F02 mci emotion pre ratings.R

2020-09-22

```
## MCI EMO PRE-RATINGS SCRIPT ##
# Pre-ratings of cloze probability, plausibility, metaphoricity, and imageability of the context stories were conducted
# on five-point rating scales. Script computes analyses of variances testing for potential differences in these ratings
# between semantic conditions. Additionally, pairwise t-tests test differences between each pair of semantic conditions
# (violation - intuitive, MCI - intuitive, MCI - violation). The Bonferroni-Holm-correction was applied to control for
# multiple comparisons.
# Load packages
library(tidyverse)
                   # Version 1.3.0
library(magrittr) # Version 1.5
library(emmeans)
                  # version 1.4.8
# Load pre-rating data from SPSS file
pilot <- haven::read_sav("FB/gesamt_2.sav")</pre>
# Trim whitespace
pilot %<>% mutate(KonzeptNr = KonzeptNr %>% trimws() %>% as.numeric(),
                  VerbBedingung = VerbBedingung %>% trimws())
# Rename conditions
pilot %<>% mutate(semantics = factor(VerbBedingung, levels = c("neutral", "sem", "mci"),
                                    labels = c("int", "vio", "mci")))
# Summarize by participants
avgs <- pilot %>%
  group_by(VP, semantics) %>%
```

```
summarise(clozeprob = mean(Frage1),
           plausibility = mean(Frage2),
           metaphoricity = mean(Frage3),
           imageability = mean(Frage4)) %>%
  mutate(VP = factor(VP))
## 'summarise()' regrouping output by 'VP' (override with '.groups' argument)
## ANOVAs ## -----
# Semantics is a within subjects factor; data are fully balanced
# Cloze probability
summary(anova cloze <- aov(clozeprob ~ semantics + Error(VP/semantics), data = avgs))</pre>
##
## Error: VP
            Df Sum Sq Mean Sq F value Pr(>F)
## Residuals 19 7.689 0.4047
## Error: VP:semantics
            Df Sum Sq Mean Sq F value Pr(>F)
## semantics 2 16.74 8.369 17.66 3.78e-06 ***
## Residuals 38 18.01 0.474
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
mean(anova_cloze$`VP:semantics`$residuals^2) # Mean squared error of the effect
## [1] 0.450286
# Plausibility
summary(anova_plausibility <- aov(plausibility ~ semantics + Error(VP/semantics), data = avgs))</pre>
##
## Error: VP
```

```
Df Sum Sq Mean Sq F value Pr(>F)
## Residuals 19 5.338 0.281
## Error: VP:semantics
            Df Sum Sq Mean Sq F value Pr(>F)
## semantics 2 13.55 6.777
                               10.74 0.000201 ***
## Residuals 38 23.98 0.631
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
mean(anova_plausibility$`VP:semantics`$residuals^2) # Mean squared error of the effect
## [1] 0.5994464
# Imageability
summary(anova_imageability <- aov(imageability ~ semantics + Error(VP/semantics), data = avgs))</pre>
##
## Error: VP
            Df Sum Sq Mean Sq F value Pr(>F)
## Residuals 19 6.643 0.3497
## Error: VP:semantics
            Df Sum Sq Mean Sq F value Pr(>F)
## semantics 2 12.43 6.215 14.61 1.96e-05 ***
## Residuals 38 16.16 0.425
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
mean(anova_imageability$`VP:semantics`$residuals^2) # Mean squared error of the effect
## [1] 0.4041053
# Metaphoricity
summary(anova_metaphoricity <- aov(metaphoricity ~ semantics + Error(VP/semantics), data = avgs))</pre>
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##
## Error: VP
           Df Sum Sq Mean Sq F value Pr(>F)
## Residuals 19 62.24 3.276
## Error: VP:semantics
           Df Sum Sq Mean Sq F value Pr(>F)
## semantics 2 5.142 2.5712 8.988 0.000636 ***
## Residuals 38 10.870 0.2861
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
mean(anova metaphoricity$\text{VP:semantics}\text{*residuals}^2) # Mean squared error of the effect
## [1] 0.2717558
## PAIRWISE TESTS ## ------
# Cloze probability
(pairwise_clozeprob <- anova_cloze %>% emmeans(specs = pairwise ~ semantics) %>% summary(adjust = "holm"))
## Note: re-fitting model with sum-to-zero contrasts
## $emmeans
## semantics emmean
                    SE df lower.CL upper.CL
## int
            3.47 0.15 56.7
                               3.10
                                        3.84
## vio
             2.65 0.15 56.7
                               2.28
                                        3.02
## mci
              2.19 0.15 56.7
                               1.82
                                        2.56
## Warning: EMMs are biased unless design is perfectly balanced
## Confidence level used: 0.95
## Conf-level adjustment: bonferroni method for 3 estimates
##
## $contrasts
## contrast estimate
                       SE df t.ratio p.value
## int - vio 0.814 0.218 38 3.739 0.0012
## int - mci 1.278 0.218 38 5.870 <.0001
```

```
## vio - mci 0.464 0.218 38 2.131 0.0396
## P value adjustment: holm method for 3 tests
# Plausibility
(pairwise plausibility <- anova plausibility %>% emmeans(specs = pairwise ~ semantics) %>% summary(adjust = "holm"))
## Note: re-fitting model with sum-to-zero contrasts
## $emmeans
## semantics emmean
                      SE
                         df lower.CL upper.CL
## int
               2.84 0.16 51.7
                                 2.44
                                          3.24
## vio
               2.13 0.16 51.7
                                 1.73
                                          2.53
## mci
              1.69 0.16 51.7
                                 1.29
                                          2.08
## Warning: EMMs are biased unless design is perfectly balanced
## Confidence level used: 0.95
## Conf-level adjustment: bonferroni method for 3 estimates
## $contrasts
## contrast estimate
                         SE df t.ratio p.value
## int - vio 0.710 0.251 38 2.826 0.0150
## int - mci 1.154 0.251 38 4.594 0.0001
## vio - mci 0.444 0.251 38 1.769 0.0850
## P value adjustment: holm method for 3 tests
# Imageability
(pairwise imageability <- anova imageability %>% emmeans(specs = pairwise ~ semantics) %>% summary(adjust = "holm"))
## Note: re-fitting model with sum-to-zero contrasts
## $emmeans
## semantics emmean
                       SE df lower.CL upper.CL
## int
               3.65 0.141 56.5
                                  3.30
                                           4.00
## vio
             2.99 0.141 56.5
                                  2.64
                                           3.34
## mci
               2.54 0.141 56.5
                                  2.19
                                           2.89
