Study 1

This file reproduces the preprocessing and analysis steps of Study 1. The data are automatically imported from Github and necessary packages will be downloaded and installed if they are not yet available.

sessionInfo()

[46] foreign_0.8-80

```
## R version 4.0.3 (2020-10-10)
## Platform: x86_64-apple-darwin17.0 (64-bit)
## Running under: macOS Big Sur 10.16
##
## Matrix products: default
           /Library/Frameworks/R.framework/Versions/4.0/Resources/lib/libRblas.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 utils
                                                datasets
                                                          methods
                                                                     base
##
## other attached packages:
   [1] psych_2.0.9
                                emmeans_1.5.3
                                                       brms_2.14.4
    [4] Rcpp_1.0.5
                                BayesFactor 0.9.12-4.2 coda 0.19-4
##
                                lme4_1.1-26
                                                       Matrix_1.2-18
  [7] afex_0.28-0
                                readbulk_1.1.3
                                                       forcats_0.5.0
## [10] jmv_1.2.23
                                dplyr_1.0.2
                                                       purrr_0.3.4
## [13] stringr_1.4.0
                                tidyr 1.1.2
                                                       tibble 3.0.4
## [16] readr 1.4.0
## [19] ggplot2_3.3.2
                                tidyverse_1.3.0
                                                       pacman_0.5.1
##
## loaded via a namespace (and not attached):
     [1] readxl_1.3.1
##
                              backports_1.2.1
                                                    plyr_1.8.6
##
     [4] igraph_1.2.6
                              splines_4.0.3
                                                    crosstalk_1.1.0.1
     [7] TH.data_1.0-10
                              inline_0.3.17
                                                    rstantools_2.1.1
##
    [10] digest_0.6.27
                              htmltools_0.5.0
                                                    rsconnect_0.8.16
##
    [13] lmerTest_3.1-3
                              fansi_0.4.1
                                                    magrittr_2.0.1
##
    [16] openxlsx_4.2.3
                              modelr_0.1.8
                                                    RcppParallel_5.0.2
                              xts_0.12.1
##
    [19] matrixStats_0.57.0
                                                    sandwich_3.0-0
##
    [22] prettyunits 1.1.1
                              colorspace 2.0-0
                                                    rvest 0.3.6
##
   [25] haven_2.3.1
                              xfun_0.19
                                                    callr_3.5.1
##
   [28] crayon 1.3.4
                              jsonlite_1.7.2
                                                    survival 3.2-7
                                                    gtable_0.3.0
##
   [31] zoo_1.8-8
                              glue_1.4.2
    [34] MatrixModels_0.4-1
                              V8_3.4.0
##
                                                    pkgbuild_1.1.0
  [37] car_3.0-10
##
                              rstan_2.21.3
                                                    abind_1.4-5
  [40] scales_1.1.1
##
                              mvtnorm 1.1-1
                                                    DBI 1.1.0
                              xtable_1.8-4
##
   [43] miniUI_0.1.1.1
                                                    tmvnsim_1.0-2
```

