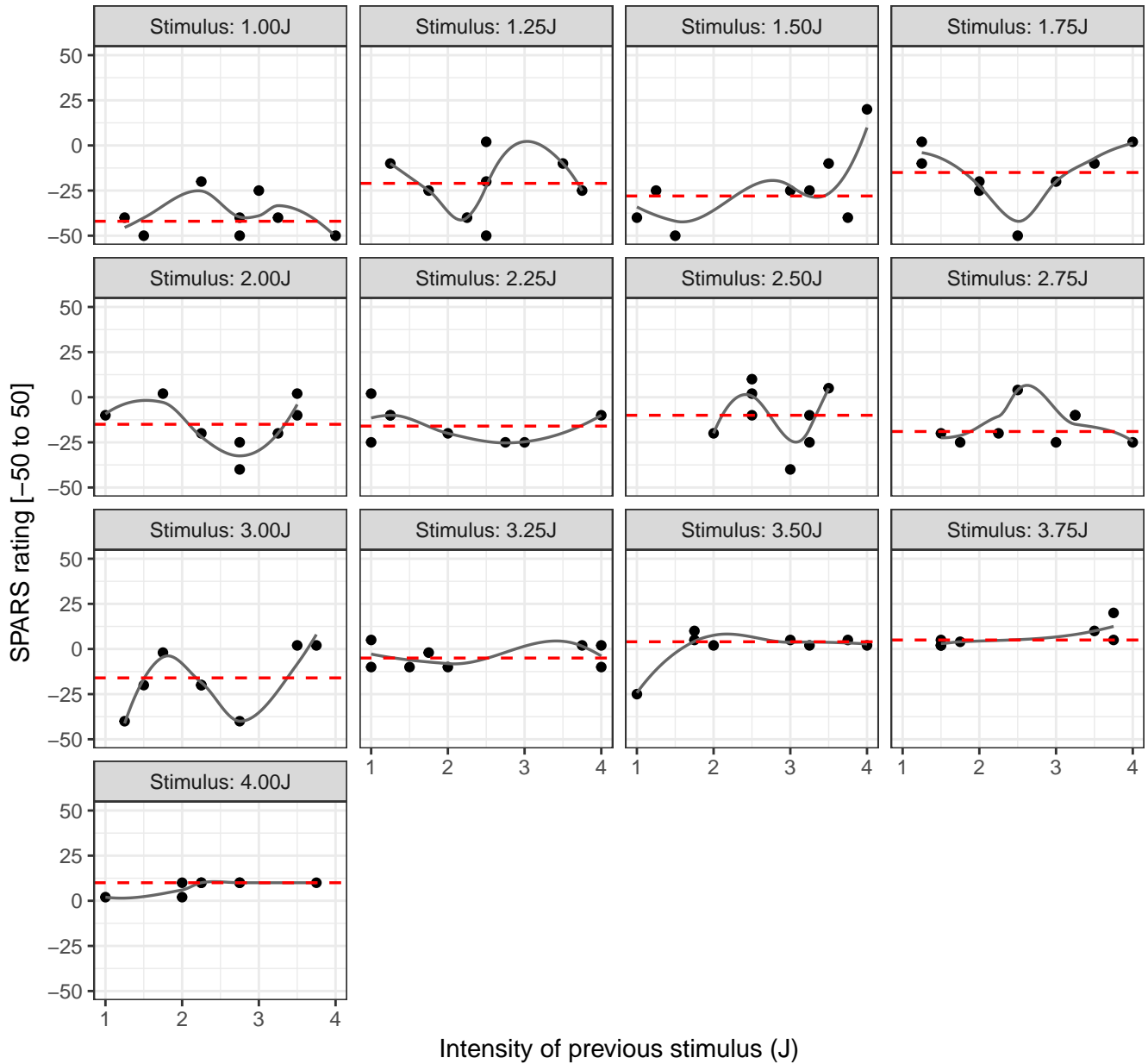
```
facet_wrap(~intensity,
           ncol = 4) +
theme_bw())
```

```
# Print plots
```

```
walk(.x = plot_trial$plot, ~ print(.x))
```

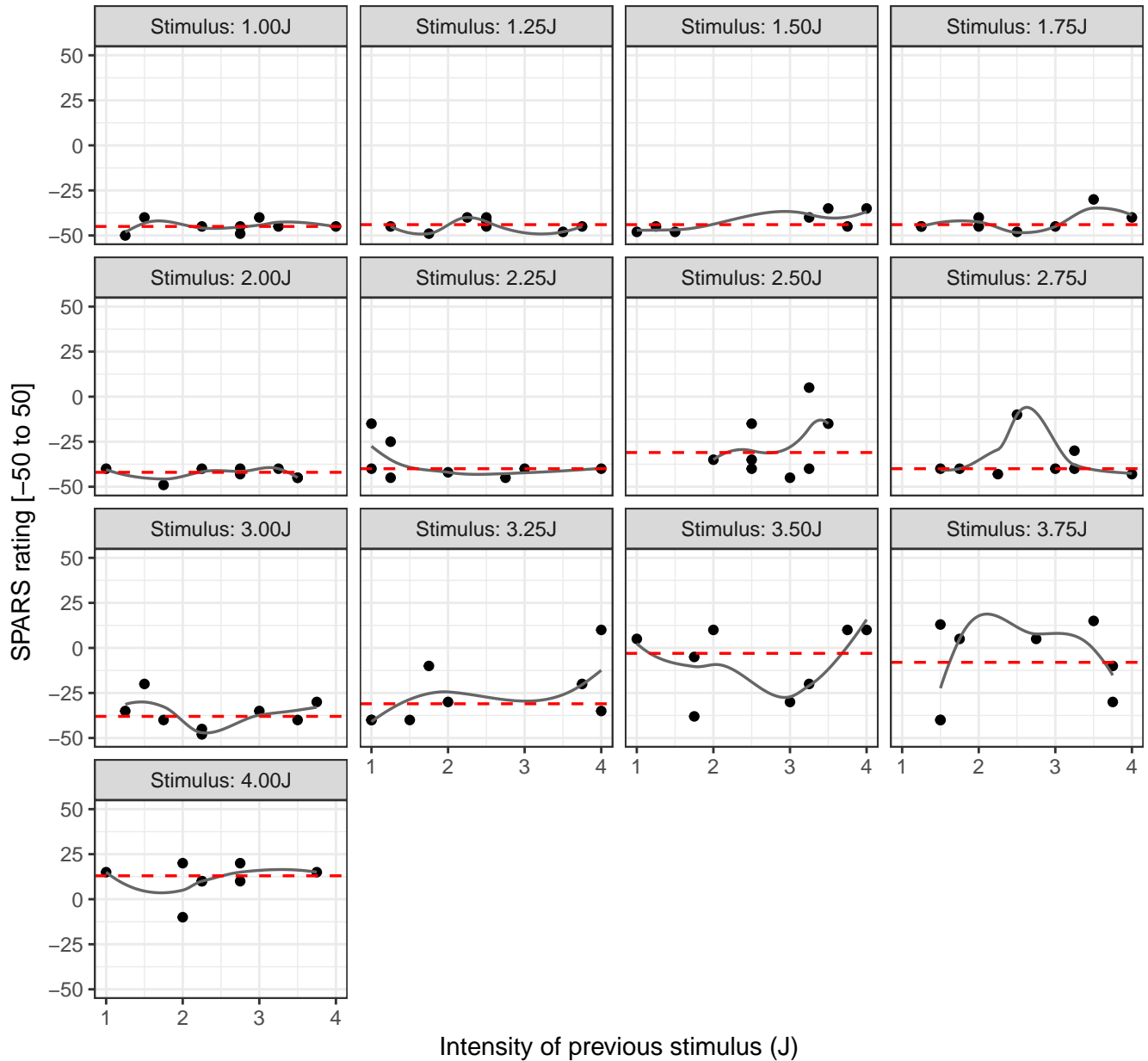
ID01: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



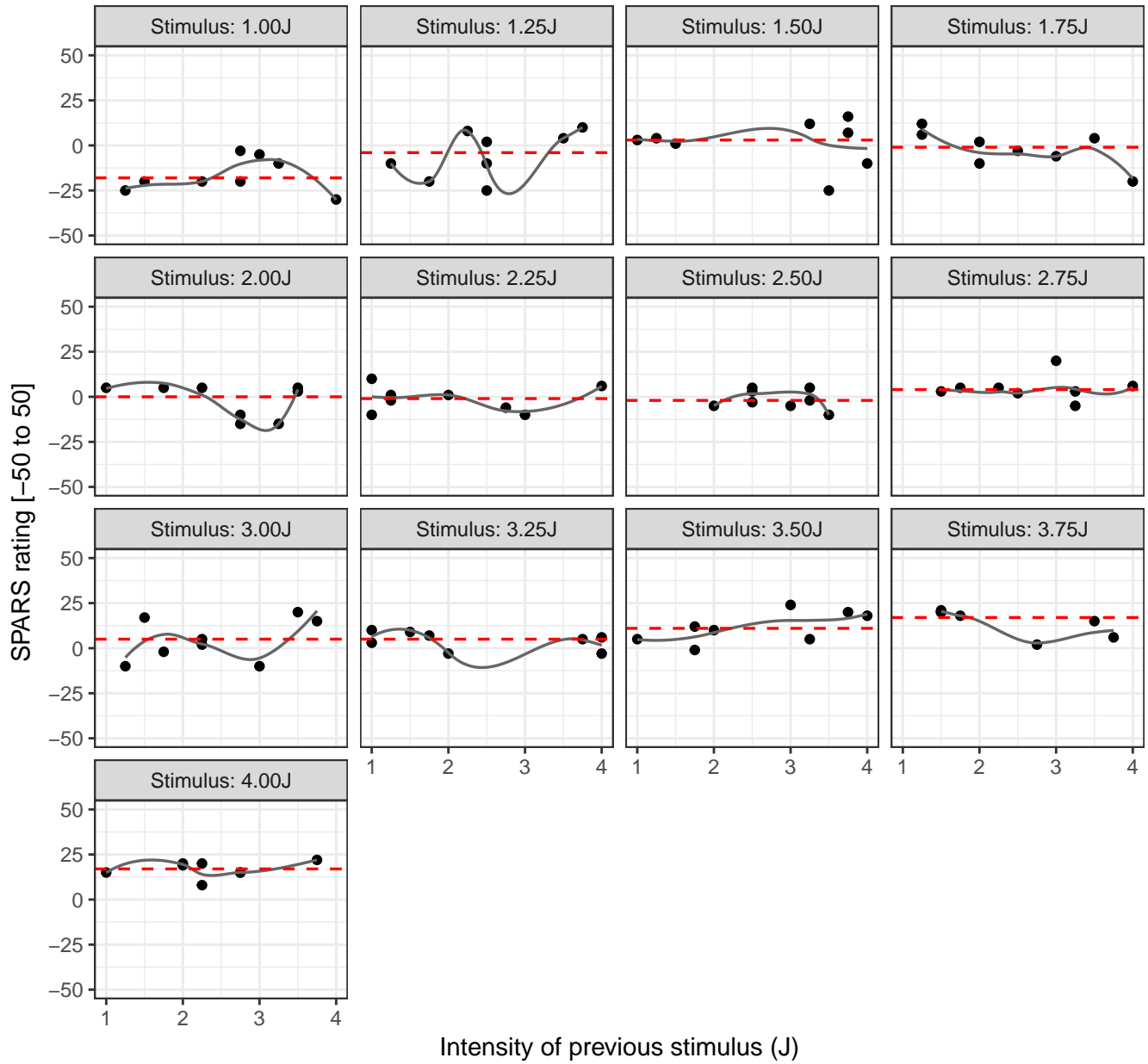
ID02: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



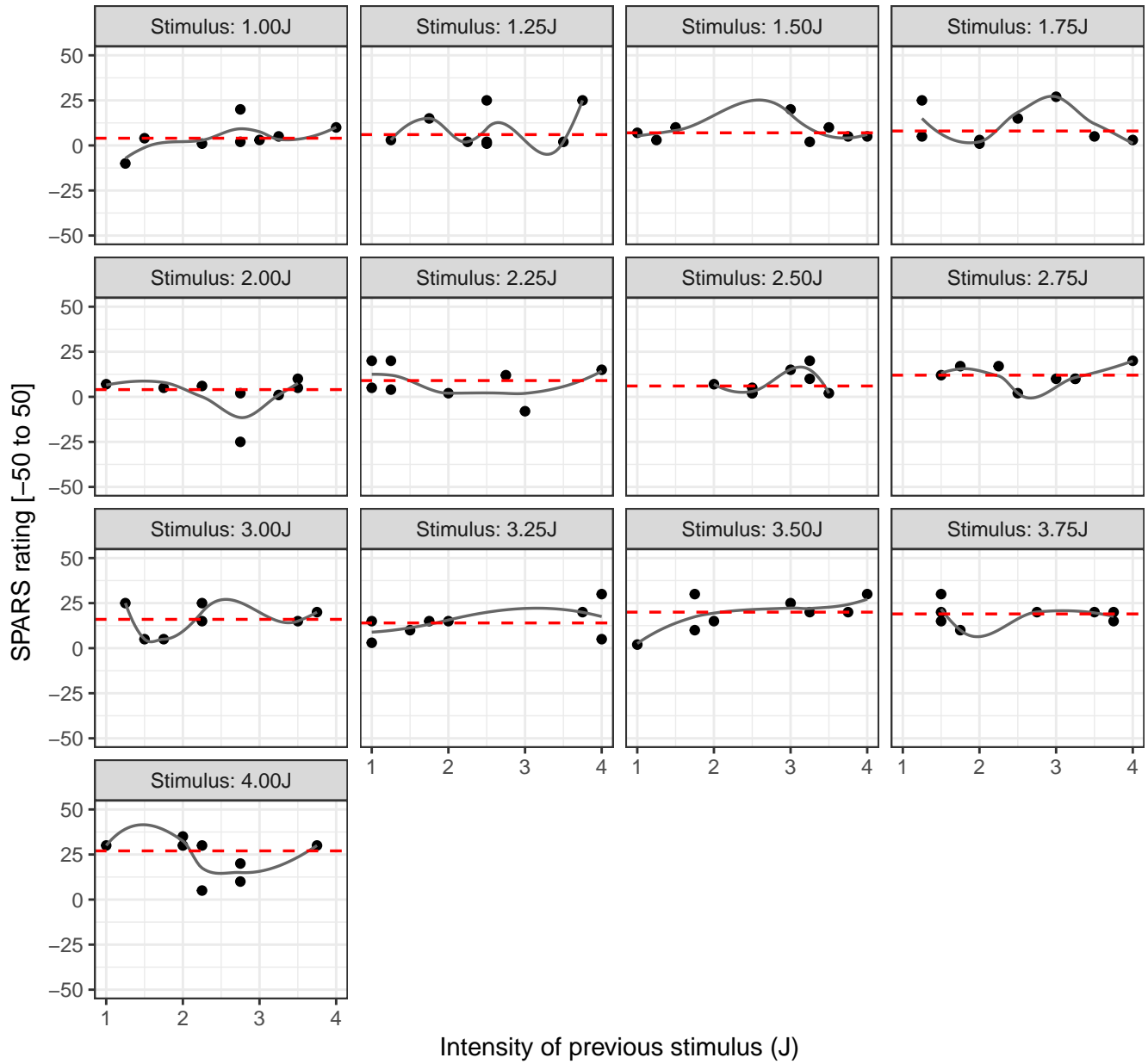
ID03: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



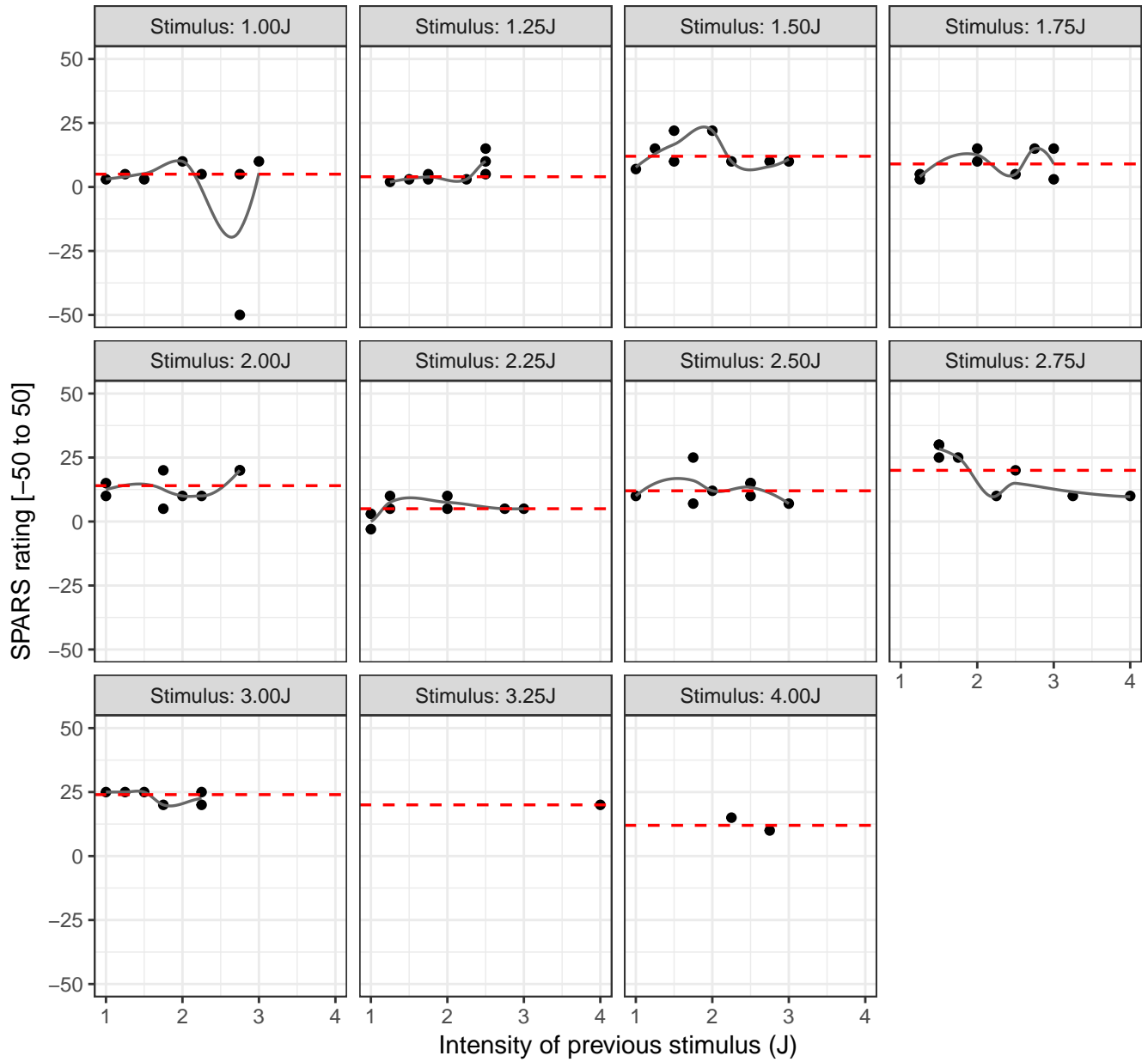
ID04: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



