Supplement 12

Experiment 2 – Descriptive plots of the stimulus-response relationship for the SPARS, SRS, and pain NRS

Peter Kamerman and Tory Madden

21 June 2018

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This script assesses and contrasts the stimulus-response characteristics of the SPARS, sensation rating scale (SRS), and pain NRS, through the use of descriptive plots.

The three scales measure the following ranges of somatic sensation:

- pain NRS: 0 (no pain) to 100 (worst pain you can imagine)
- SRS: 0 (no sensation) to 100 (pain)
- SPARS: -50 (no sensation), 0 (pain threshold), +50 (worst pain you can imagine)

Because the stimulus range was centred on the pre-determined pain threshold of each participant (compared to the fixed range of intensities used in Trial A), all analyses use the rank order of the nine stimulus intensities each participant was exposed to rather than the absolute intensities of the stimuli used.

The experimental design involved exposing each participant to four successive experimental blocks of 27 trials (laser stimulations) each for each of the three measurement scales. The sequence of stimulus intensities used within each block was pre-determined, and differed between blocks. The order of in which the measurement scales were assessed was randomized, but for convenience of reporting, the plots are always shown in the order: pain NRS_P, SRS, and SPARS.

Import and inspect data

```
# Import
data <- read_rds('./data-cleaned/SPARS_B.rds')
# Inspect
glimpse(data)</pre>
```

```
## Observations: 2,256
## Variables: 10
## $ PID
                      <chr> "ID01", "ID01", "ID01", "ID01", "ID01", "ID0...
## $ scale
                      <chr> "SPARS", "SPARS", "SPARS", "SPARS", "SPARS", ...
## $ block number
                      <int> 2, 2, 2, 4, 4, 4, 6, 6, 6, 8, 8, 8, 11, 11, ...
                     <int> 9, 15, 23, 7, 20, 25, 9, 18, 22, 3, 17, 23, ...
## $ trial number
                      <dbl> 2.25, 2.25, 2.25, 2.25, 2.25, 2.25, 2.25, 2.25, 2....
## $ intensity
                     <chr> "2.25", "2.25", "2.25", "2.25", "2.25", "2.25", "2.2...
## $ intensity_char
                     ## $ intensity rank
                      <dbl> -31, -20, -48, -48, -21, -23, -48, -45, -47,...
## $ rating
                      <dbl> 19, 30, 2, 2, 29, 27, 2, 5, 3, 0, 1, 3, 50, ...
## $ rating positive
## $ rating_equivalent <dbl> -31, -20, -48, -48, -21, -23, -48, -45, -47,...
```

Clean and transform data

```
#
                                           #
#
                   Clean
                                          #
data %<>%
 # Select required columns
 select(PID, scale, block_number, intensity_rank, rating, rating_equivalent) %>%
 # Recode scales to CNRS NP = NRS, NRS NP = SRS
 mutate(scale = case when(
    scale == 'NRS NP' ~ 'SRS',
    scale == 'CNRS_P' ~ 'NRS';
    scale == 'SPARS' ~ 'SPARS'
 ))
#
#
            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 <- 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
#
# Add sequential block numbers, tagged by scale
data %<>%
   # Group by ID and scale
   group_by(PID, scale) %>%
   # Arrange blocks
   arrange(block number) %>%
   # Add sequential
   mutate(block sequential = dense_rank(block number),
          block_sequential = pasteO(scale, ' - ', block_sequential)) %>%
   ungroup()
# Calculate the participant trimean for each scale (using 'rating equivalent')
data_tm <- data %>%
 group_by(PID, scale, intensity rank) %>%
 summarise(tri_mean = tri.mean(rating_equivalent)) %>%
 ungroup()
```

Exploratory plots

Raw intensity ratings (participant-level)

Data are shown on their original scales: pain NRS: 0 to 100, SRS: 0 to 100, and SPARS: -50 to +50.

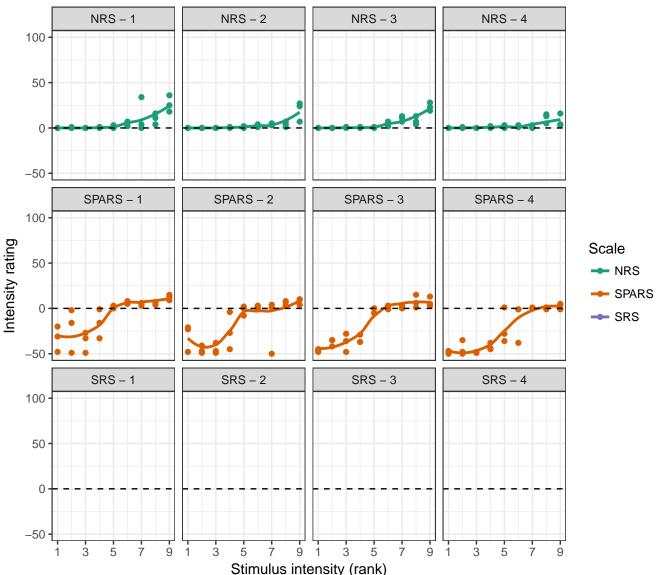
```
# Generate plots
plot raw <- data %>%
  group_by(PID) %>%
 nest() %>%
 mutate(plots = map2(.x = data,
                      y = PID,
                      ~ ggplot(data = .x) +
                          aes(x = intensity rank,
                              y = rating,
                              colour = scale,
                              fill = scale) +
                          geom_smooth(se = FALSE) +
                          geom_point() +
                          scale_y_continuous(limits = c(-50, 100)) +
                          scale_x_continuous(breaks = c(1, 3, 5, 7, 9)) +
                          scale_fill_brewer(name = 'Scale',
                                             type = 'qual',
                                             palette = 'Dark2') +
                          scale_colour_brewer(name = 'Scale',
                                               type = 'qual',
                                               palette = 'Dark2') +
                          labs(title = paste0(.y, ': Raw participant-level stimulus-respong)
```

```
subtitle = 'Plots are conditioned on measurement scale and
x = 'Stimulus intensity (rank)',
y = 'Intensity rating') +
geom_hline(yintercept = 0, linetype = 2) +
facet_wrap(~ block_sequential)))

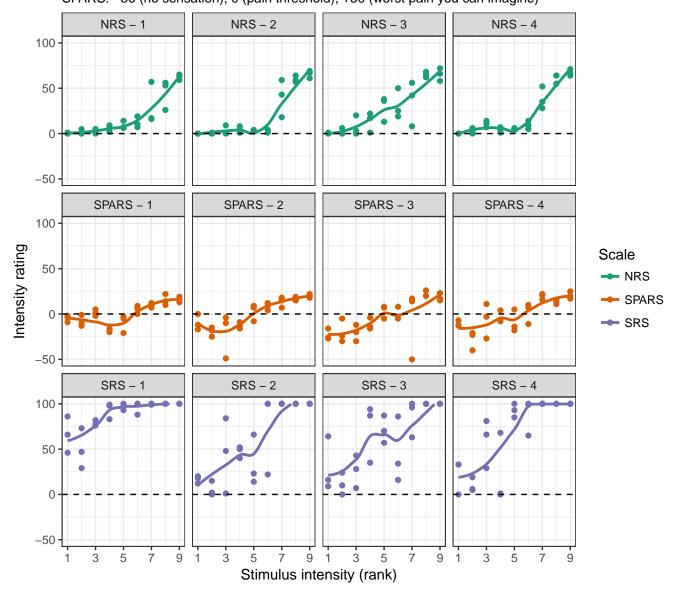
# Print plots
walk(plot_raw$plots, ~ print(.x))
```