StanHeaders_2.21.0-6 stats4_4.0.3

```
[49] DT 0.16
                              htmlwidgets 1.5.3
                                                   httr 1.4.2
## [52] threejs_0.3.3
                              ellipsis_0.3.1
                                                   pkgconfig_2.0.3
                              dbplyr 2.0.0
## [55] loo 2.4.1
                                                   tidyselect 1.1.0
                              reshape2_1.4.4
                                                   later_1.1.0.1
## [58] rlang_0.4.9
## [61] munsell 0.5.0
                              cellranger_1.1.0
                                                   tools_4.0.3
## [64] cli 2.2.0
                              jmvcore 1.2.23
                                                   generics 0.1.0
## [67] broom 0.7.2
                              ggridges 0.5.2
                                                   evaluate 0.14
## [70] fastmap 1.0.1
                              yaml_2.2.1
                                                   processx_3.4.5
## [73] knitr 1.30
                              fs 1.5.0
                                                   zip_2.1.1
## [76] pbapply_1.4-3
                              nlme_3.1-149
                                                   mime_0.9
## [79] projpred_2.0.2
                              xml2_1.3.2
                                                   shinythemes_1.1.2
## [82] compiler_4.0.3
                              bayesplot_1.7.2
                                                   rstudioapi_0.13
## [85] curl_4.3
                              gamm4_0.2-6
                                                   reprex_0.3.0
## [88] statmod_1.4.35
                              stringi_1.5.3
                                                   ps_1.5.0
## [91] Brobdingnag_1.2-6
                              lattice_0.20-41
                                                   nloptr_1.2.2.2
## [94] markdown_1.1
                              shinyjs_2.0.0
                                                   vctrs_0.3.5
## [97] pillar_1.4.7
                              lifecycle_0.2.0
                                                   bridgesampling_1.0-0
## [100] estimability 1.3
                              data.table 1.13.4
                                                   httpuv 1.5.4
## [103] R6_2.5.0
                              promises_1.1.1
                                                   gridExtra_2.3
## [106] rio 0.5.16
                              codetools 0.2-16
                                                   boot 1.3-25
## [109] colourpicker_1.1.0
                             MASS_7.3-53
                                                   gtools_3.8.2
## [112] assertthat_0.2.1
                              withr_2.3.0
                                                   mnormt_2.0.2
## [115] shinystan_2.5.0
                              multcomp_1.4-15
                                                   mgcv_1.8-33
## [118] parallel 4.0.3
                             hms 0.5.3
                                                   grid 4.0.3
                              rmarkdown_2.6
## [121] minga 1.2.4
                                                   carData_3.0-4
## [124] numDeriv 2016.8-1.1 shiny 1.5.0
                                                   lubridate_1.7.9.2
## [127] base64enc_0.1-3
                              dygraphs_1.1.1.6
```

Import data

```
file = 'https://raw.githubusercontent.com/mertensu/thinking-in-ratios/master/data_total_study1.csv'
df = read.csv(file)
```

Preprocessing

Demographics

```
psych::describe(df$age)
             n mean
                       sd median trimmed mad min max range skew kurtosis
## X1
      1 1600 25.15 9.56
                              23
                                   23.12 4.45 19 64
                                                        45 3.3
df %>% distinct(File, .keep_all = T) %>% group_by(gender) %% summarise(
 N = n(),
 Min =
   min(age),
 Max =
   max(age),
 Mean =
   mean(age),
 Sd =
   sd(age)
)
## 'summarise()' ungrouping output (override with '.groups' argument)
## # A tibble: 2 x 6
                         Max Mean
    gender
              N Min
   <chr> <int> <int> <int> <dbl> <dbl>
## 1 m
               4
                    20
                          64 34.8 20.0
## 2 w
              16
                    19
                          30 22.8 3.30
df %>% distinct(File, .keep_all = T) %>% count(student)
##
    student n
## 1
          0 4
## 2
          1 16
df %>% distinct(File, .keep_all = T) %% filter(student == 1) %>% count(psycho)
## psycho n
      0 12
## 1
## 2
         1 4
Analysis
ANOVA I (within 2 (method) x 8(luminance))
df$cd_factor = factor(df$cd, levels = c(1, 1.8, 3.2, 5.7, 17.9, 32.0, 57.2, 100.0))
(fit = aov ez(
dv = 'log_brightness_rating',
```

```
within = c('condition', 'cd_factor'),
id = 'File',
data = df
))
```

frequentist fit

```
## Anova Table (Type 3 tests)
##
## Response: log_brightness_rating
##
                 Effect
                                 df MSE
                                                 F ges p.value
## 1
              condition
                              1, 19 0.26 50.41 *** .090
## 2
              cd_factor 1.36, 25.75 3.15 69.00 *** .692
                                                          <.001
## 3 condition:cd_factor 1.47, 28.02 1.35 13.39 *** .168
                                                         <.001
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '+' 0.1 ' ' 1
## Sphericity correction method: GG
```

```
df_sub = df %>% select(log_brightness_rating, condition, cd_factor, File)
df_sub$condition = factor(df_sub$condition)
df_sub$File = factor(df_sub$File)
bfs = anovaBF(
 log_brightness_rating ~ condition * cd_factor + File,
 whichRandom = 'File',
 whichModels = 'top',
 data = df_sub
# BF cd_stepwise_distance
bf_1 = lmBF(log_brightness_rating ~ condition + File,
            whichRandom = 'File',
            data = df_sub)
bf_2 = lmBF(
 log_brightness_rating ~ cd_factor + condition + File,
 whichRandom = 'File'.
 data = df_sub
(bf_cd_factor = bf_2 / bf_1)
```