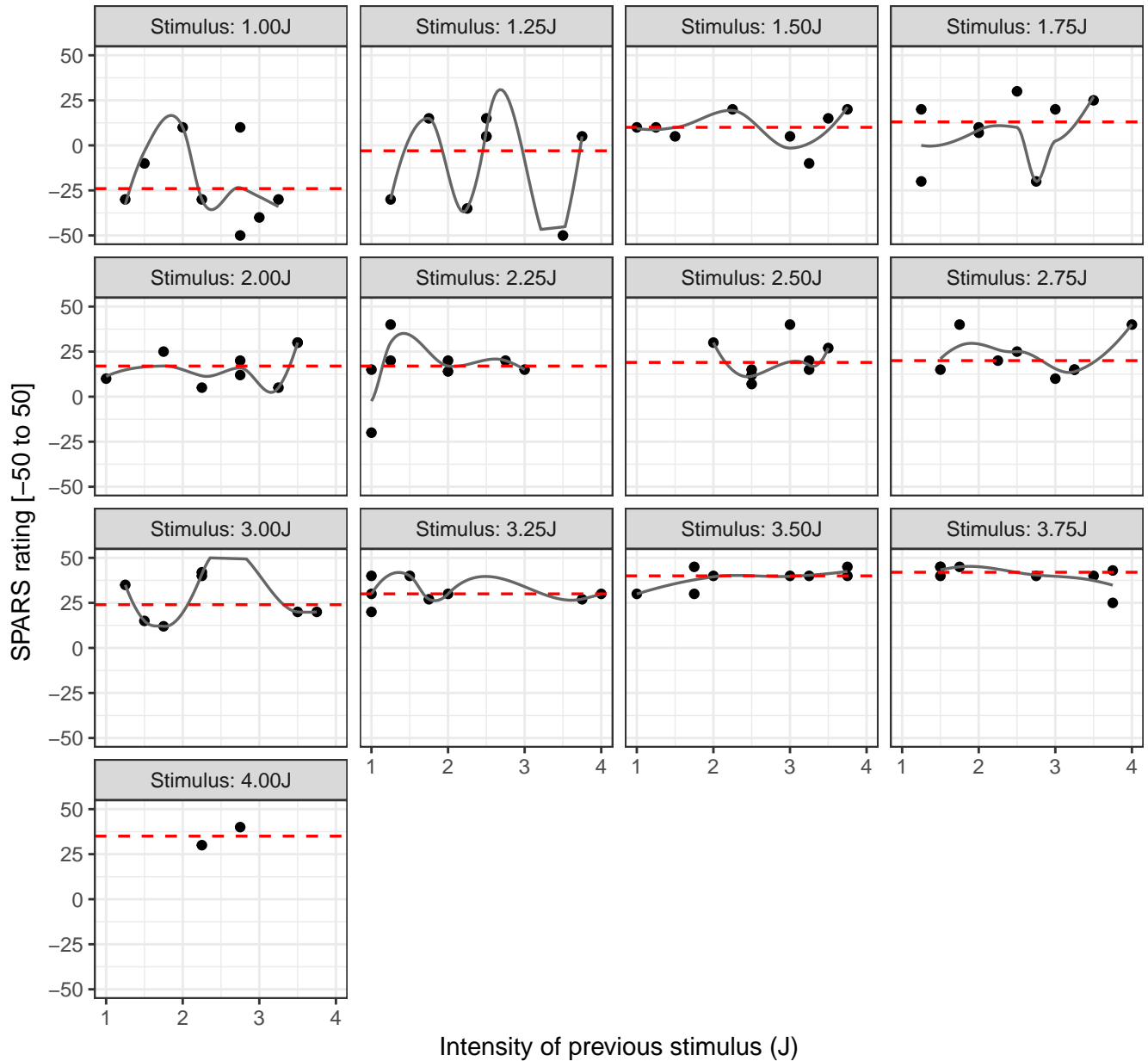
ID05: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



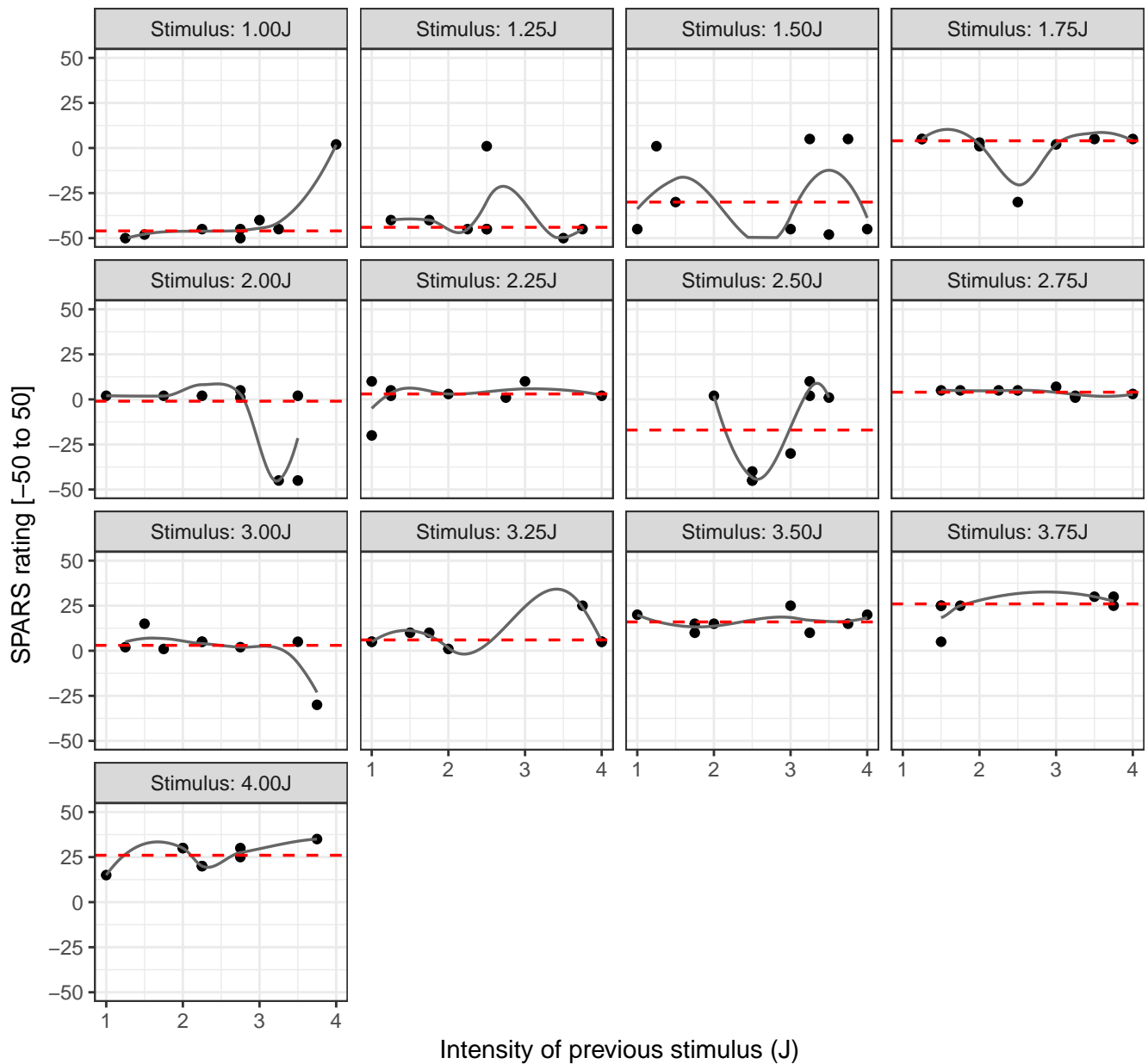
ID06: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



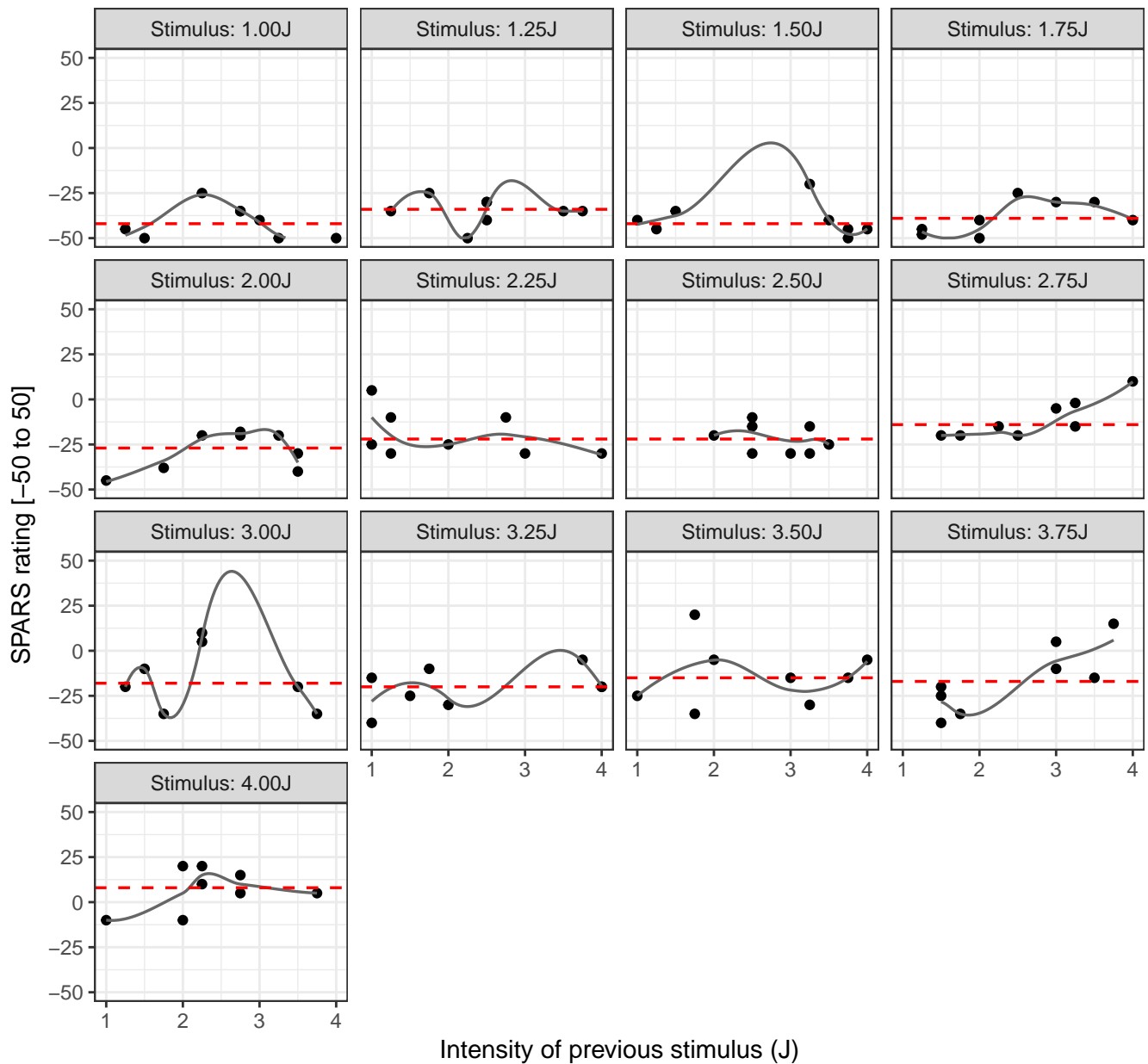
ID07: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



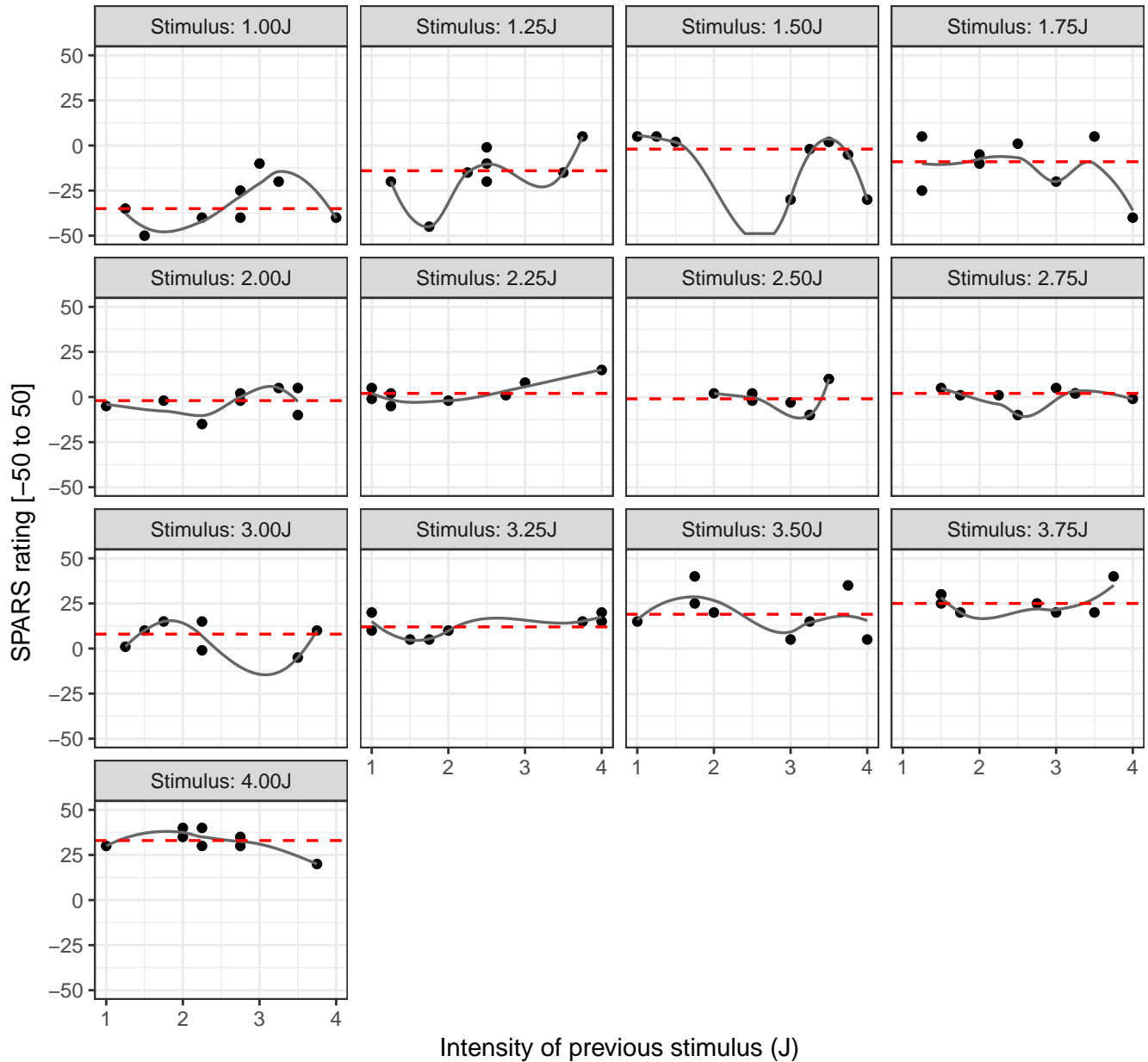
ID08: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



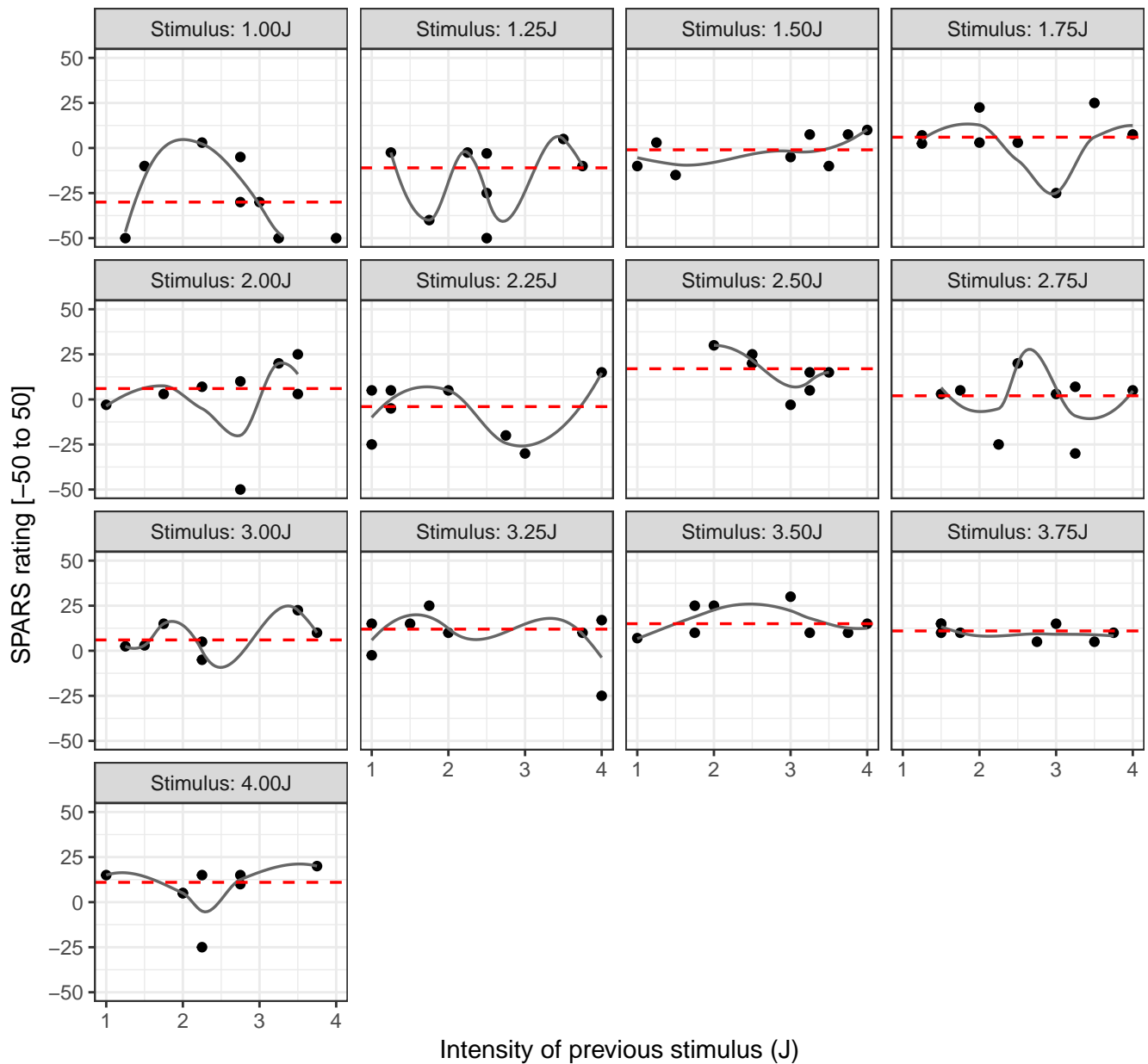
ID09: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