ID01: Raw participant-level stimulus-response ratings on the pain NRS, SRS an

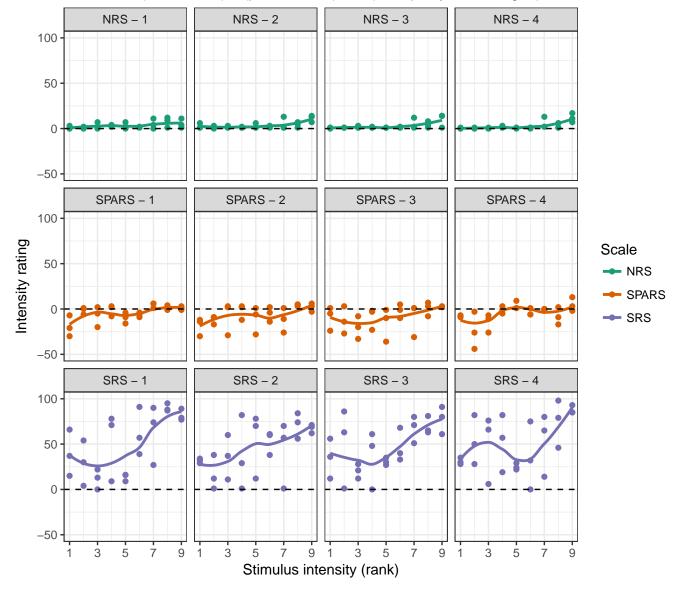
Plots are conditioned on measurement scale and experimental block pain NRS: 0 (no pain) to 100 (worst pain you can imagine) SRS: 0 (no sensation) to 100 (pain)



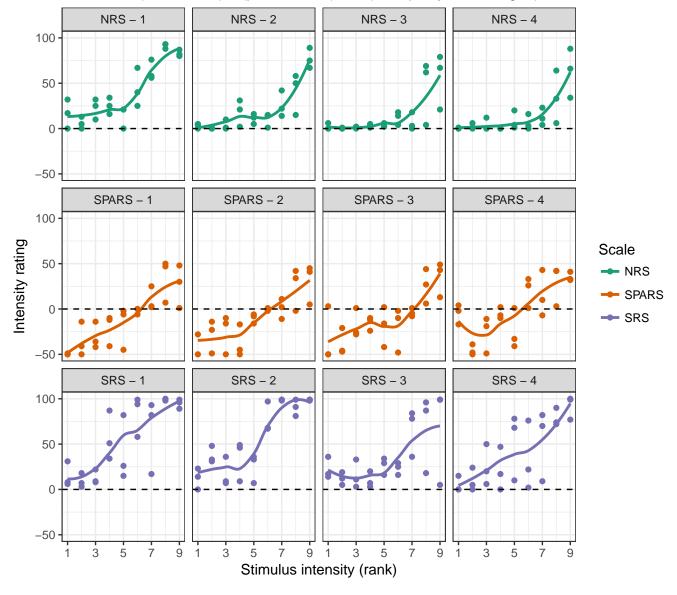
ID02: Raw participant-level stimulus-response ratings on the pain NRS, SRS an



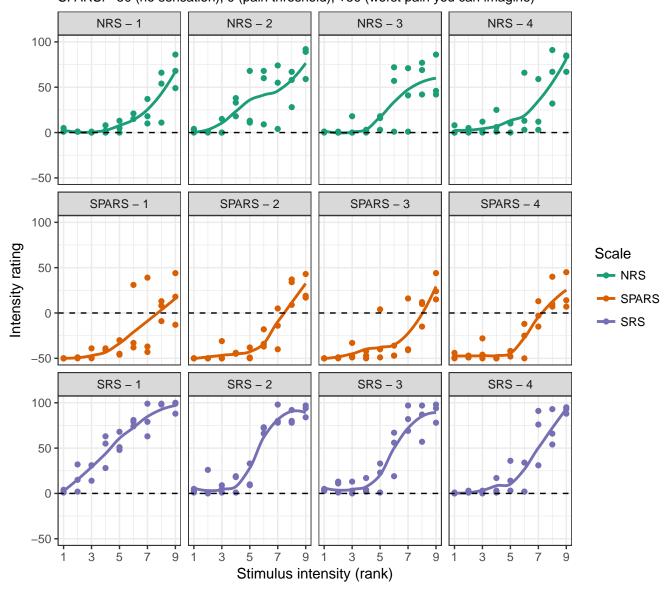
ID03: Raw participant-level stimulus-response ratings on the pain NRS, SRS an