bayesian fit

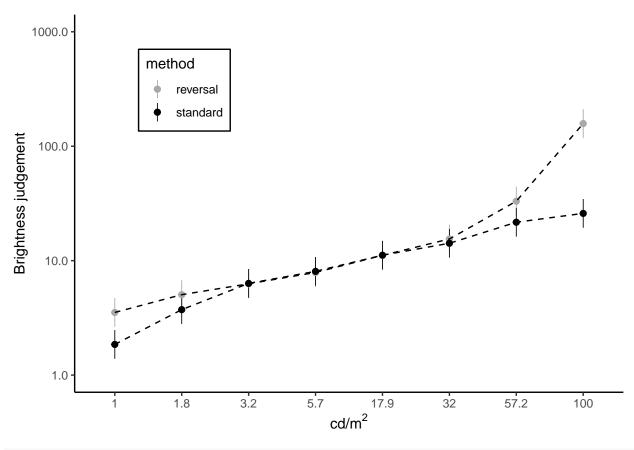
```
## Bayes factor analysis
## -----
## [1] cd_factor + condition + File : 6.903914e+297 ±1.62%
##
## Against denominator:
## log_brightness_rating ~ condition + File
## ---
## Bayes factor type: BFlinearModel, JZS
```

```
print(paste0('logBF ', bf_cd_factor@bayesFactor$bf))
## [1] "logBF 685.799861158154"
# BF condition
bf_1 = lmBF(log_brightness_rating ~ cd_factor + File,
            whichRandom = 'File',
            data = df_sub)
bf_2 = lmBF(
  log_brightness_rating ~ cd_factor + condition + File,
  whichRandom = 'File',
  data = df_sub
(bf_condition = bf_2 / bf_1)
## Bayes factor analysis
## [1] cd_factor + condition + File : 6.516522e+19 ±1.44%
## Against denominator:
    log_brightness_rating ~ cd_factor + File
## ---
## Bayes factor type: BFlinearModel, JZS
print(paste0('logBF ', bf_condition@bayesFactor$bf))
## [1] "logBF 45.6234576226107"
# BF interaction
bf_1 = lmBF(
 log_brightness_rating ~ cd_factor + condition + File,
 whichRandom = 'File',
 data = df sub
)
bf_2 = lmBF(
  log_brightness_rating ~ cd_factor * condition + File,
  whichRandom = 'File',
 data = df_sub
(bf_interaction = bf_2 / bf_1)
## Bayes factor analysis
## -----
## [1] cd_factor * condition + File : 2.965686e+40 ±1.71%
##
## Against denominator:
   log_brightness_rating ~ cd_factor + condition + File
## Bayes factor type: BFlinearModel, JZS
```

```
print(paste0('logBF ', bf_interaction@bayesFactor$bf))
## [1] "logBF 93.1905119234688"
```

Figure 1

```
scaleFUN <- function(x)</pre>
  sprintf("%.1f", x)
grid = data.frame(emmeans(fit, ~ cd_factor + condition))
ggplot(grid, aes(
 x = cd_factor,
 y = \exp(emmean),
 group = condition
)) +
  geom_pointrange(aes(
    ymin = exp(lower.CL),
   ymax = exp(upper.CL),
   color = condition
  ), size =
    0.3) +
  scale_y_continuous(
   trans = 'log2',
    breaks = c(1.0, 10.0, 100.0, 1000.0),
   limits = c(1.0, 1000.0),
   labels = scaleFUN
  ) +
  geom_line(linetype = 'dashed') +
  labs(y = 'Brightness judgement') +
  xlab(expression(paste("cd/", m ^ 2, sep = ""))) +
  scale_color_manual(
   values = c("darkgrey", "black"),
   name = "method",
   labels = c("reversal", "standard")
  scale_x_discrete(labels = substring(grid$cd_factor, 2)) +
  theme_classic() +
  theme(
    legend.position = c(0.2, 0.8),
    legend.background = element_rect(color = "black")
```



```
ggsave(
  paste0("final_plots/study1_figure1.png"),
  dpi = 600,
  height = 4,
  width = 5,
  units = "in"
)
```