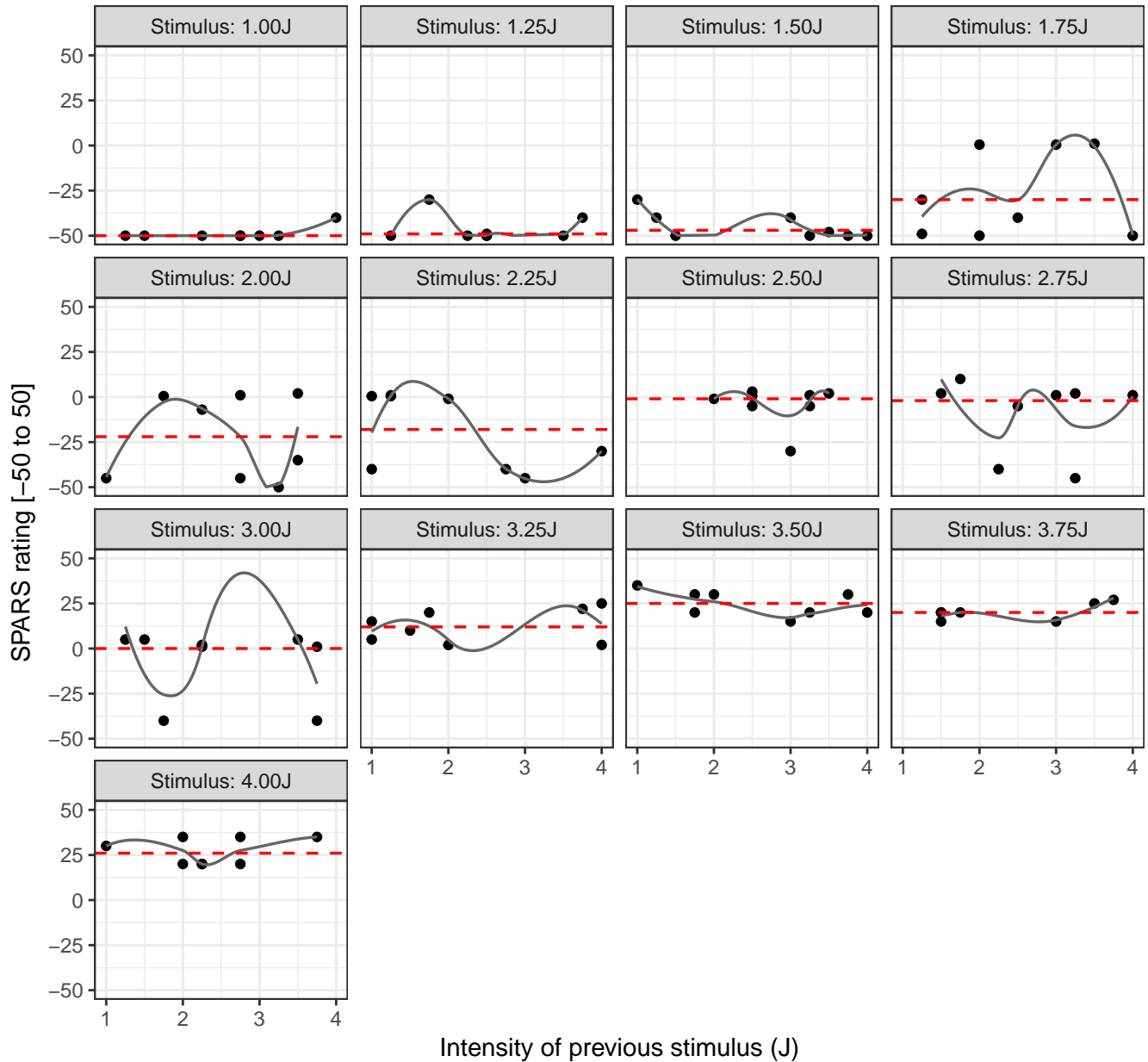
ID10: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



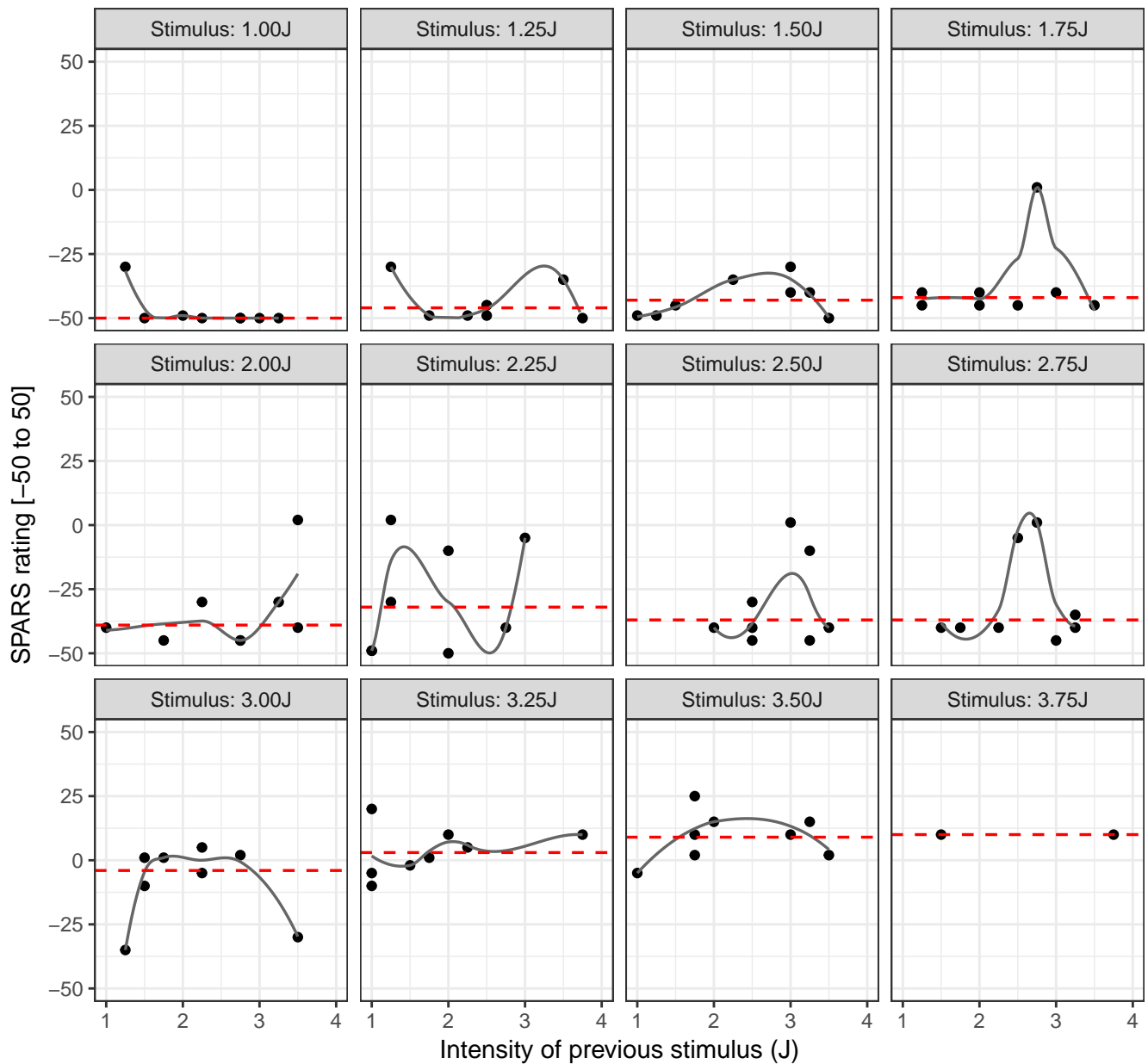
ID11: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



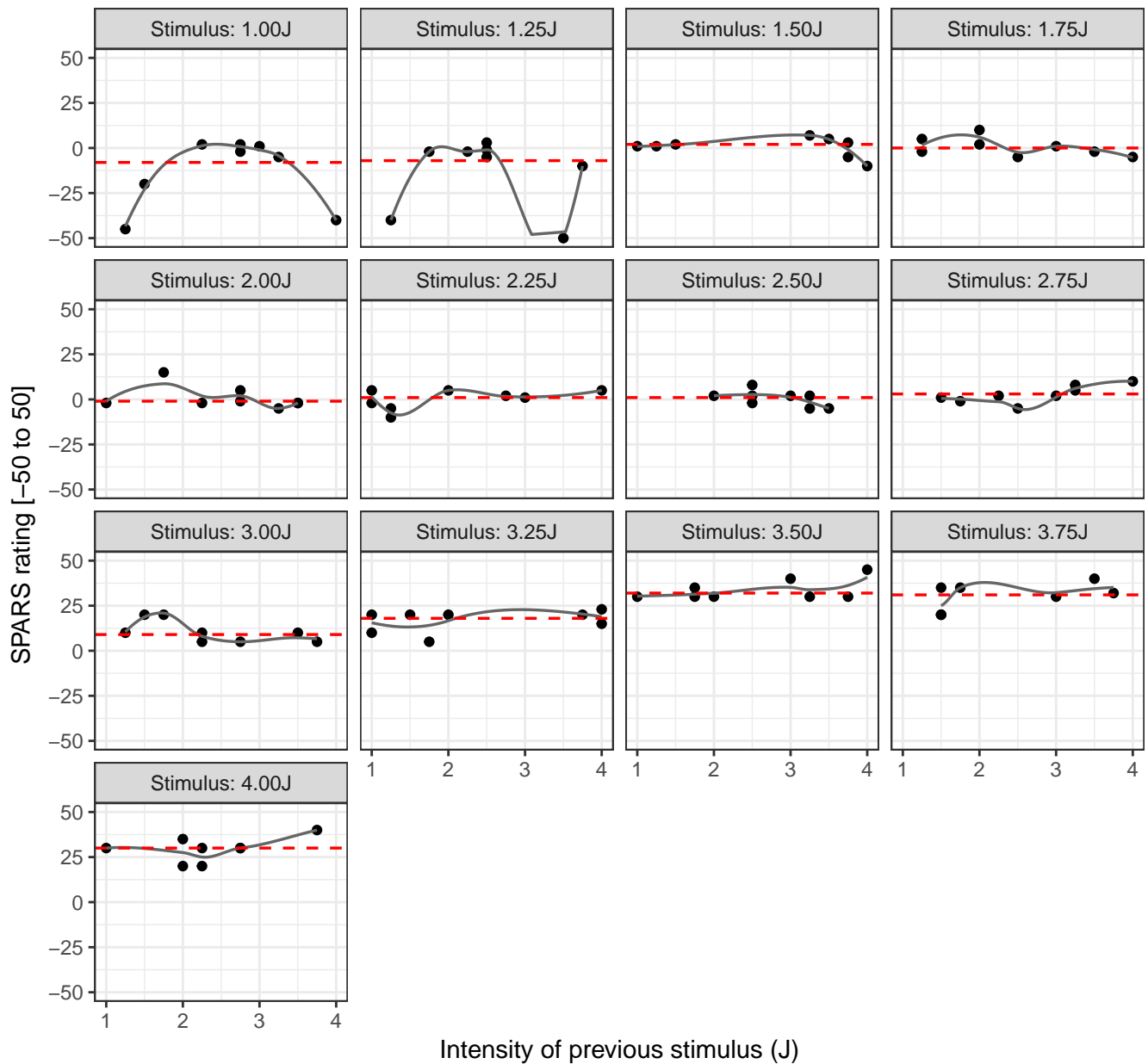
ID12: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



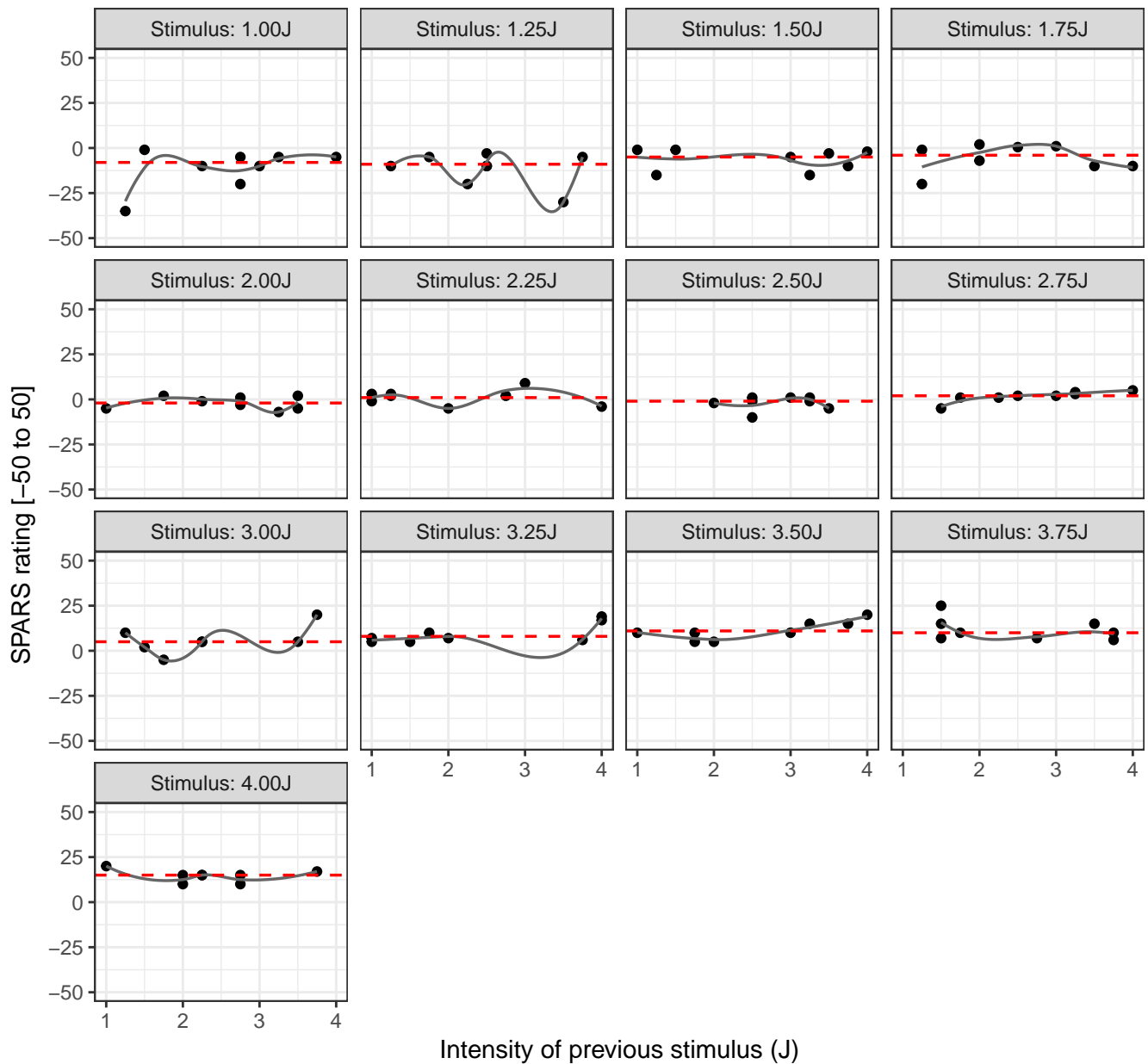
ID13: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



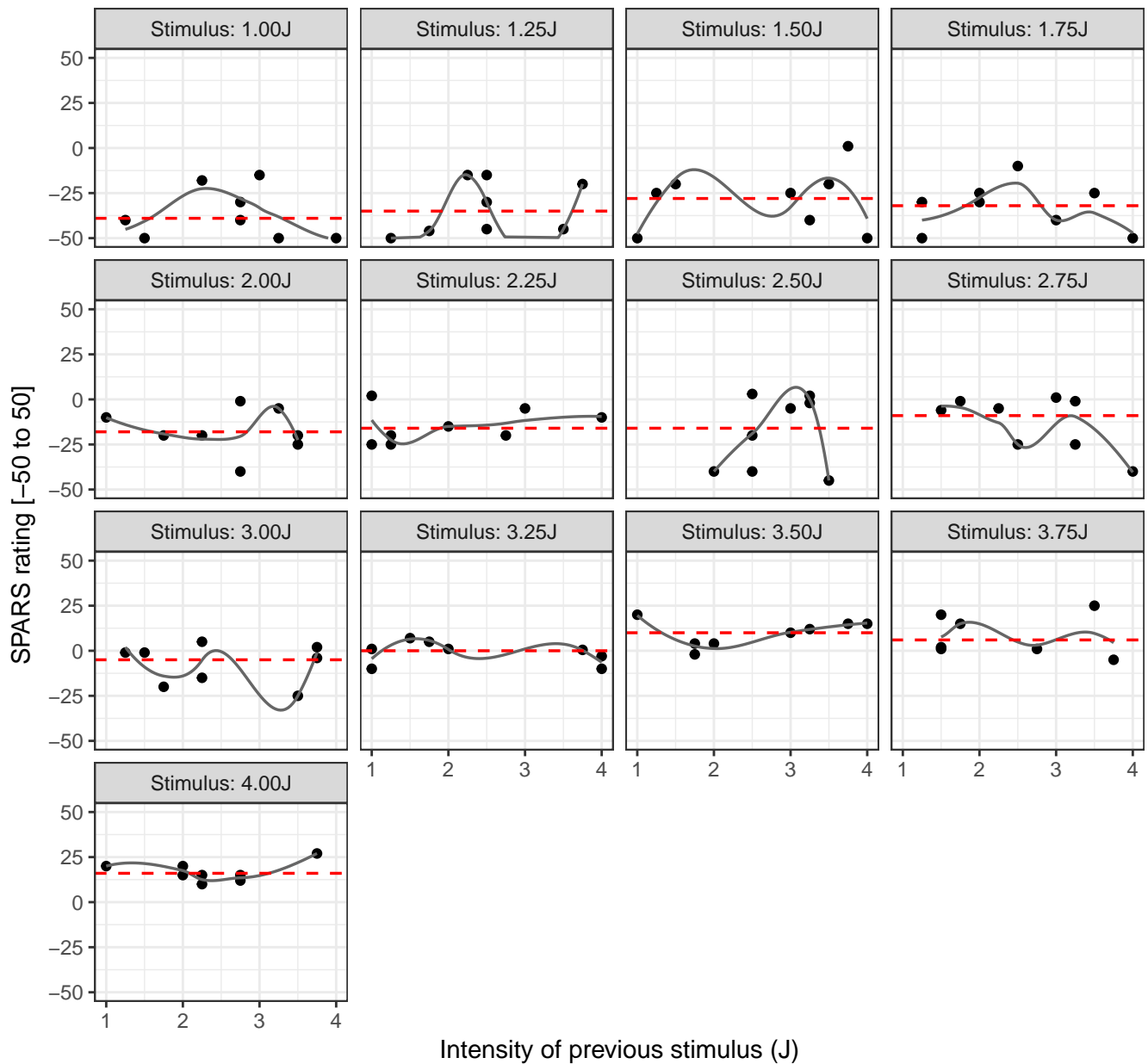
ID14: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