ID04: Raw participant-level stimulus-response ratings on the pain NRS, SRS an

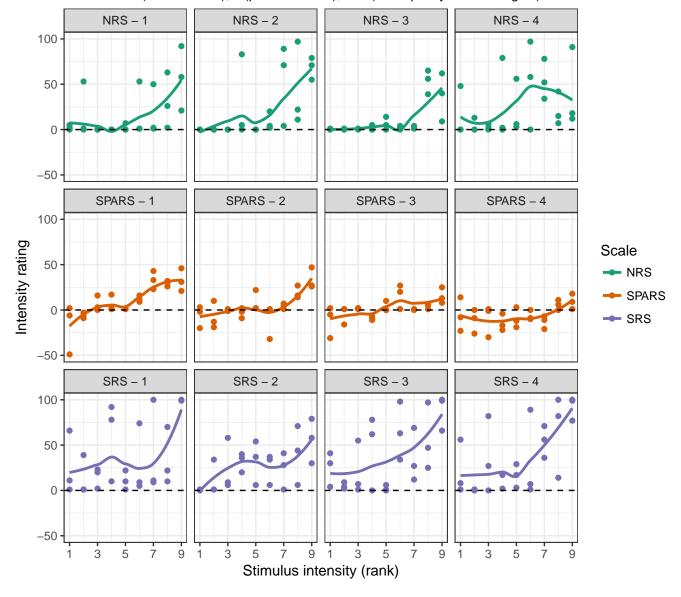


ID05: Raw participant-level stimulus-response ratings on the pain NRS, SRS an



ID06: Raw participant-level stimulus-response ratings on the pain NRS, SRS an

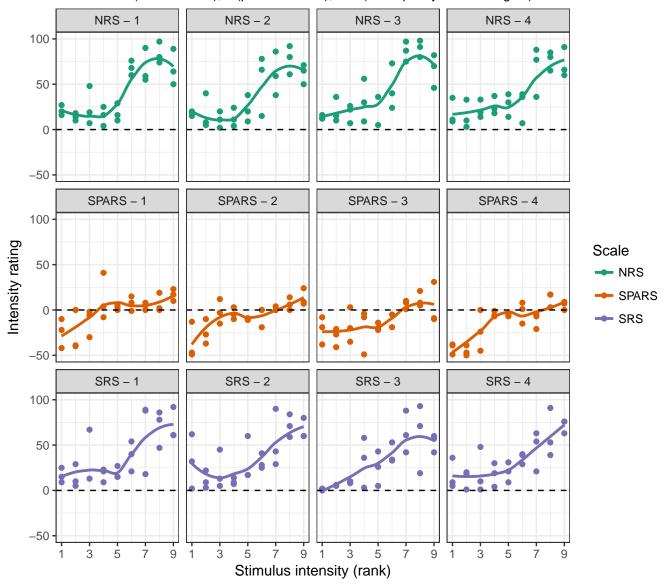
Plots are conditioned on measurement scale and experimental block pain NRS: 0 (no pain) to 100 (worst pain you can imagine) SRS: 0 (no sensation) to 100 (pain)



ID07: Raw participant-level stimulus-response ratings on the pain NRS, SRS an

Plots are conditioned on measurement scale and experimental block pain NRS: 0 (no pain) to 100 (worst pain you can imagine) SRS: 0 (no sensation) to 100 (pain)

SPARS: -50 (no sensation), 0 (pain threshold), +50 (worst pain you can imagine)



Equivalent units ratings (participant-level)

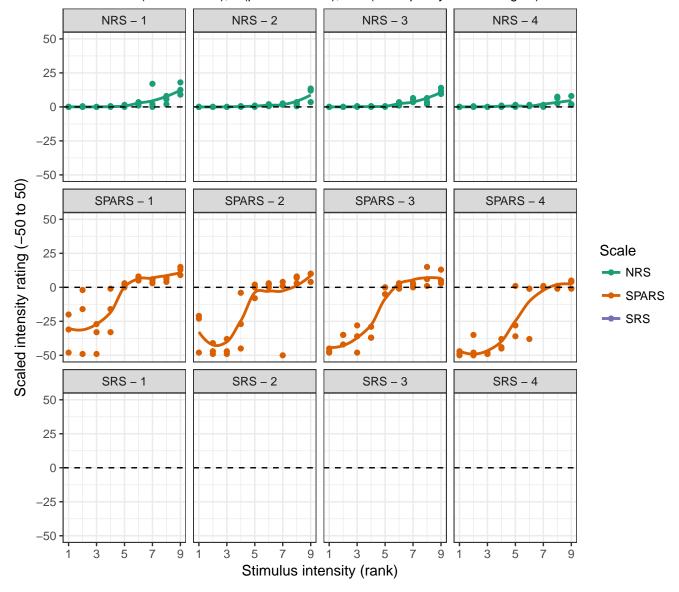
Raw scores for the pain NRS and SRS scales (both rated on 0 to 100 scales) were converted to SPARS equivalent ranges (-50 to +50). The scaling of the pain NRS and SRS were as follows:

- pain NRS: raw 0 to 100 scores were converted to a 0 to 50 range by dividing 2, such that after the scaling, 0 = no pain and 50 = worst pain you can imagine. This is equivalent to the positive range of the SPARS.
- SRS: raw 0 to 100 scores were converted to a -50 to 0 range by subtracting 100, and then dividing 2, such that after the scaling, -50 = *no sensation* and 0 = *pain*. This is equivalent to the negative range of the SPARS.