```

```
## Warning: EMMs are biased unless design is perfectly balanced
## Confidence level used: 0.95
## Conf-level adjustment: bonferroni method for 3 estimates
## $contrasts
## contrast estimate
                         SE df t.ratio p.value
## int - vio 0.660 0.206 38 3.199 0.0056
## int - mci 1.108 0.206 38 5.373 <.0001
## vio - mci 0.448 0.206 38 2.174 0.0360
## P value adjustment: holm method for 3 tests
# Metaphoricity
(pairwise metaphoricity <- anova metaphoricity %>% emmeans(specs = pairwise ~ semantics) %>% summary(adjust = "holm"))
## Note: re-fitting model with sum-to-zero contrasts
## $emmeans
## semantics emmean
                       SE df lower.CL upper.CL
## int
               2.30 0.253 25.8
                                   1.65
                                           2.95
## vio
               2.49 0.253 25.8
                                   1.84
                                           3.13
## mci
               2.99 0.253 25.8
                                   2.34
                                           3.64
## Warning: EMMs are biased unless design is perfectly balanced
## Confidence level used: 0.95
## Conf-level adjustment: bonferroni method for 3 estimates
## $contrasts
## contrast estimate
                         SE df t.ratio p.value
## int - vio -0.189 0.169 38 -1.115 0.2720
## int - mci -0.693 0.169 38 -4.100 0.0006
## vio - mci -0.505 0.169 38 -2.985 0.0099
## P value adjustment: holm method for 3 tests
```

System specs and package versions sessionInfo()

```
## R version 4.0.2 (2020-06-22)
## Platform: x86_64-apple-darwin17.0 (64-bit)
## Running under: macOS Catalina 10.15.6
## Matrix products: default
          /System/Library/Frameworks/Accelerate.framework/Versions/A/Frameworks/vecLib.framework/Versions/A/libBLAS.dylib
## LAPACK: /Library/Frameworks/R.framework/Versions/4.0/Resources/lib/libRlapack.dylib
##
## locale:
## [1] en US.UTF-8/en US.UTF-8/en US.UTF-8/c/en US.UTF-8/en US.UTF-8
##
## attached base packages:
## [1] stats
                 graphics grDevices datasets utils
                                                         methods
                                                                   base
##
## other attached packages:
   [1] emmeans_1.4.8
                        magrittr_1.5
                                        forcats_0.5.0
                                                        stringr_1.4.0
                                                                        dplyr_1.0.0
                                                                                         purrr_0.3.4
                                                                                                         readr_1.3.1
                                                                                                                         tidyr_1.1.0
## [10] ggplot2_3.3.2
                       tidyverse_1.3.0
##
## loaded via a namespace (and not attached):
## [1] tidyselect_1.1.0 xfun_0.16
                                          haven_2.3.1
                                                           colorspace_1.4-1 vctrs_0.3.2
                                                                                              generics_0.0.2
                                                                                                               htmltools_0.5.0
                                                                                                                                yaml_2.2.1
## [10] rlang_0.4.7
                                                           withr_2.2.0
                                                                                              dbplyr_1.4.4
                                                                                                               modelr_0.1.8
                         pillar_1.4.6
                                          glue_1.4.1
                                                                             DBI 1.1.0
                                                                                                                                readxl_1.3.
## [19] lifecycle 0.2.0 munsell 0.5.0
                                          gtable 0.3.0
                                                           cellranger 1.1.0 rvest 0.3.5
                                                                                              mvtnorm 1.1-1
                                                                                                               evaluate 0.14
                                                                                                                                knitr 1.29
## [28] highr_0.8
                         broom_0.7.0.9001 Rcpp_1.0.5
                                                           xtable 1.8-4
                                                                            renv 0.12.0
                                                                                              scales 1.1.1
                                                                                                               backports 1.1.8
                                                                                                                                jsonlite_1.
                                                           grid 4.0.2
## [37] hms 0.5.3
                         digest 0.6.25
                                          stringi 1.4.6
                                                                             cli 2.0.2
                                                                                              tools 4.0.2
                                                                                                               crayon 1.3.4
                                                                                                                                 pkgconfig 2
## [46] xml2 1.3.2
                         estimability 1.3 reprex 0.3.0
                                                           lubridate 1.7.9 assertthat 0.2.1 rmarkdown 2.3
                                                                                                               httr 1.4.2
                                                                                                                                rstudioapi_
## [55] compiler 4.0.2
```