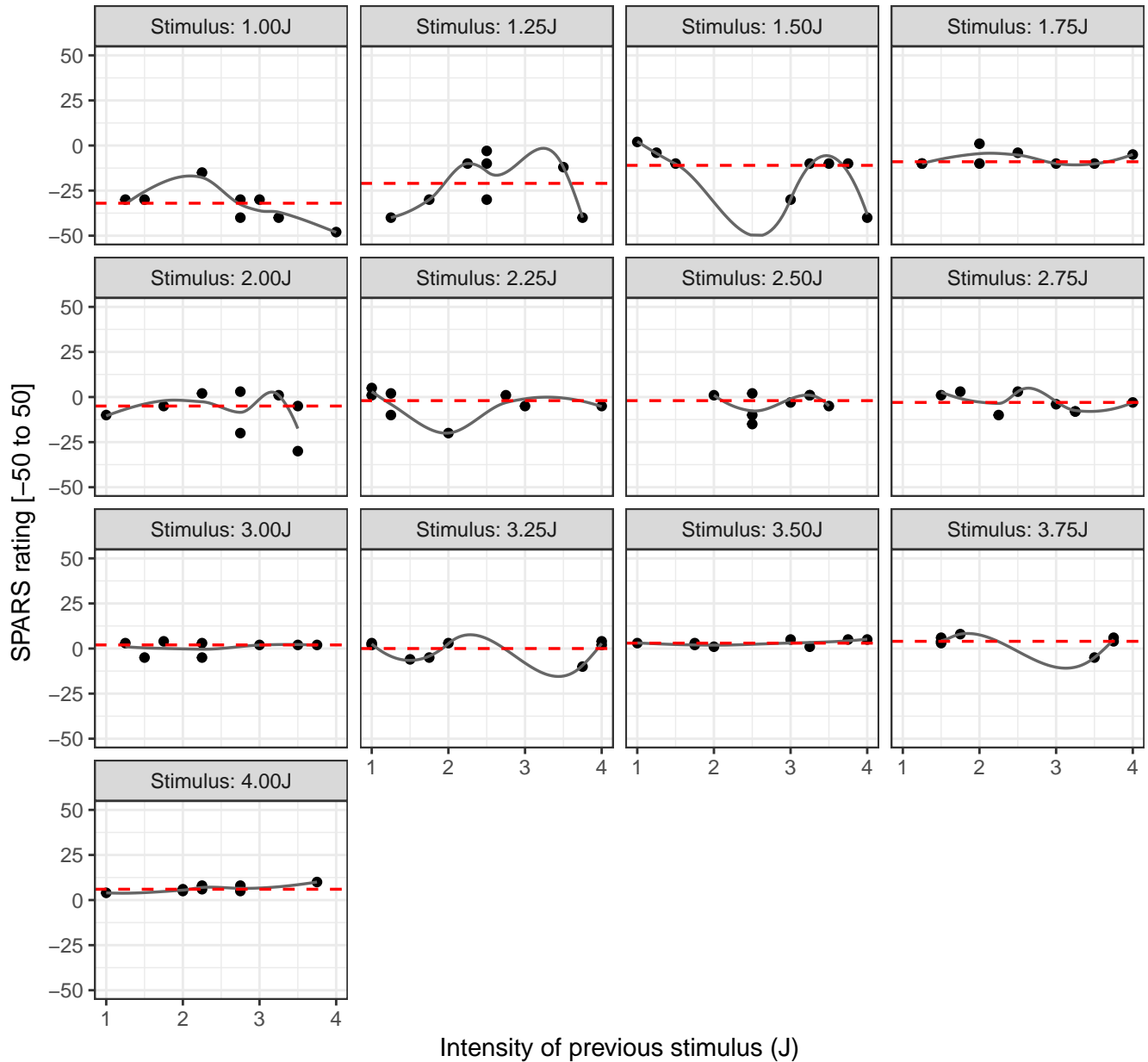
ID15: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



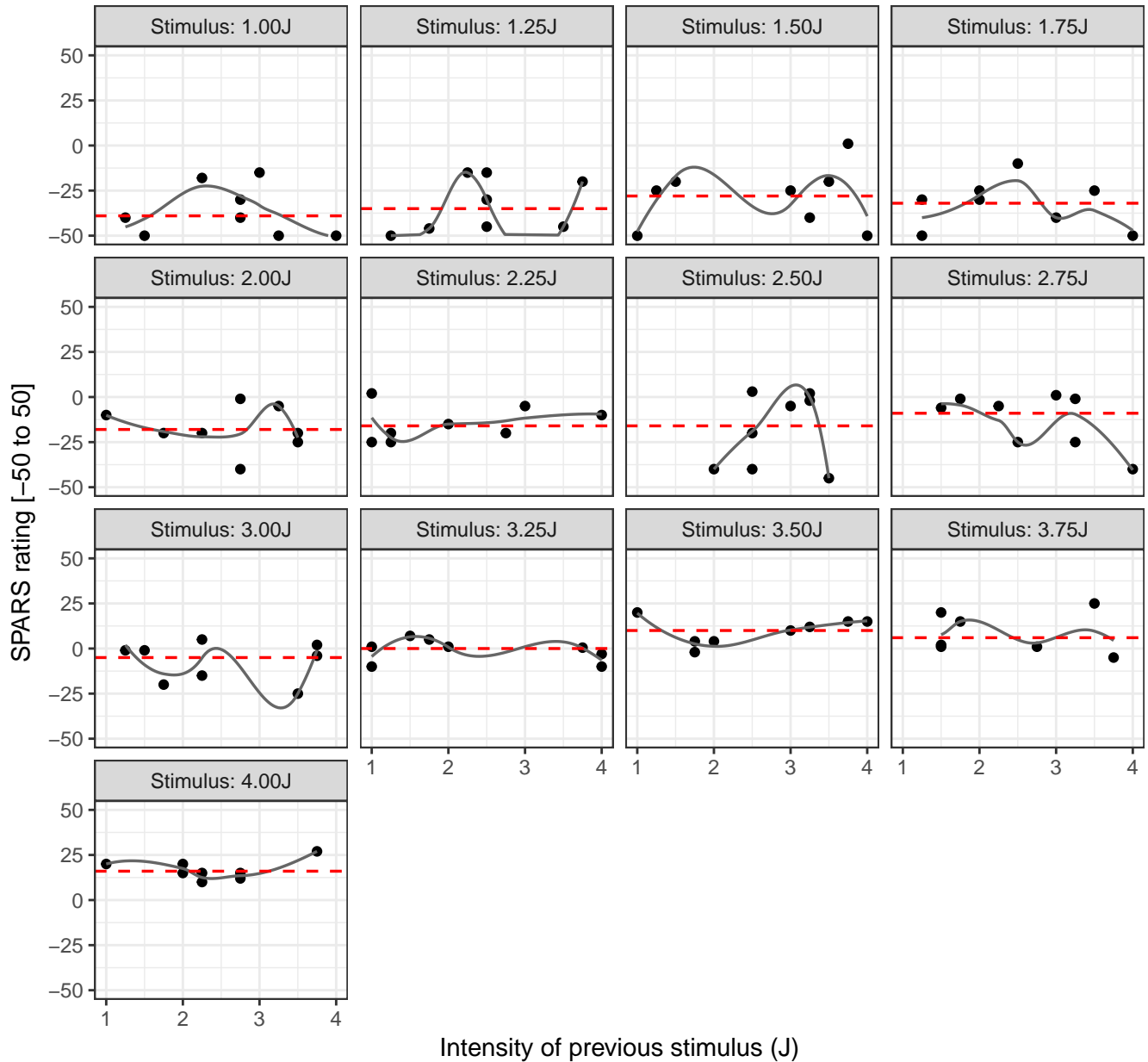
ID16: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



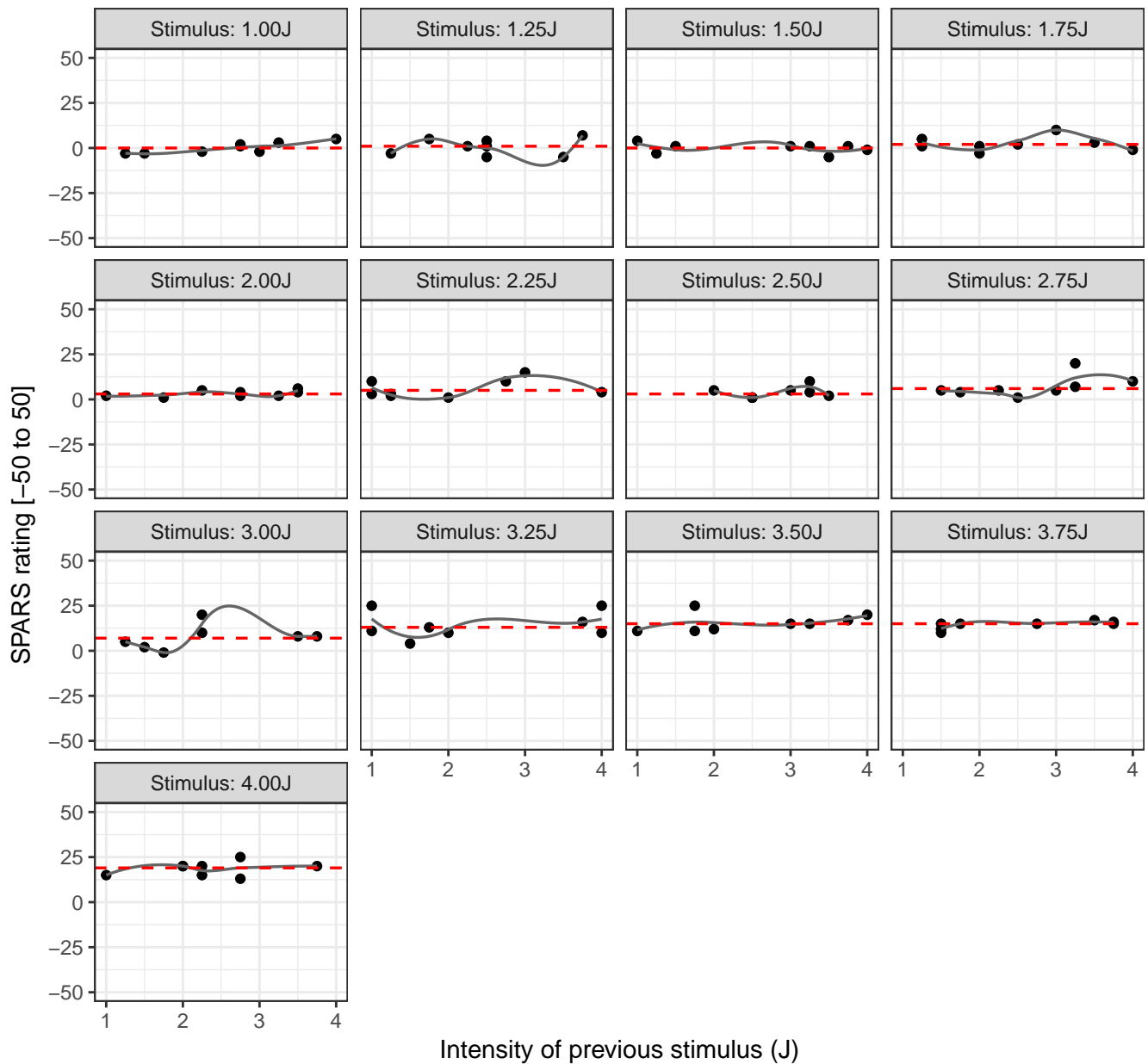
ID17: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



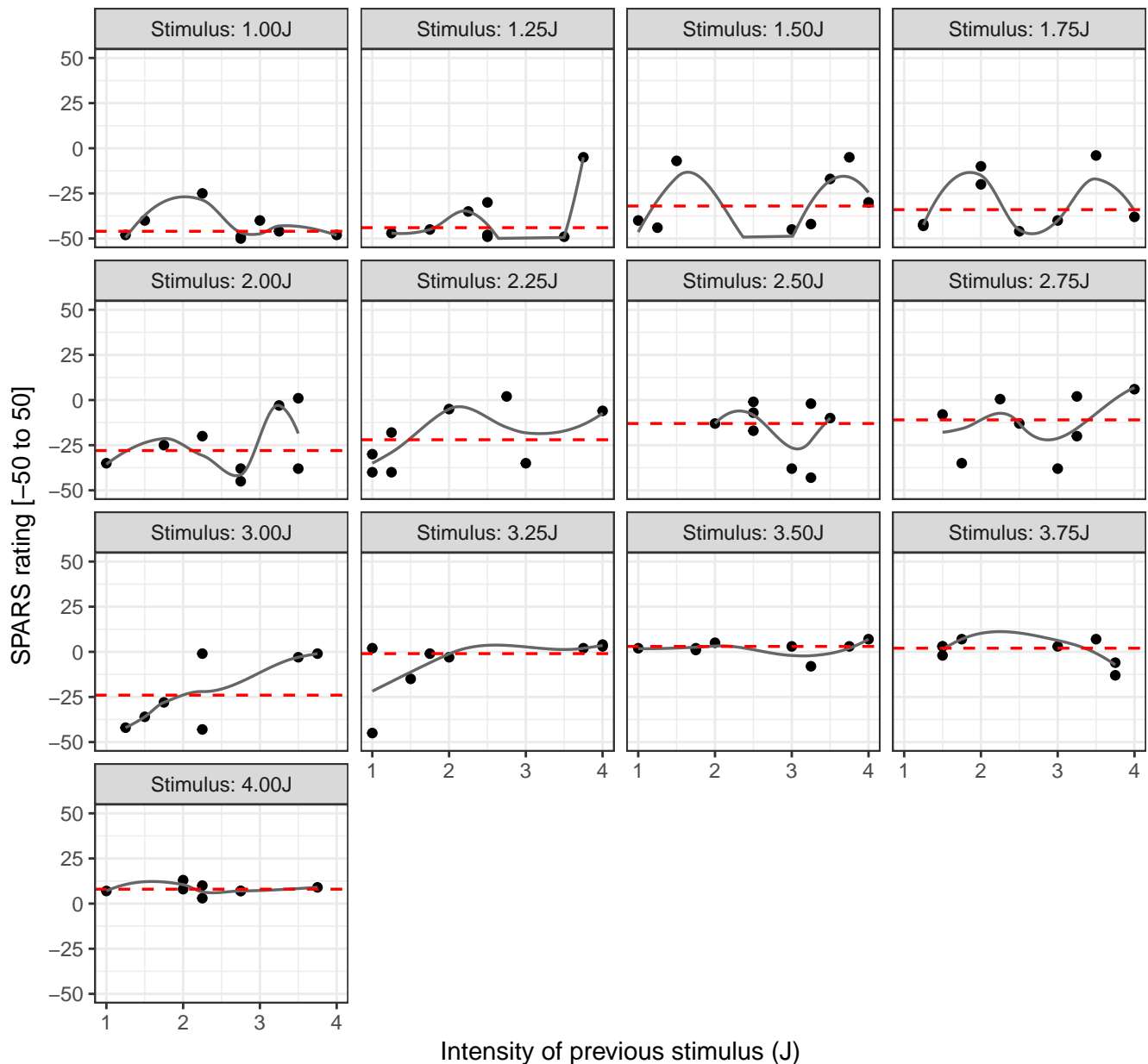
ID18: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



ID19: SPARS rating versus intensity of the previous stimulus, at different stimulus

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



Conclusion

Visual inspection of the figures shows the loess curve (grey curve) oscillating around the Tukey trimean (dashed red line), for all participants and across almost all stimulus intensities, indicating no systematic relationship between the rating of a stimulus and the intensity of the previous stimulus.

No further analysis undertaken.

Block order effects

Question: Is the SPARS rating for a given stimulus intensity associated with experimental block?

The analysis includes data from experimental blocks, ignoring trials. Because each stimulus intensity is repeated multiple times in each participant, a hierarchical analysis by study participant and stimulus intensity was conducted.