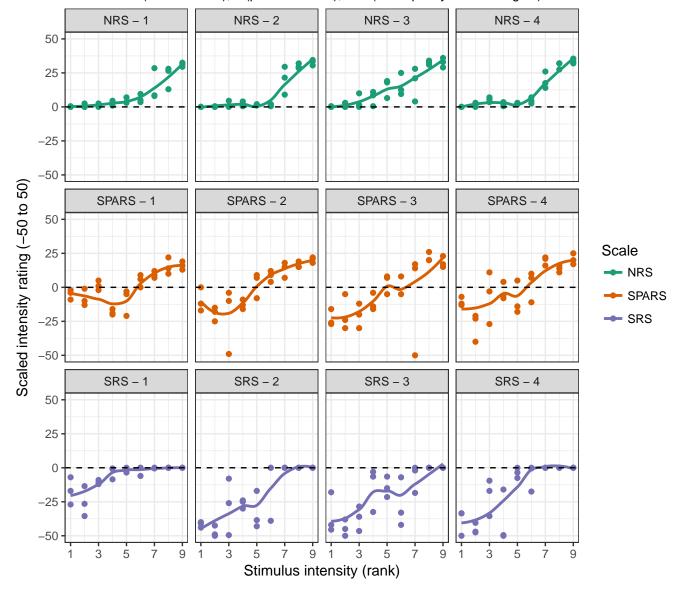
```
~ ggplot(data = .x) +
                          aes(x = intensity_rank,
                              y = rating_equivalent,
                              colour = scale,
                              fill = scale) +
                          geom_smooth(se = FALSE) +
                          geom_point() +
                          scale_y_continuous(limits = c(-50, 50)) +
                          scale_x_continuous(breaks = c(1, 3, 5, 7, 9)) +
                          scale_fill_brewer(name = 'Scale',
                                            type = 'qual',
                                            palette = 'Dark2') +
                          scale_colour_brewer(name = 'Scale',
                                              type = 'qual',
                                              palette = 'Dark2') +
                          labs(title = paste0(.y, ': Scaled participant-level stimulus-res
                               subtitle = 'Plots are conditioned on measurement scale and
                               x = 'Stimulus intensity (rank)',
                               y = 'Scaled intensity rating (-50 to 50)') +
                          geom_hline(yintercept = 0, linetype = 2) +
                          facet_wrap(~ block_sequential)))
# Print plots
walk(plot equi$plots, ~ print(.x))
```

ID01: Scaled participant-level stimulus-response ratings on the pain NRS, SRS

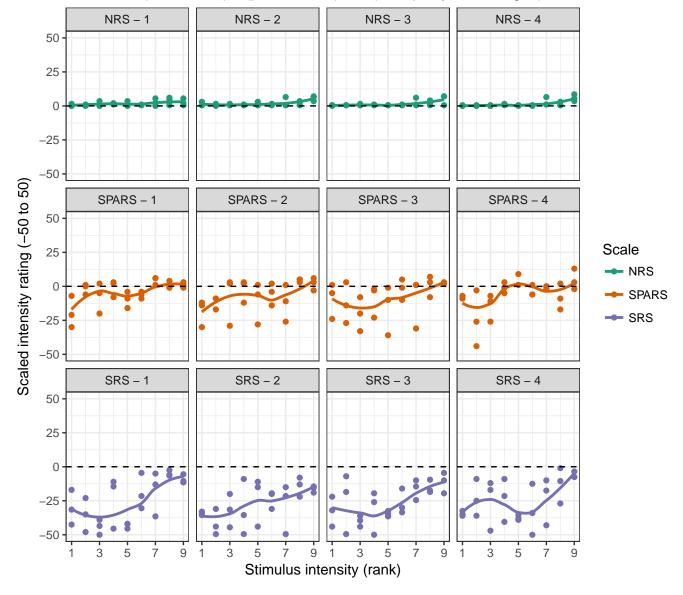
Plots are conditioned on measurement scale and experimental block pain NRS: 0 (no pain) to 50 (worst pain you can imagine) SRS: –50 (no sensation) to 0 (pain)



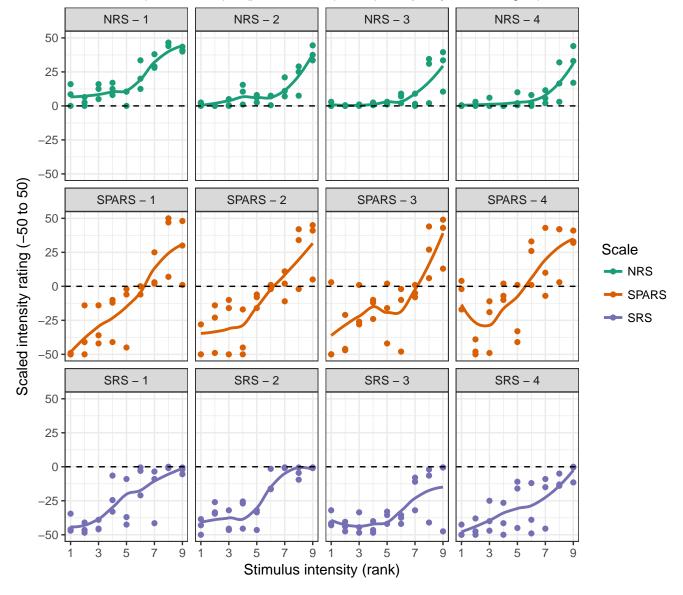
ID02: Scaled participant-level stimulus-response ratings on the pain NRS, SRS



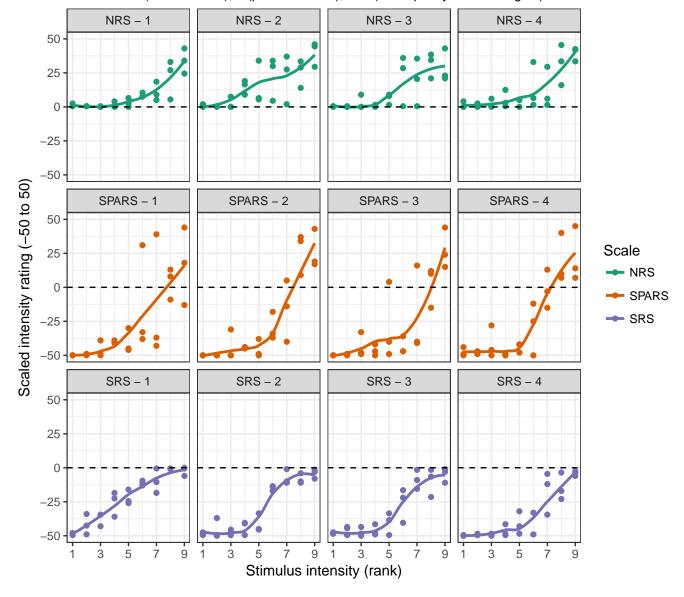
ID03: Scaled participant-level stimulus-response ratings on the pain NRS, SRS