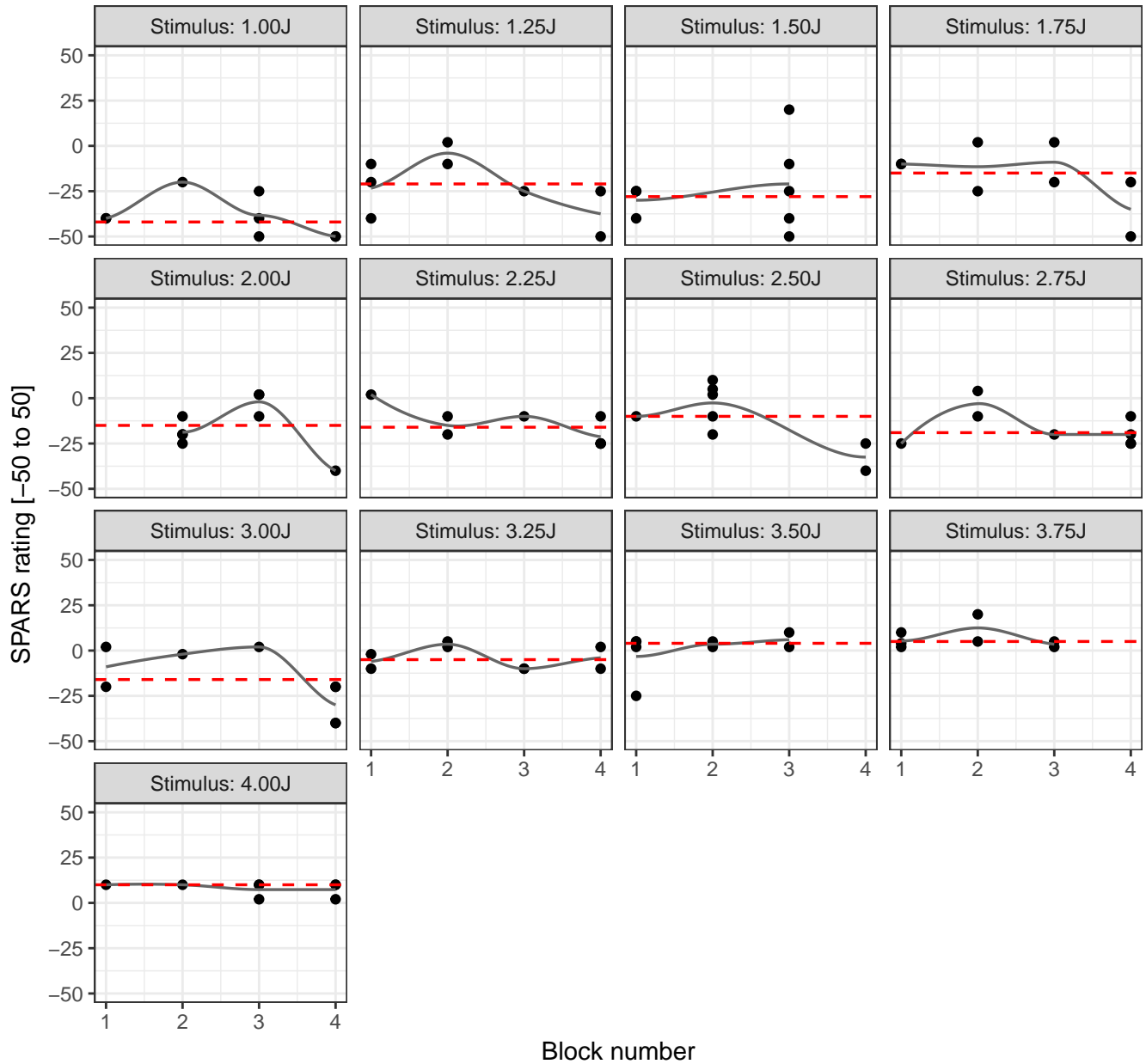
Participant-level exploratory plots

```
# Plot
plot_block <- plot_data %>%
  mutate(plot = pmap(.l = list(data, trimean, unique(PID)),
    ~ ggplot(data = ..1) +
      aes(x = block_order,
        y = rating) +
      geom_point() +
      geom_smooth(se = FALSE,
        colour = '#666666',
        size = 0.6,
        na.rm = TRUE) +
      geom_hline(data = ..2,
        aes(yintercept = tm),
        colour = 'red',
        size = 0.6,
        linetype = 2) +
      labs(title = paste0(..3, ': SPARS rating versus experimental block n',
        subtitle = 'Red line (dashed): Tukey trimean of SPARS rating |
        y = 'SPARS rating [-50 to 50]',
        x = 'Block number') +
      scale_y_continuous(limits = c(-50, 50)) +
      scale_x_continuous(limits = c(1, 4)) +
      facet_wrap(~intensity,
        ncol = 4) +
      theme_bw()))

# Print plots
walk(.x = plot_block$plot, ~ print(.x))
```

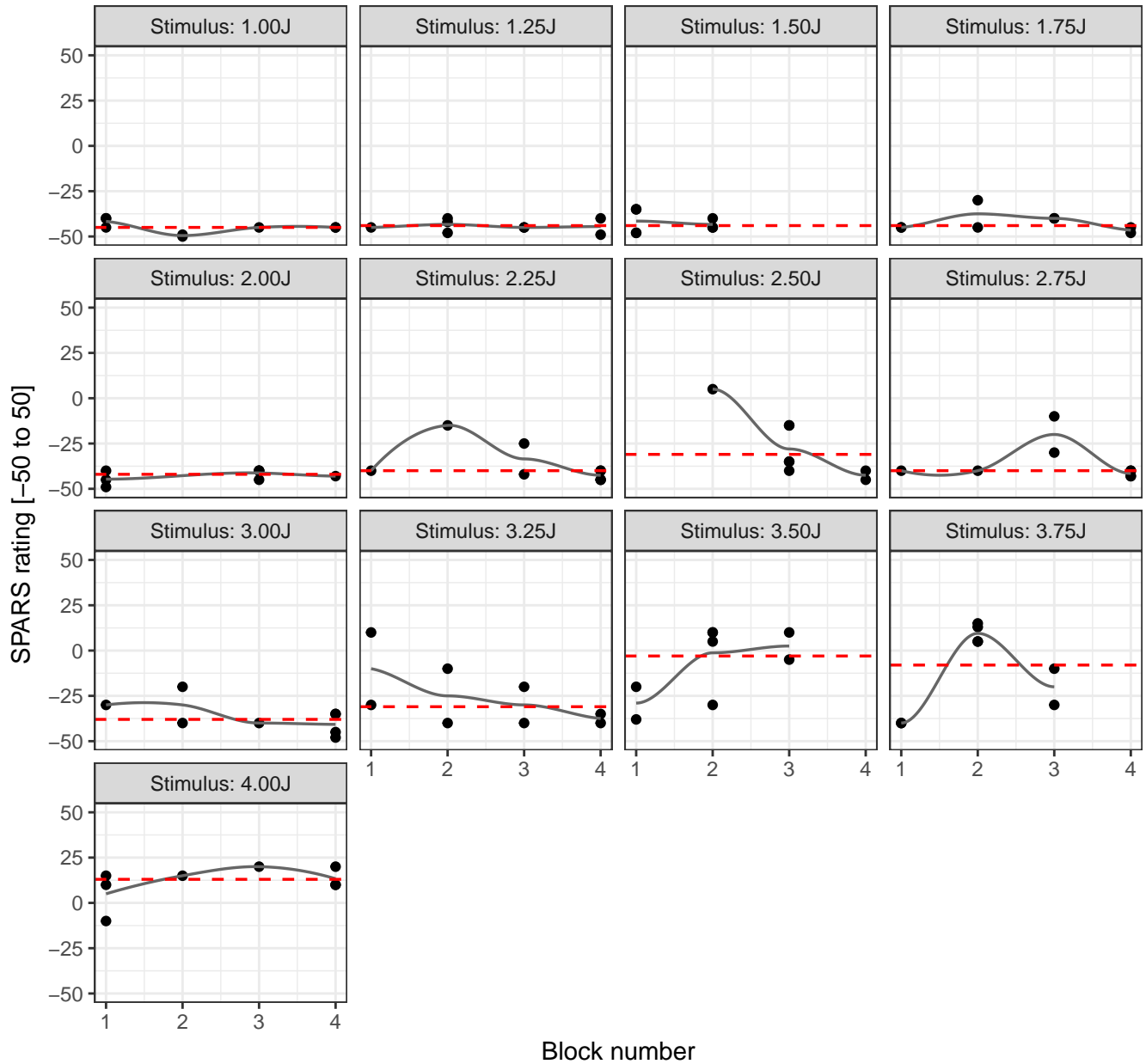
ID01: SPARS rating versus experimental block number, at different stimulus intensities

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



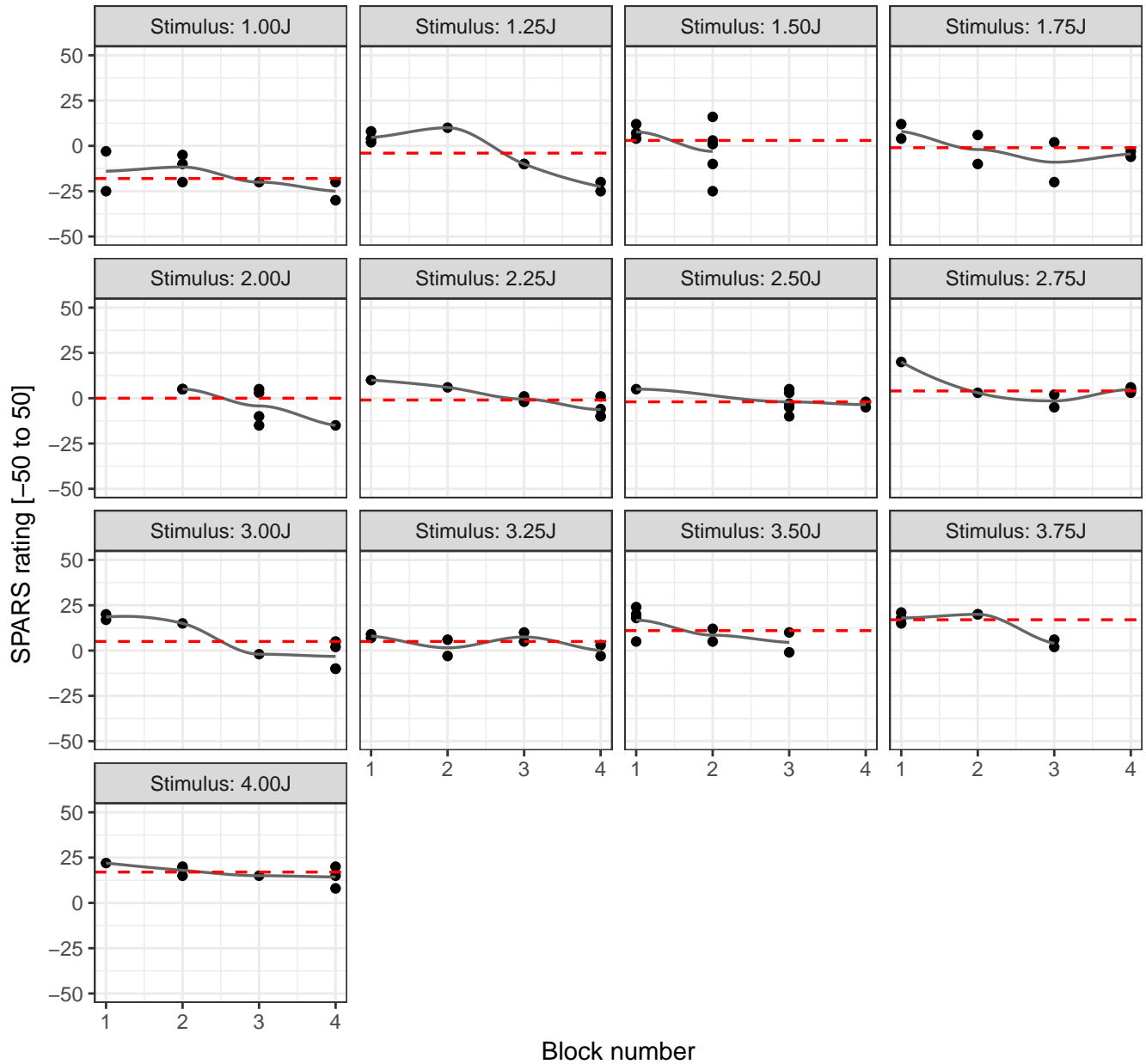
ID02: SPARS rating versus experimental block number, at different stimulus intens

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



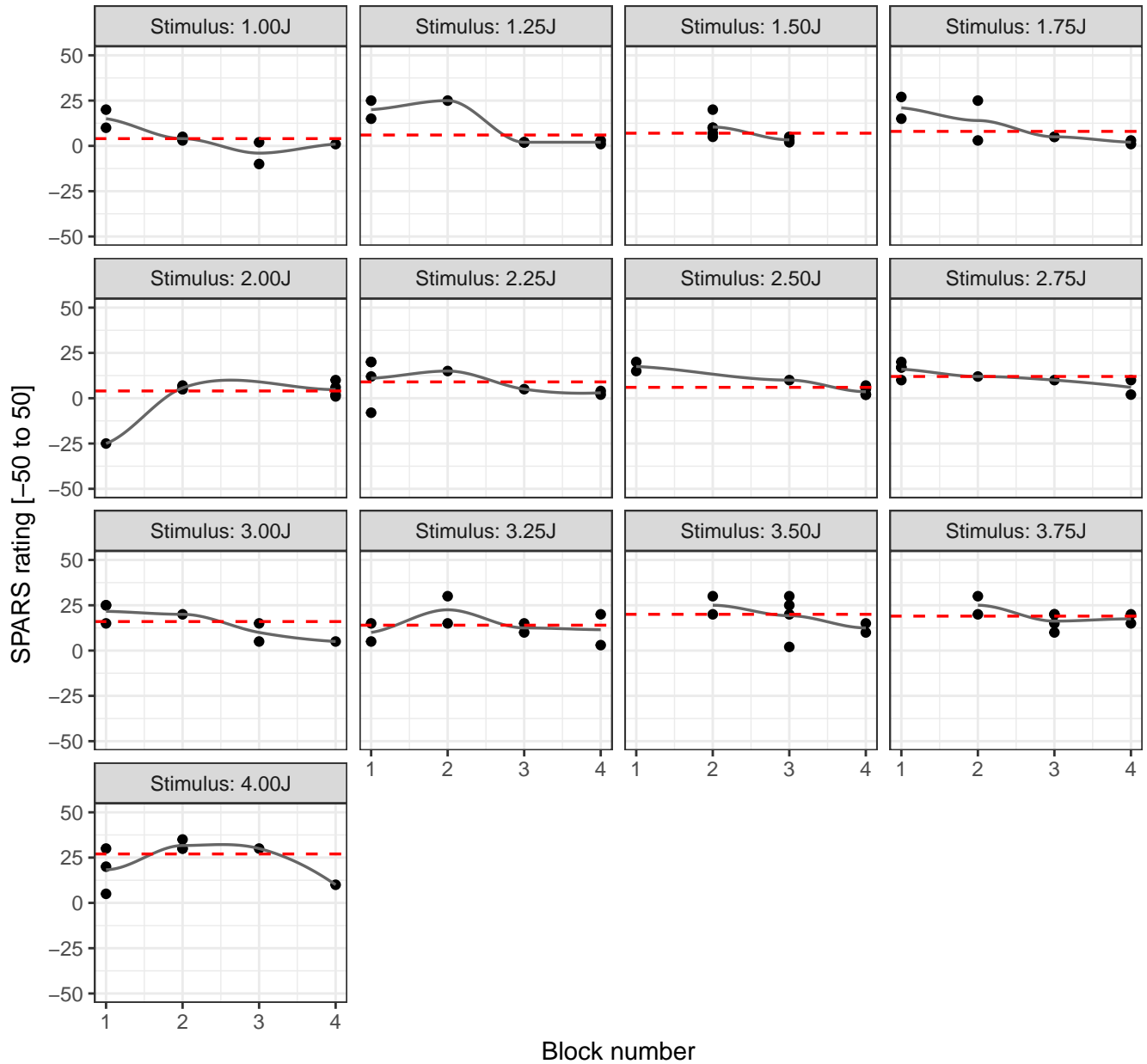
ID03: SPARS rating versus experimental block number, at different stimulus intens

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



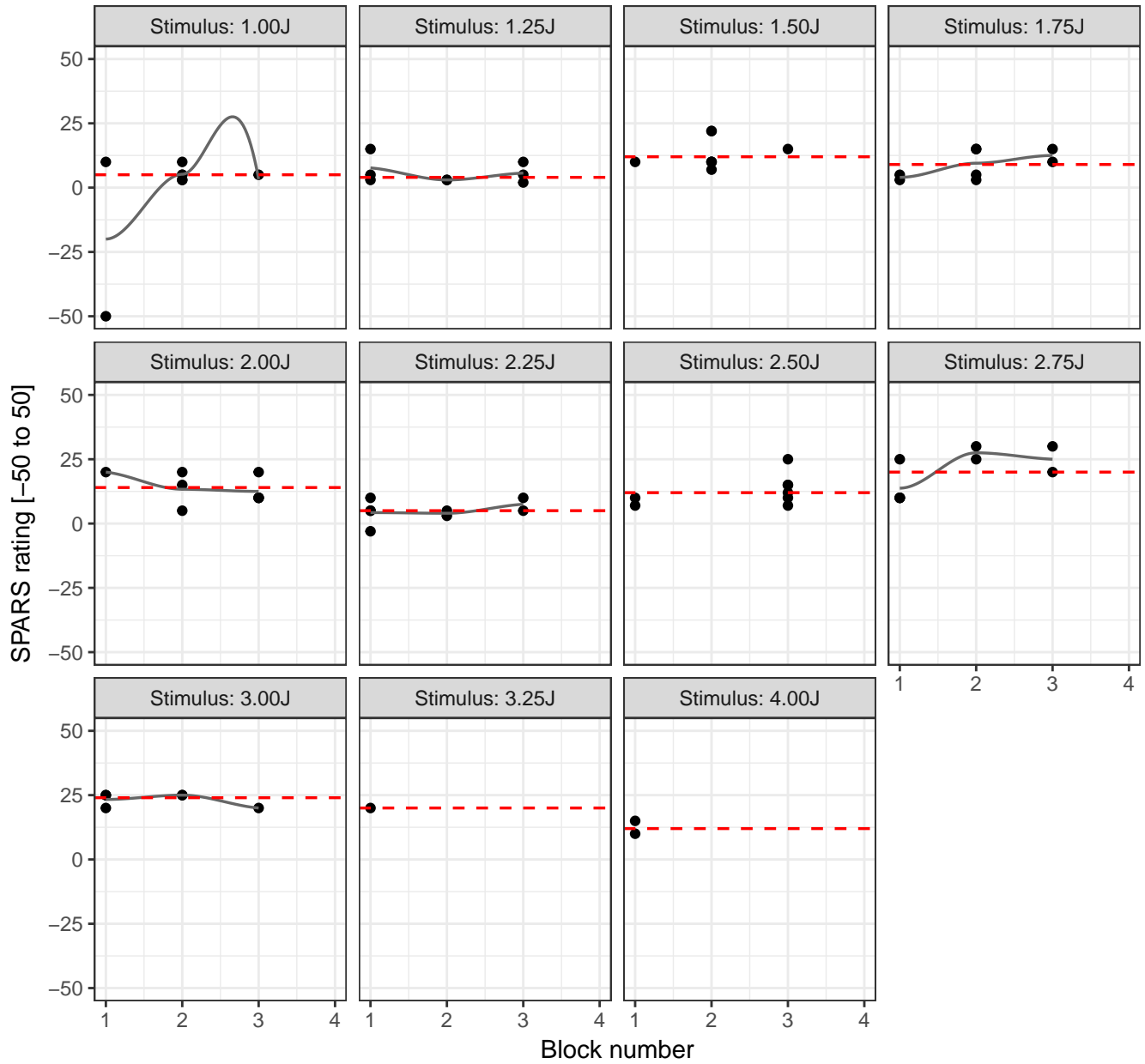
ID04: SPARS rating versus experimental block number, at different stimulus intens

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



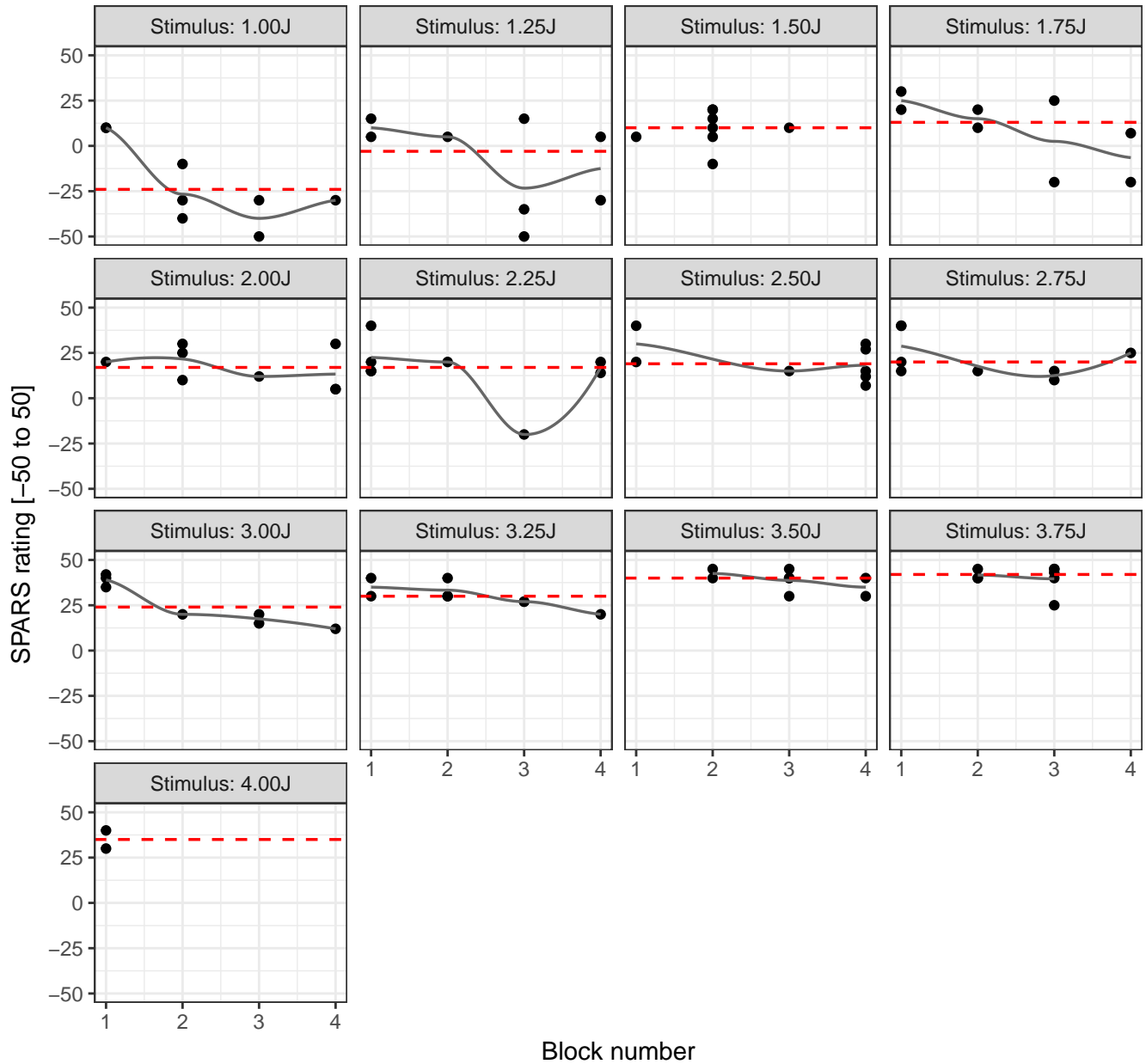
ID05: SPARS rating versus experimental block number, at different stimulus intensities

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



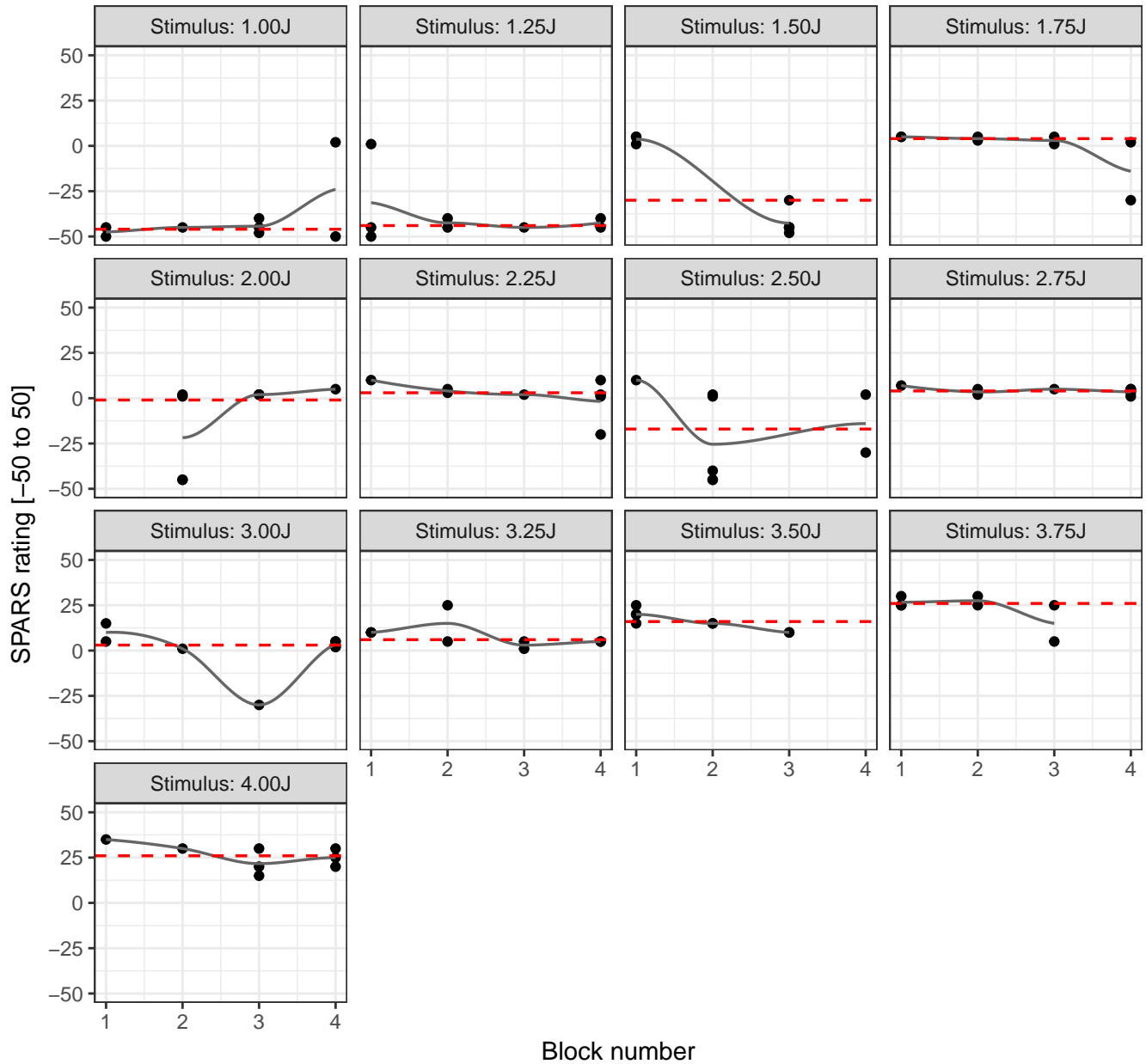
ID06: SPARS rating versus experimental block number, at different stimulus intens

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



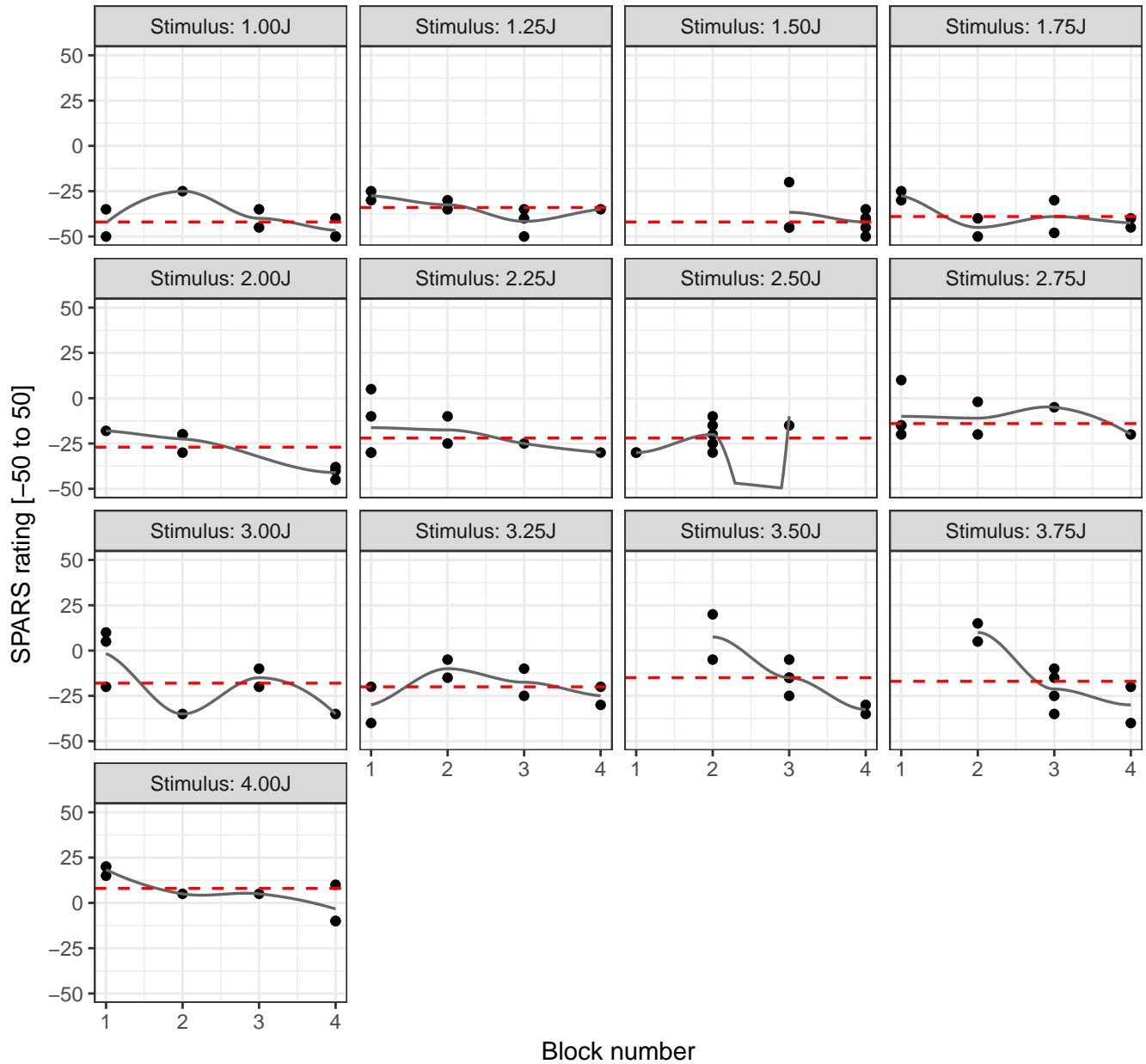
ID07: SPARS rating versus experimental block number, at different stimulus intens

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



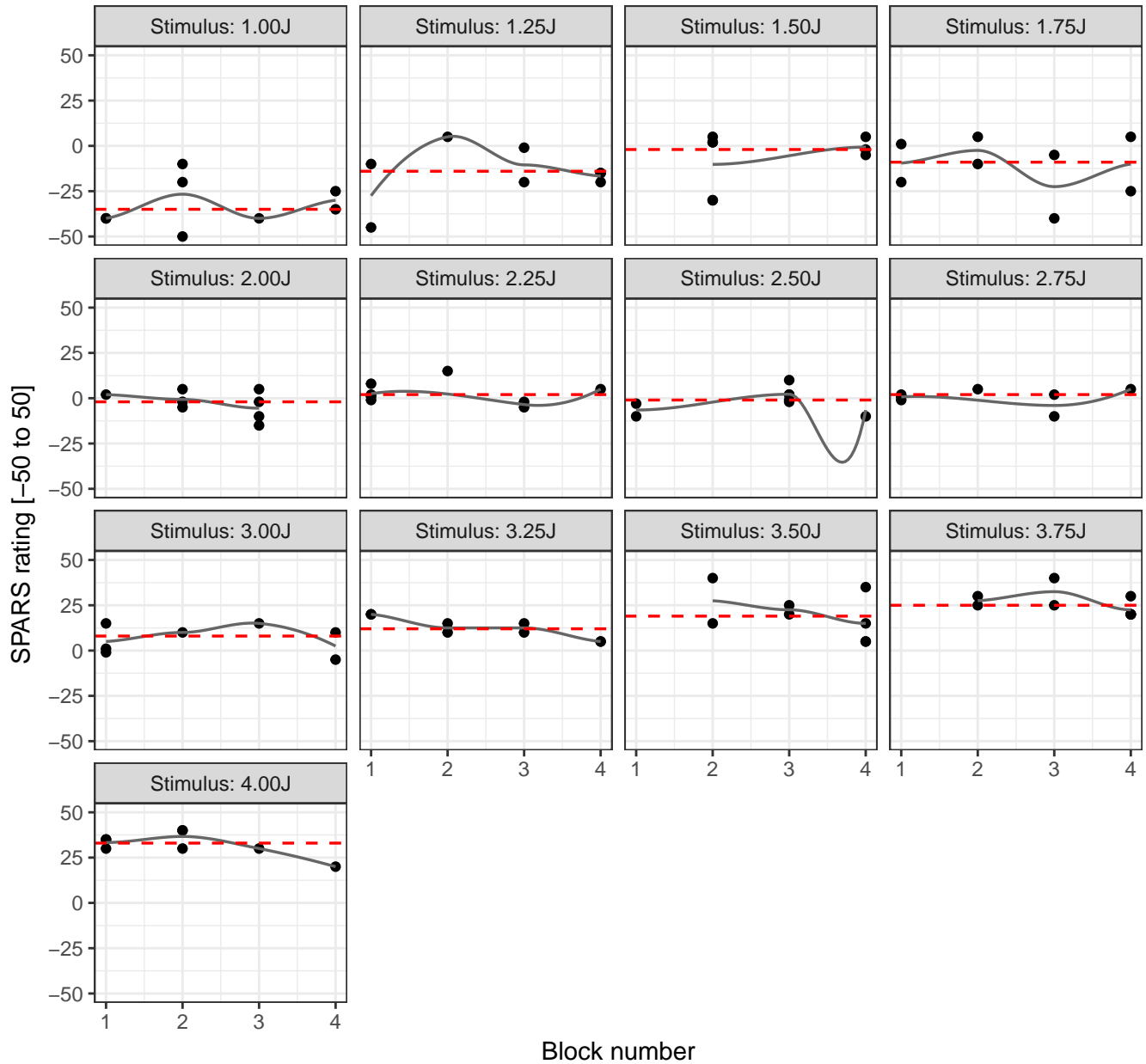
ID08: SPARS rating versus experimental block number, at different stimulus intens

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



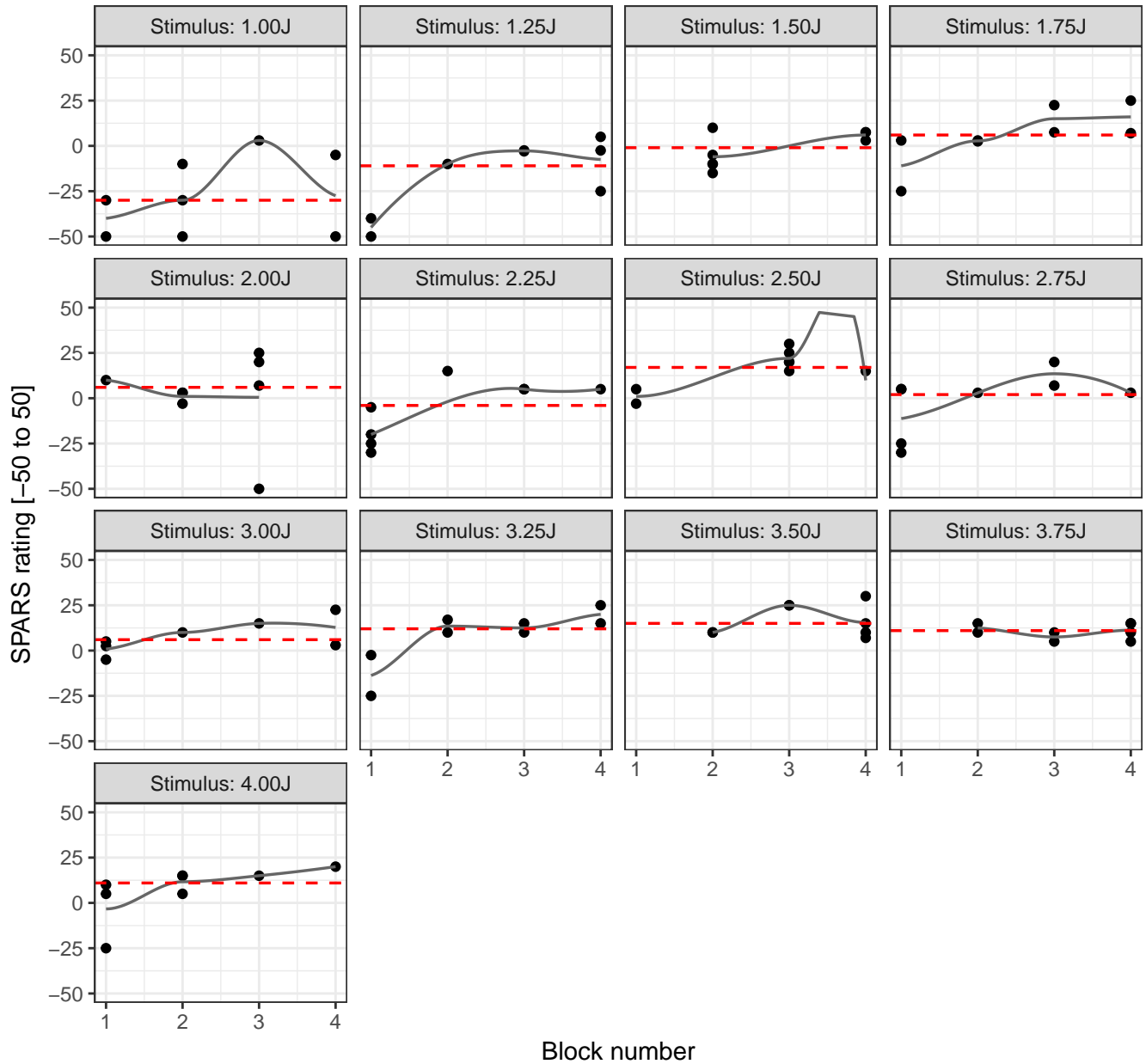
ID09: SPARS rating versus experimental block number, at different stimulus intens