ID04: Scaled participant-level stimulus-response ratings on the pain NRS, SRS

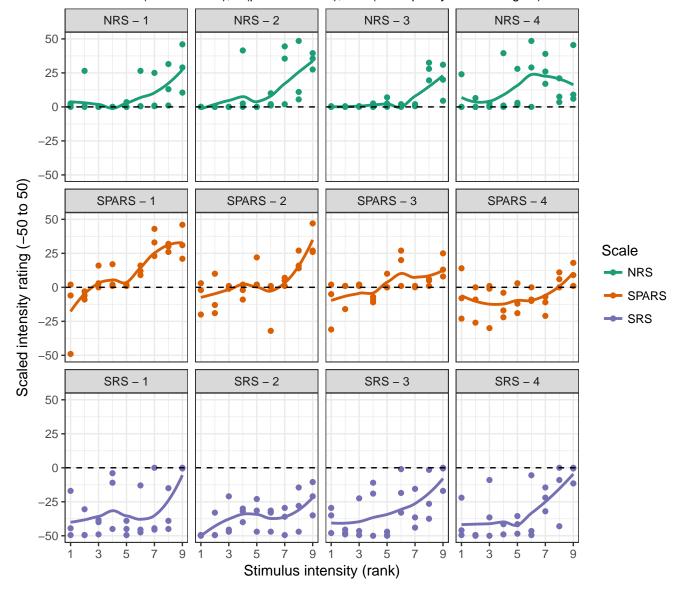


ID05: Scaled participant-level stimulus-response ratings on the pain NRS, SRS



ID06: Scaled participant-level stimulus-response ratings on the pain NRS, SRS

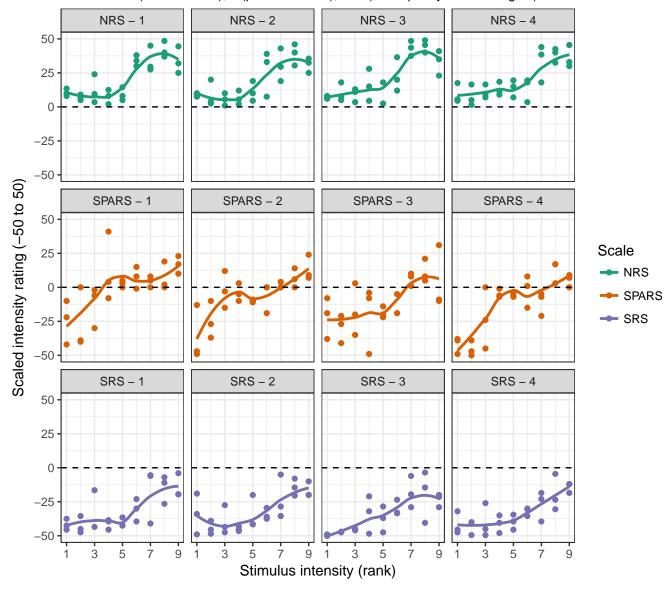
Plots are conditioned on measurement scale and experimental block pain NRS: 0 (no pain) to 50 (worst pain you can imagine) SRS: –50 (no sensation) to 0 (pain)



ID07: Scaled participant-level stimulus-response ratings on the pain NRS, SRS

Plots are conditioned on measurement scale and experimental block pain NRS: 0 (no pain) to 50 (worst pain you can imagine) SRS: –50 (no sensation) to 0 (pain)

SPARS: -50 (no sensation), 0 (pain threshold), +50 (worst pain you can imagine)



Tukey trimean plots (participant-level)

For each participant, we calculated the Tukey trimean of the intensity rating at each stimulus intensity and for each of the measurement scales. The scaled versions of the pain NRS and SRS were used.

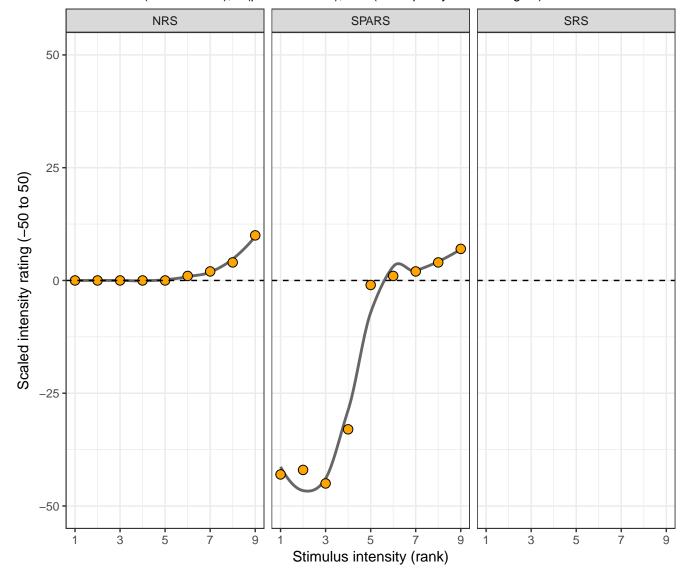
```
geom_smooth(method = 'loess',
                                       se = FALSE,
                                       colour = '#666666') +
                           geom_point(shape = 21,
                                      size = 3,
                                      fill = 'orange') +
                           scale_fill_brewer(name = 'Scale',
                                             type = 'qual',
                                             palette = 'Dark2') +
                           scale_colour_brewer(name = 'Scale',
                                               type = 'qual',
                                               palette = 'Dark2') +
                           scale_x_continuous(breaks = c(1, 3, 5, 7, 9)) +
                           scale_y_continuous(limits = c(-50, 50)) +
                           labs(title = paste0(.y, ': Scaled participant-level stimulus-re
                                subtitle = 'Orange points: Tukey trimeans | Grey curve: lo
                                x = 'Stimulus intensity (rank)',
                                y = 'Scaled intensity rating (-50 to 50)') +
                           geom_hline(yintercept = 0, linetype = 2) +
                           facet_wrap(~ scale, ncol = 3)))
# Print plots
walk(plot_tm$plot, ~ print(.x))
```