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



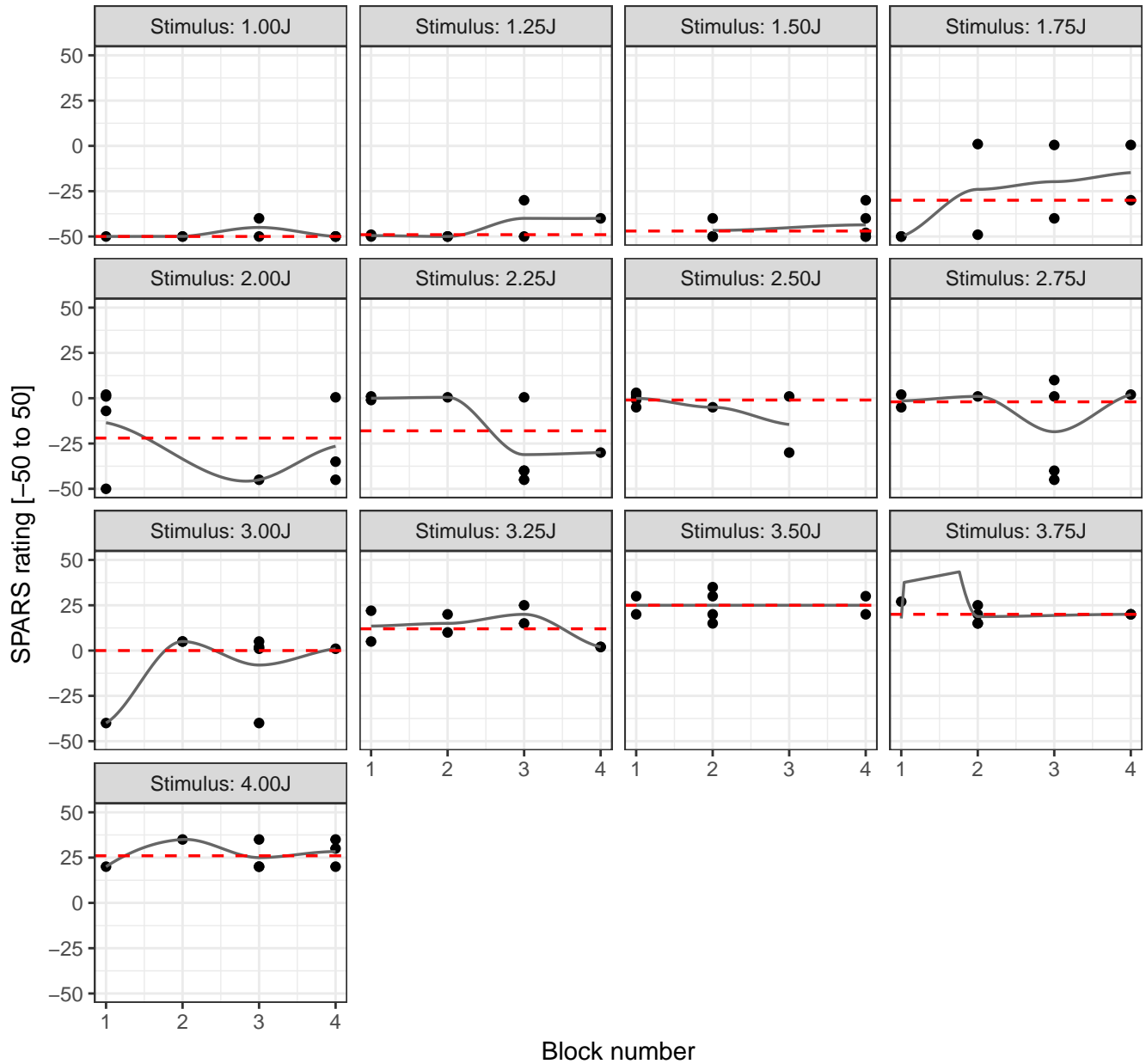
ID10: SPARS rating versus experimental block number, at different stimulus intensities

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



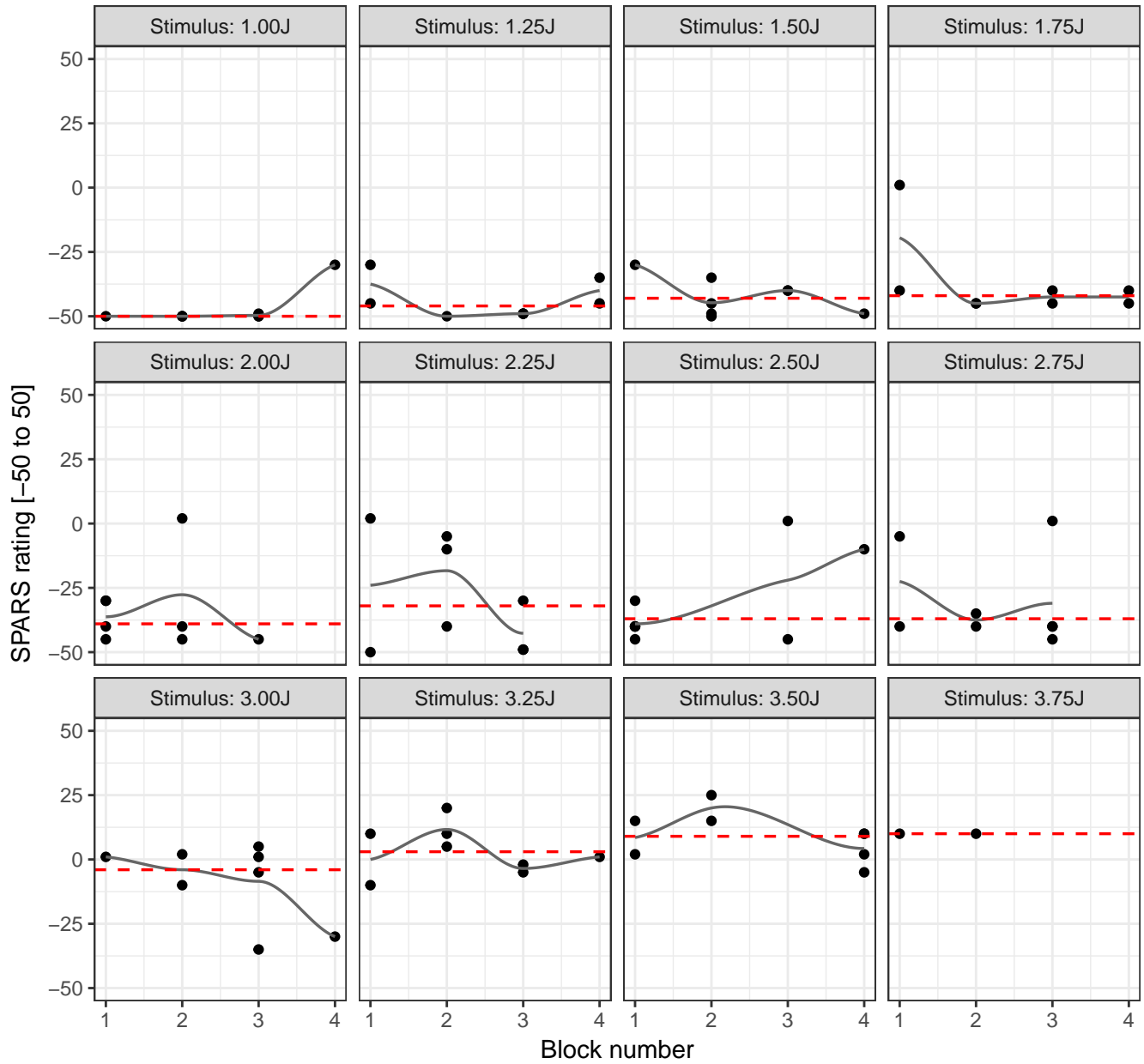
ID11: SPARS rating versus experimental block number, at different stimulus intens

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



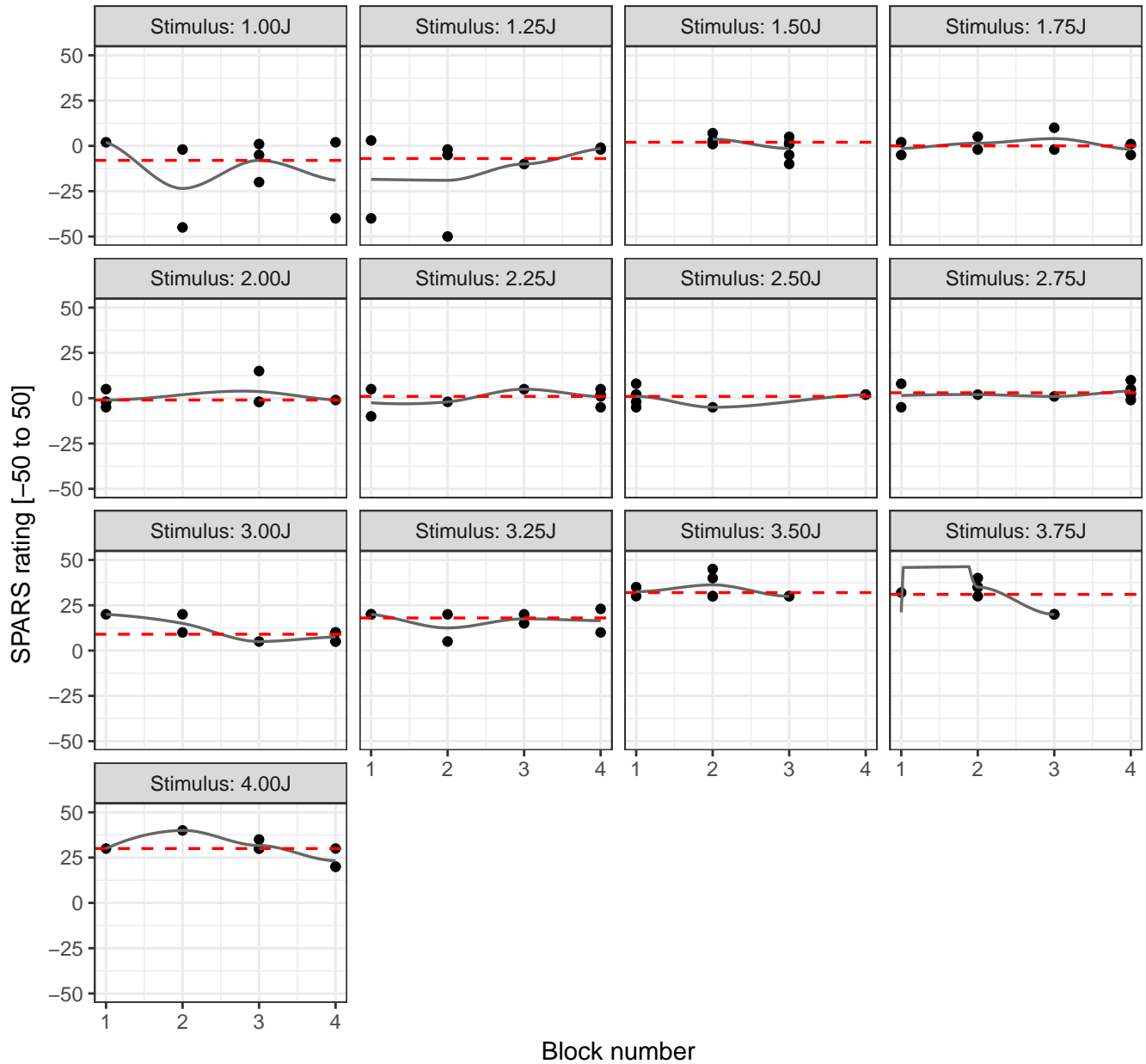
ID12: SPARS rating versus experimental block number, at different stimulus intensities

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



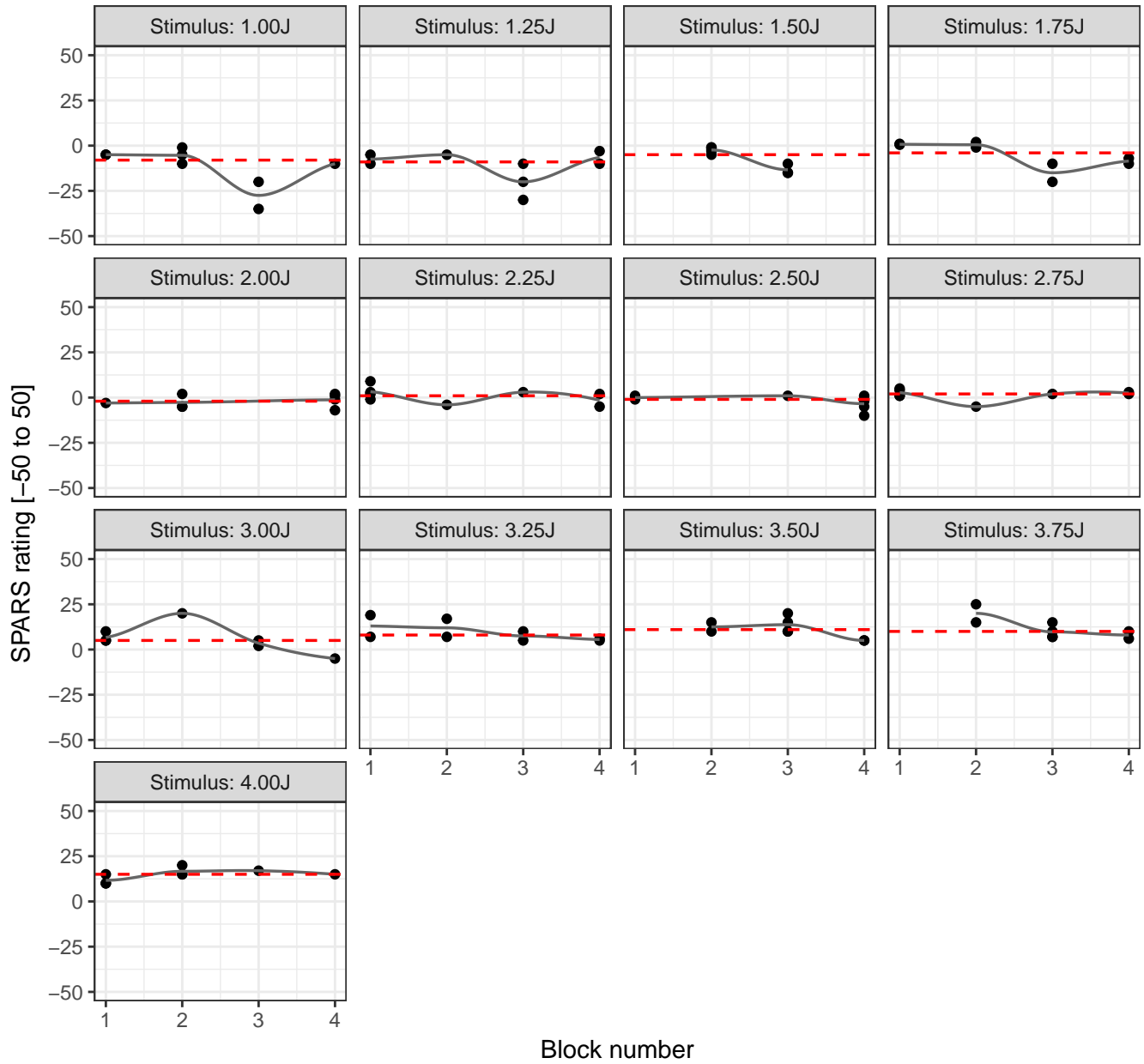
ID13: SPARS rating versus experimental block number, at different stimulus intensities

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



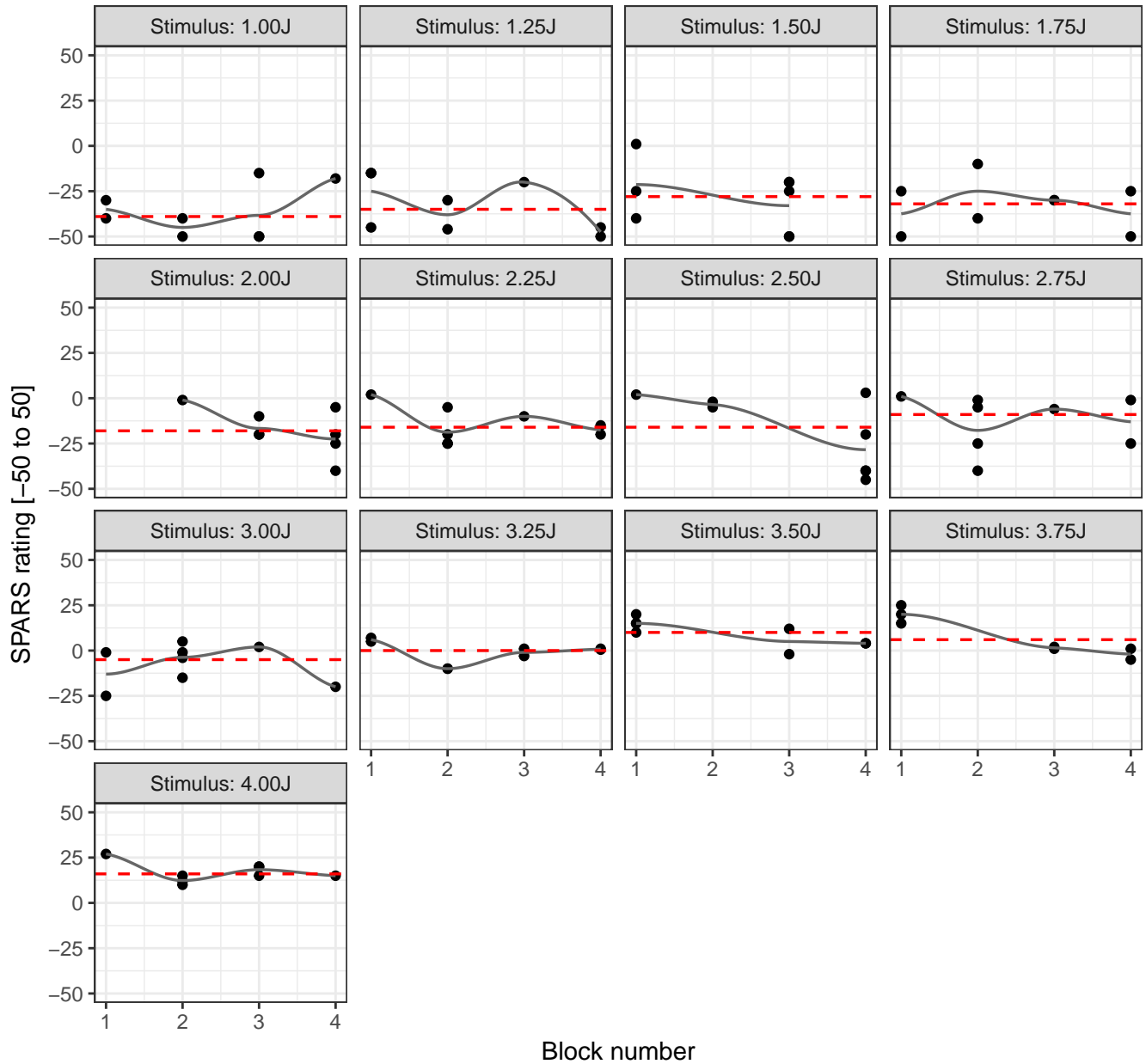
ID14: SPARS rating versus experimental block number, at different stimulus intensities