ID01: Scaled participant-level stimulus-response rating Tukey trimeans on the participant-level stimulus-response rating Tukey trimeans response rating Tukey trimeans respo

Orange points: Tukey trimeans | Grey curve: loess curve

Plots are conditioned on the three scales pain NRS: 0 (no pain) to 50 (worst pain you can imagine)

SRS: -50 (no sensation) to 0 (pain)

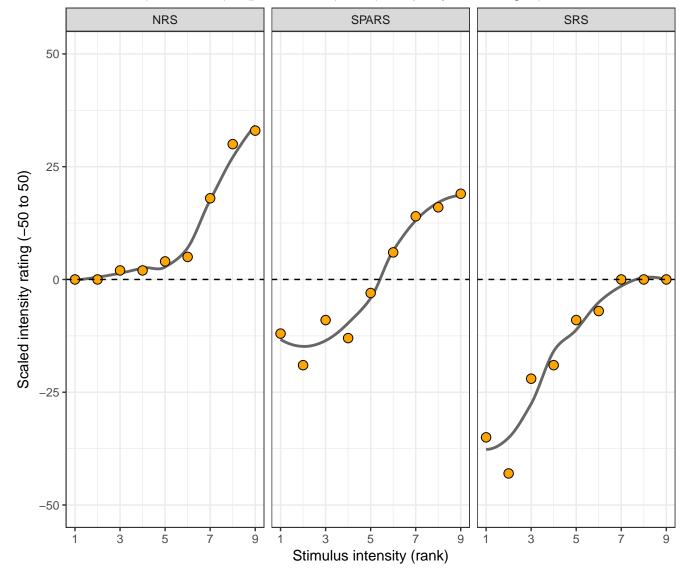


ID02: Scaled participant-level stimulus-response rating Tukey trimeans on the participant-level stimulus-response rating Tukey trimeans response rating Tukey trimeans respo

Orange points: Tukey trimeans | Grey curve: loess curve

Plots are conditioned on the three scales pain NRS: 0 (no pain) to 50 (worst pain you can imagine)

SRS: –50 (no sensation) to 0 (pain) SPARS: –50 (no sensation), 0 (pain threshold), +50 (worst pain you can imagine)

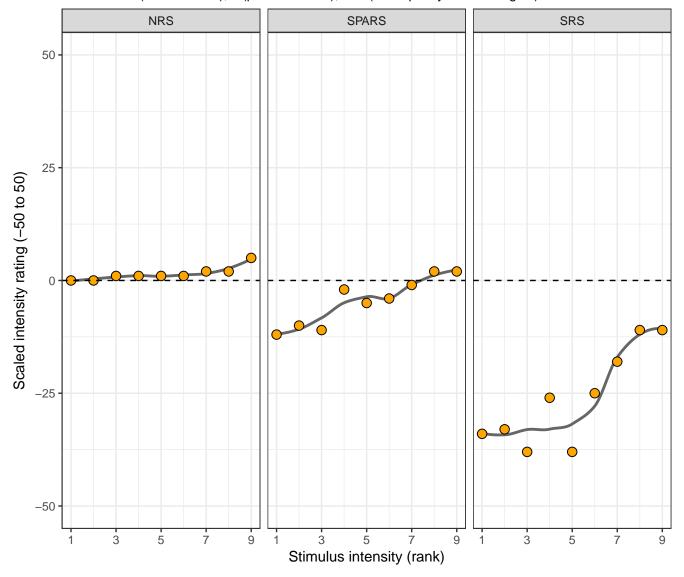


ID03: Scaled participant-level stimulus-response rating Tukey trimeans on the participant-level stimulus-response rating Tukey trimeans response rating Tukey trimeans respo

Orange points: Tukey trimeans | Grey curve: loess curve

Plots are conditioned on the three scales pain NRS: 0 (no pain) to 50 (worst pain you can imagine)

SRS: -50 (no sensation) to 0 (pain)

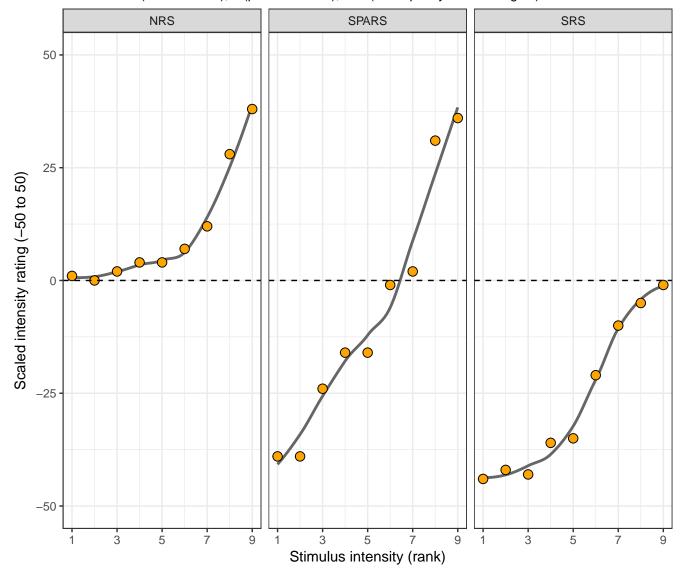


ID04: Scaled participant-level stimulus-response rating Tukey trimeans on the participant-level stimulus-response rating Tukey trimeans of the participant-level stimulus-response rating Tukey trimeans on the participant-level stimulus-response rating Tukey trimeans response rating Tukey trimeans respo

Orange points: Tukey trimeans | Grey curve: loess curve

Plots are conditioned on the three scales pain NRS: 0 (no pain) to 50 (worst pain you can imagine)

SRS: –50 (no sensation) to 0 (pain) SPARS: –50 (no sensation), 0 (pain threshold), +50 (worst pain you can imagine)

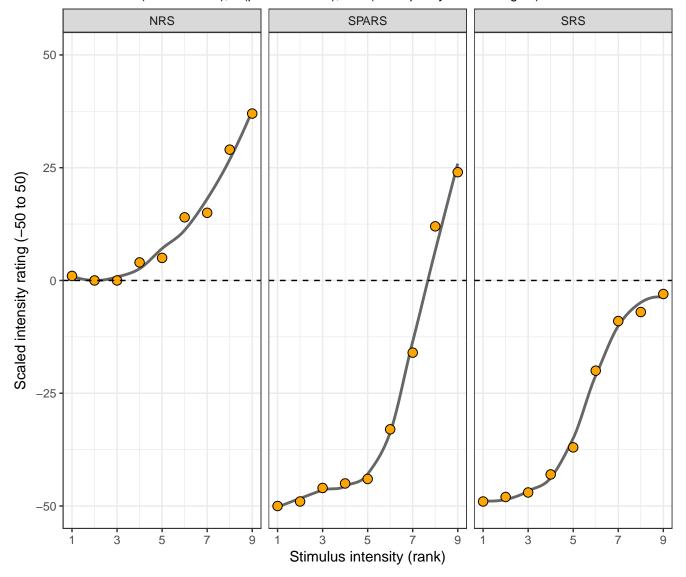


ID05: Scaled participant-level stimulus-response rating Tukey trimeans on the participant-level stimulus-response rating Tukey trimeans of the participant-level stimulus-response rating Tukey trimeans of the participant-level stimulus-response rating Tukey trimeans response rating Tukey trimeans respo

Orange points: Tukey trimeans | Grey curve: loess curve

Plots are conditioned on the three scales pain NRS: 0 (no pain) to 50 (worst pain you can imagine)

SRS: –50 (no sensation) to 0 (pain) SPARS: –50 (no sensation), 0 (pain threshold), +50 (worst pain you can imagine)

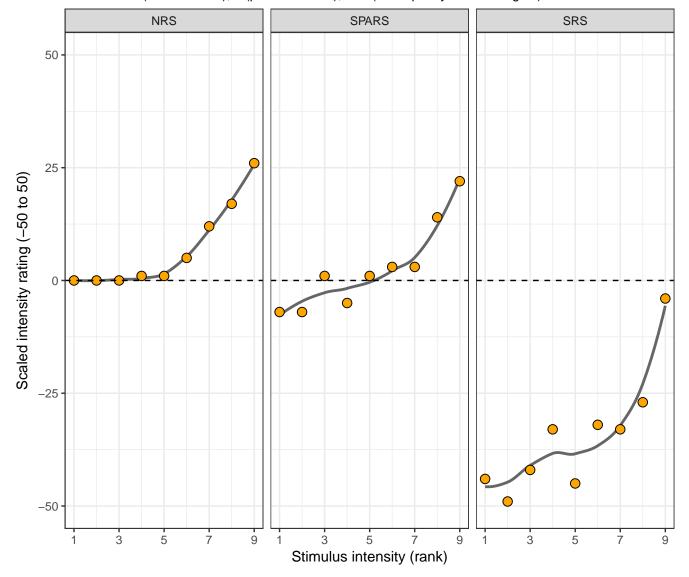


ID06: Scaled participant-level stimulus-response rating Tukey trimeans on the participant-level stimulus-response rating Tukey trimeans of the participant-level stimulus-response rating Tukey trimeans on the participant-level stimulus-response rating Tukey trimeans response rating Tukey trimeans respo

Orange points: Tukey trimeans | Grey curve: loess curve

Plots are conditioned on the three scales pain NRS: 0 (no pain) to 50 (worst pain you can imagine)

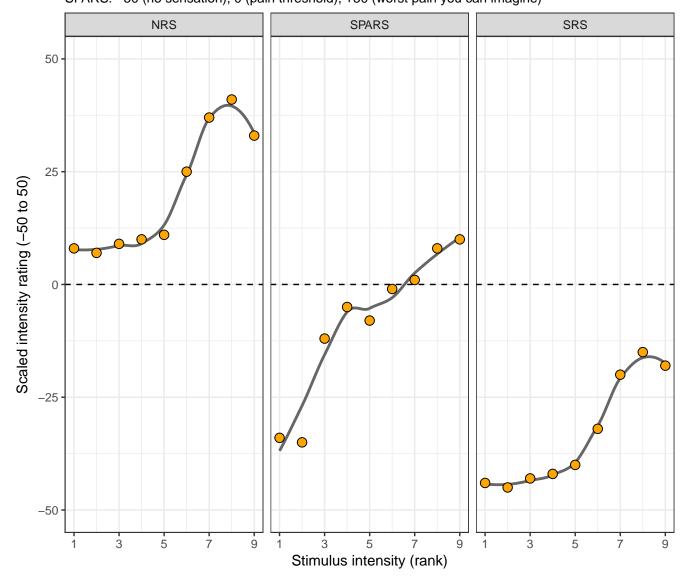
SRS: -50 (no sensation) to 0 (pain)



ID07: Scaled participant-level stimulus-response rating Tukey trimeans on the participant-level stimulus-response rating Tukey trimeans response rating Tukey trimeans respo

Orange points: Tukey trimeans | Grey curve: loess curve Plots are conditioned on the three scales pain NRS: 0 (no pain) to 50 (worst pain you can imagine) SRS: –50 (no sensation) to 0 (pain)