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



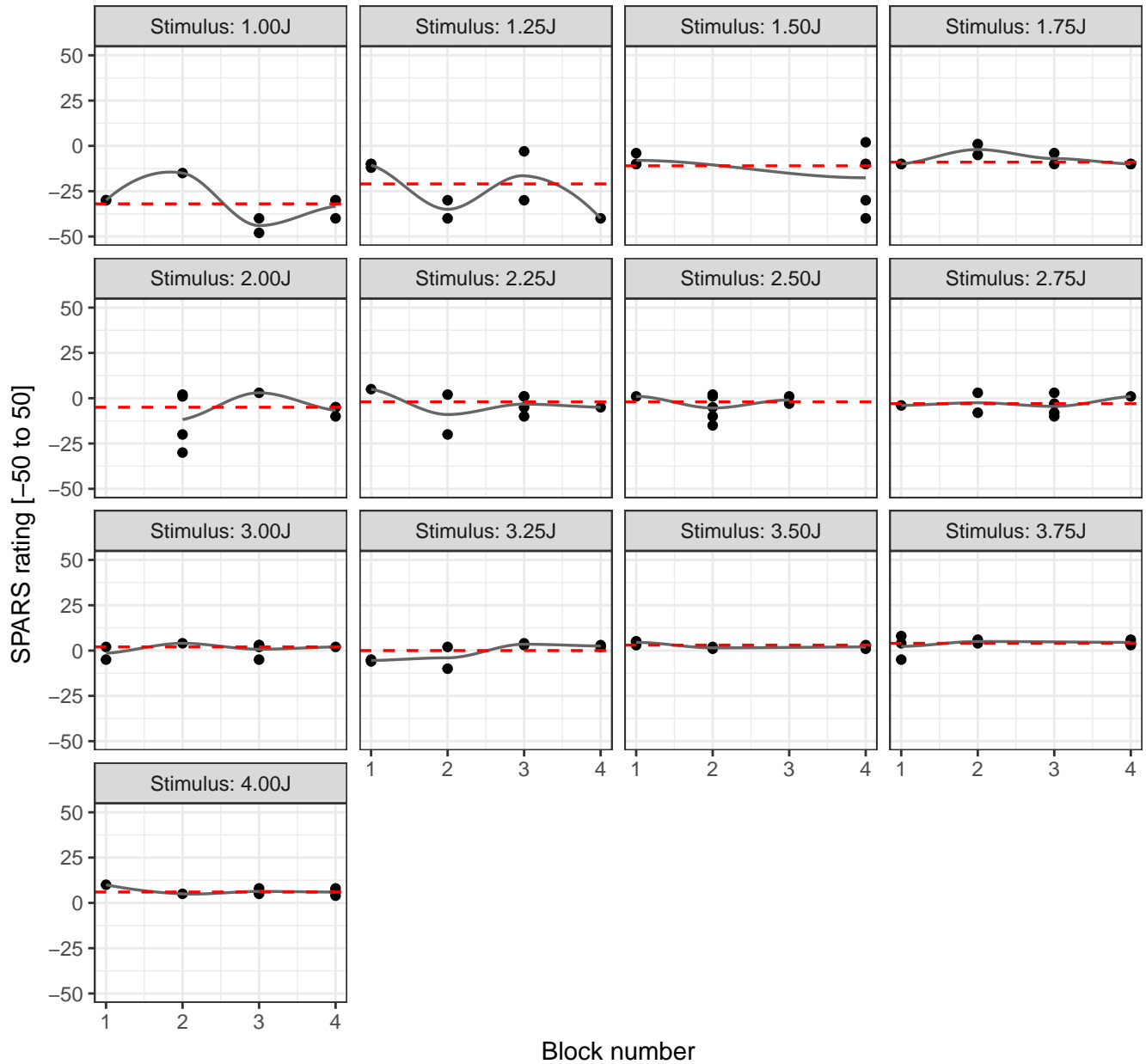
ID15: SPARS rating versus experimental block number, at different stimulus intensities

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



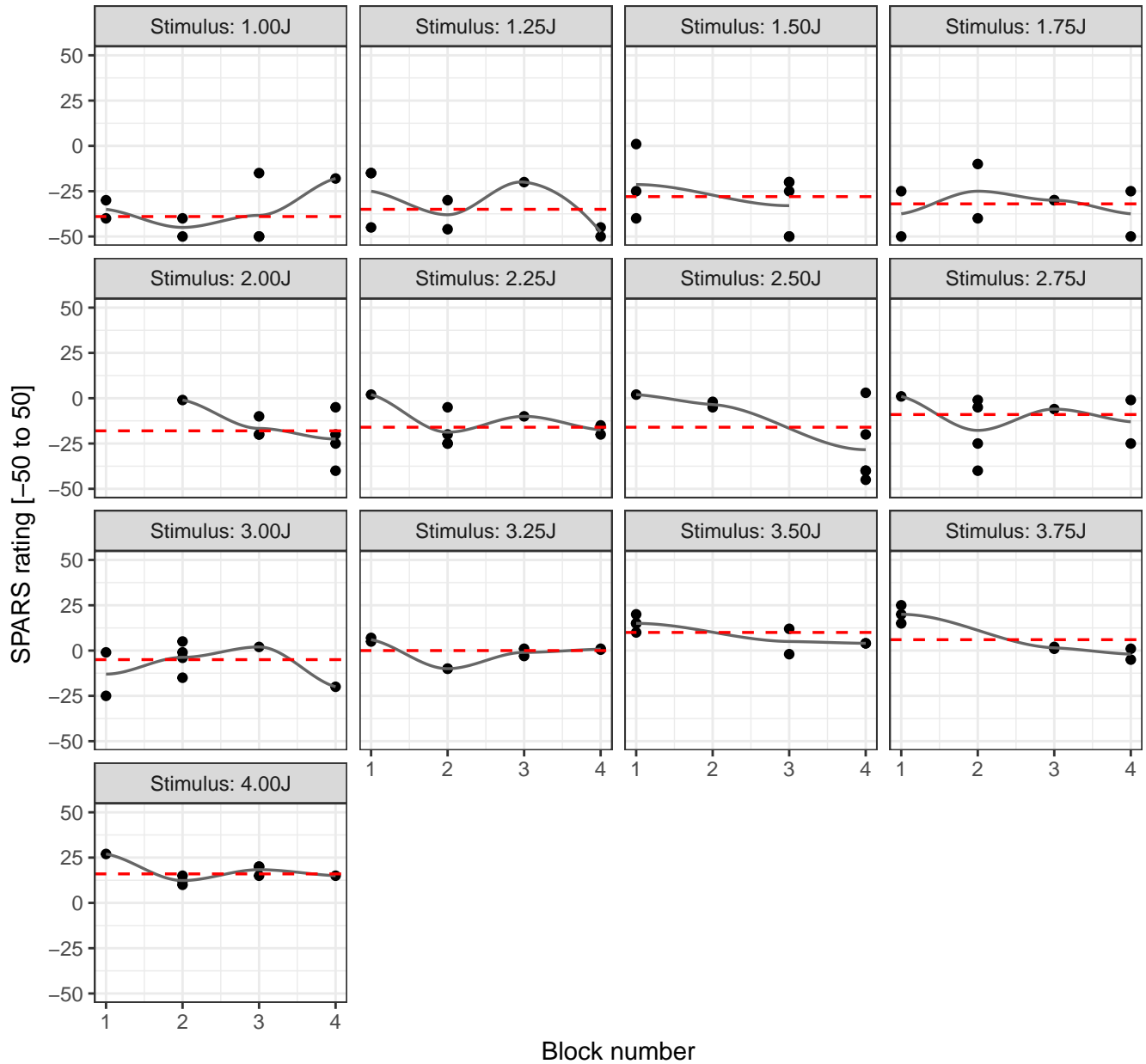
ID16: SPARS rating versus experimental block number, at different stimulus intensities

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



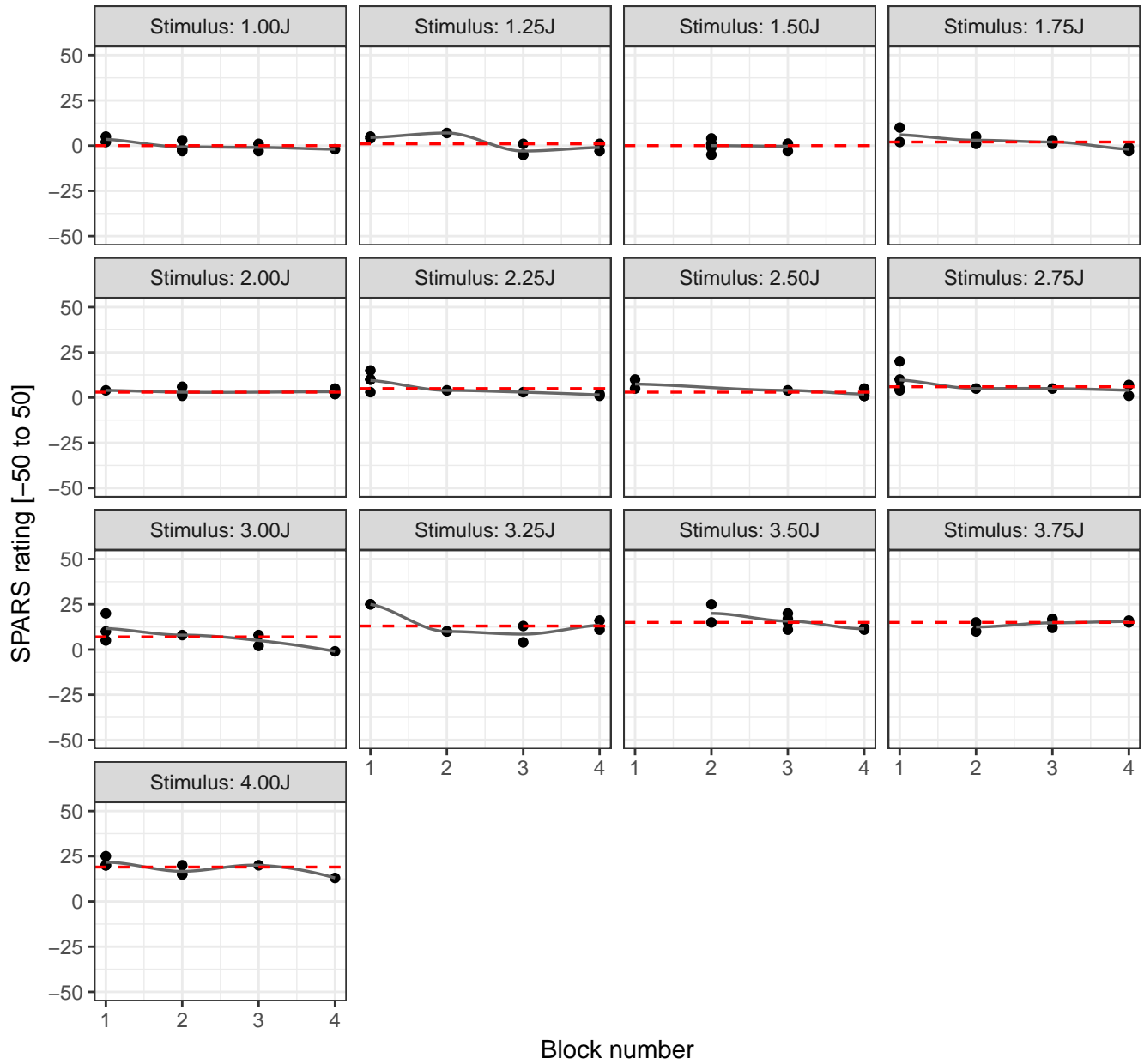
ID17: SPARS rating versus experimental block number, at different stimulus intensities

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



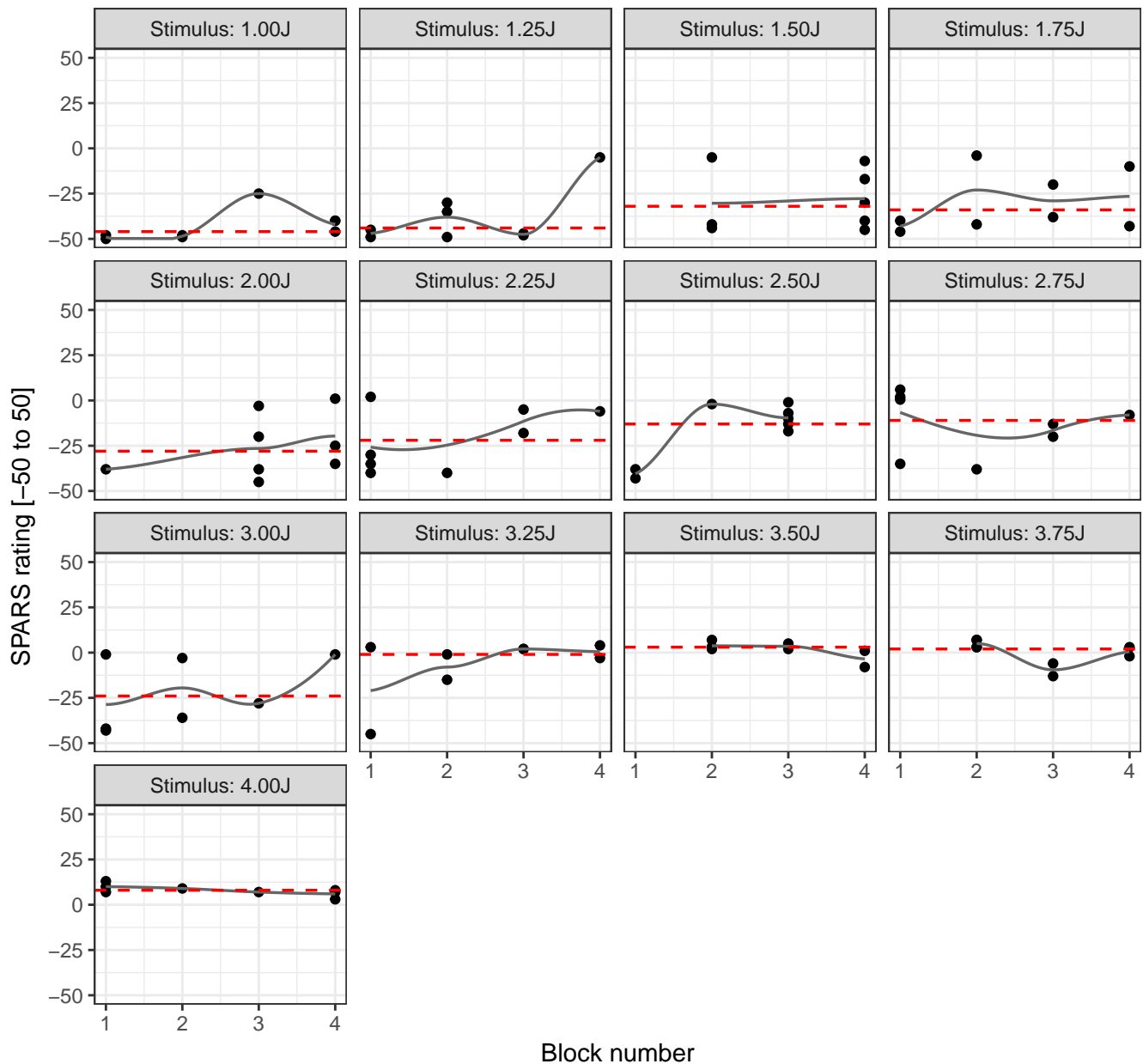
ID18: SPARS rating versus experimental block number, at different stimulus intensities

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



ID19: SPARS rating versus experimental block number, at different stimulus intensities

Red line (dashed): Tukey trimean of SPARS rating | Grey line (solid): Loess curve



Conclusion

Visual inspection of the figures shows the loess curve (grey curve) oscillating around the Tukey trimean (dashed red line), for all participants and across almost all stimulus intensities, indicating no systematic relationship between the rating of a stimulus and the experimental block.

No further analysis undertaken.

Session information

```
## R version 3.5.0 (2018-04-23)
## Platform: x86_64-apple-darwin15.6.0 (64-bit)
## Running under: macOS High Sierra 10.13.5
```

```
##
## 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
##
## attached base packages:
## [1] stats      graphics  grDevices  utils      datasets  methods   base
##
## other attached packages:
## [1] bindrcpp_0.2.2      robustlmm_2.2-1      lme4_1.1-17
## [4] Matrix_1.2-14       forcats_0.3.0        stringr_1.3.1
## [7] dplyr_0.7.5         purrr_0.2.5          readr_1.1.1
## [10] tidyr_0.8.1         tibble_1.4.2         ggplot2_2.2.1.9000
## [13] tidyverse_1.2.1     magrittr_1.5
##
## loaded via a namespace (and not attached):
## [1] Rcpp_0.12.17         lubridate_1.7.4      lattice_0.20-35
## [4] assertthat_0.2.0     rprojroot_1.3-2      digest_0.6.15
## [7] psych_1.8.4          R6_2.2.2             cellranger_1.1.0
## [10] plyr_1.8.4           backports_1.1.2      evaluate_0.10.1
## [13] httr_1.3.1           pillar_1.2.3         rlang_0.2.1
## [16] lazyeval_0.2.1       readxl_1.1.0         rstudioapi_0.7
## [19] minqa_1.2.4          nloptr_1.0.4         rmarkdown_1.9
## [22] labeling_0.3         splines_3.5.0        foreign_0.8-70
## [25] munsell_0.4.3        broom_0.4.4          compiler_3.5.0
## [28] modelr_0.1.2         pkgconfig_2.0.1      mnormt_1.5-5
## [31] htmltools_0.3.6      fastGHQuad_0.2       tidyselect_0.2.4
## [34] codetools_0.2-15     viridisLite_0.3.0    crayon_1.3.4
## [37] withr_2.1.2          MASS_7.3-50          grid_3.5.0
## [40] nlme_3.1-137         jsonlite_1.5         xtable_1.8-2
## [43] gtable_0.2.0         scales_0.5.0.9000    cli_1.0.0
## [46] stringi_1.2.2        reshape2_1.4.3       robustbase_0.93-0
## [49] xml2_1.2.0           tools_3.5.0          glue_1.2.0
## [52] DEoptimR_1.0-8       hms_0.4.2            parallel_3.5.0
## [55] yaml_2.1.19          colorspace_1.3-2     rvest_0.3.2
## [58] knitr_1.20           bindr_0.1.1          haven_1.1.1
```