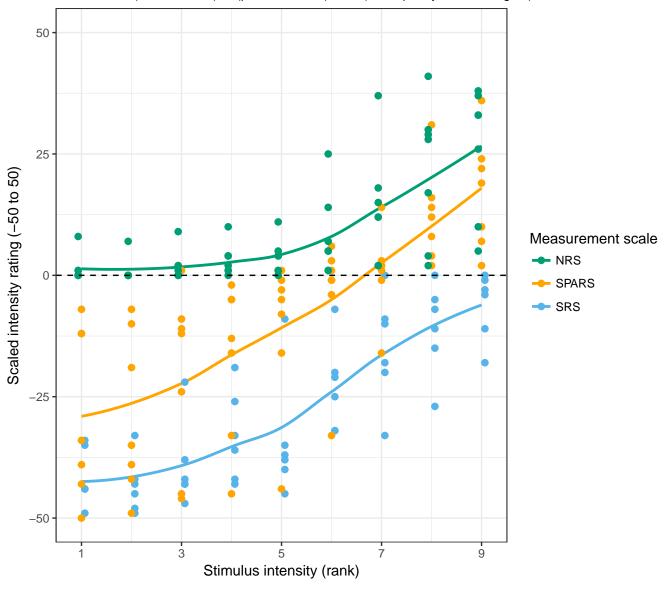
SPARS: -50 (no sensation), 0 (pain threshold), +50 (worst pain you can imagine)



Summary plots (group-level)

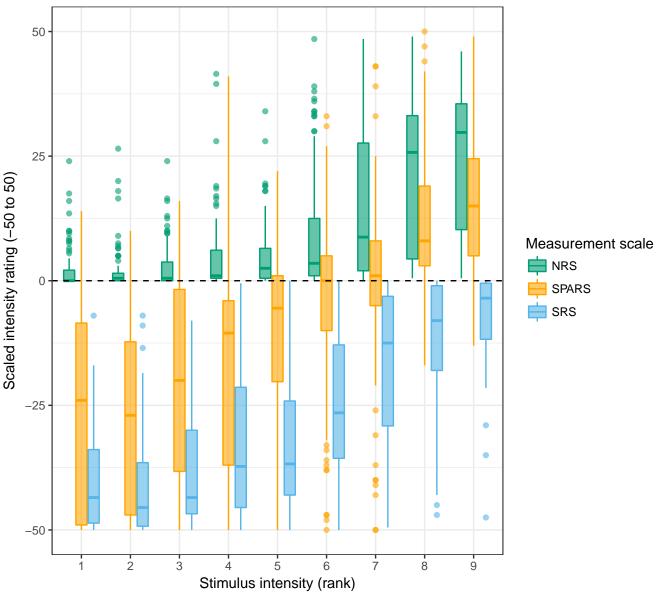
Scaled group-level stimulus-response ratings on the pain NRS, SRS and SPARS

Points (dodged for clarity): Tukey trimeans | Curves: loess lines pain NRS: 0 (no pain) to 50 (worst pain you can imagine) SRS: –50 (no sensation) to 0 (pain) SPARS: –50 (no sensation), 0 (pain threshold), +50 (worst pain you can imagine)



Scaled group-level stimulus-response ratings on the pain NRS, SRS and SPARS

pain NRS: 0 (no pain) to 50 (worst pain you can imagine) SRS: -50 (no sensation) to 0 (pain)



Summary

In general, the ratings provided on the SPARS were less than those provided on the pain NRS (even at higher stimulus intensities), and greater than those provided on the SRS (even at lower stimulus intensities). This indicates that the extra dimensionality of the SPARS compared to the other two scales (the SPARS allowing the ratings of stimulus intensities from *no sensation* to *worst pain imaginable*) leads to a compression of the numeric value of the ratings into a narrower band around 0 (pain threshold) compared to the two polar scales (measuring either noxious or non-noxious stimulus ranges).

However, the trend lines indicate that the SPARS scale is reporting the same information as each of the other two scales in their respective domains (sensation of noxious and non-noxious stimuli), but the sensitivity of the SPARS, in each of the two domains, may be lower than they achieved from the two domain-specific scales.

(Please also see: *outputs/6B-scale-agreement.html*)

Session information

```
sessionInfo()
```

```
## 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/en_GB.UTF-8
##
## attached base packages:
##
   [1] stats
                 graphics grDevices utils
                                                datasets
                                                        methods
                                                                    base
##
## other attached packages:
##
    [1] bindrcpp 0.2.2
                           forcats 0.3.0
                                               stringr 1.3.1
    [4] dplyr_0.7.5
                           purrr_0.2.5
                                               readr 1.1.1
##
    [7] tidyr 0.8.1
                           tibble 1.4.2
                                               ggplot2_2.2.1.9000
##
   [10] tidyverse 1.2.1
##
                           magrittr 1.5
##
                          (and not attached):
##
  loaded via a namespace
##
    [1] tidyselect_0.2.4
                           reshape2_1.4.3
                                               haven_1.1.1
    [4] lattice 0.20-35
##
                           colorspace_1.3-2
                                               htmltools 0.3.6
                                               pillar 1.2.3
   [7] yaml 2.1.19
                           rlang 0.2.1
##
## [10] foreign 0.8-70
                           glue 1.2.0
                                               withr 2.1.2
## [13] RColorBrewer_1.1-2 modelr_0.1.2
                                               readxl_1.1.0
## [16] bindr 0.1.1
                           plyr 1.8.4
                                               munsell 0.4.3
## [19] gtable 0.2.0
                           cellranger 1.1.0
                                               rvest_0.3.2
## [22] psych_1.8.4
                           evaluate_0.10.1
                                               labeling_0.3
## [25] knitr_1.20
                           parallel_3.5.0
                                               broom_0.4.4
## [28] Rcpp 0.12.17
                           scales 0.5.0.9000
                                               backports 1.1.2
```

##	[31]	jsonlite_1.5	mnormt_1.5-5	hms_0.4.2
##	[34]	digest_0.6.15	stringi_1.2.2	grid_3.5.0
##	[37]	rprojroot_1.3-2	cli_1.0.0	tools_3.5.0
##	[40]	lazyeval_0.2.1	crayon_1.3.4	pkgconfig_2.0.1
##	[43]	xml2_1.2.0	<pre>lubridate_1.7.4</pre>	assertthat_0.2.0
##	[46]	rmarkdown_1.9	httr_1.3.1	rstudioapi_0.7
##	[49]	R6_2.2.2	nlme_3.1-137	compiler_3.5.0