贝叶斯因子序列分析在 R 语言中的实现

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1 下载和安装需要的 R 语言程序包	
	导入数据 2.1 数据长宽数据转换

install.packages(c("tidyverse", "BayesFactor", "here")) library(BayesFactor)# 计算 T 检验和方差分析的贝叶斯因子

载入需要的程辑包: coda

载入需要的程辑包: Matrix

Welcome to BayesFactor 0.9.12-4.3. If you have questions, please contact Richard Mor

Type BFManual() to open the manual.

2 导入数据 2

```
## *******
library(tidyverse)
## -- Attaching packages -----
                                    ----- tidyverse 1.3.1 --
## v ggplot2 3.3.6
                            0.3.4
                 v purrr
## v tibble 3.1.7
                  v dplyr 1.0.9
## v tidyr
         1.2.0
                  v stringr 1.4.0
## v readr
                 v forcats 0.5.1
           2.1.2
## -- Conflicts ----- tidyverse_conflicts() --
## x tidyr::expand() masks Matrix::expand()
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                  masks stats::lag()
## x tidyr::pack()
                  masks Matrix::pack()
## x tidyr::unpack() masks Matrix::unpack()
# library(here)
```

2 导入数据

```
df <- read_csv("/Users/zhengyuanrui/Desktop/SBFA/data/df.js.sum_jasp.csv")

## Rows: 20 Columns: 10

## -- Column specification -------

## Delimiter: ","

## chr (1): subj_idx

## dbl (9): Match_Bad, Match_Good, Match_Neutral, Mismatch_Bad, Mismatch_Good, ...

##

## i Use `spec()` to retrieve the full column specification for this data.

## i Specify the column types or set `show_col_types = FALSE` to quiet this message.

## 检验使用数据
```

2.1 数据长宽数据转换

```
df_anova <- df %>%
  select(subj_idx:Mismatch_Good) %>%
  pivot_longer(
    cols = Match_Bad:Mismatch_Good,
    names_to = c("Matchness", "Valence"),
    names_sep = "_", values_to = "rt"
) #anova 使用数据
```

3 查看数据的被试信息

```
subj_num <- unique(df$subj_idx) # 每个被试的编号
n <- length(unique(df$subj_idx)) # 一共有 20 个被试
```

#(配对样本)T 检验的 R 语言实现 ## good_match 条件与 bad_match 条件的对比先建立一个空的列表,用来储存后续的贝叶斯因子。列表长度为目前数据的样本量

bf_output <- rep(NA, length(subj_num)) ### 先建立一个列表

```
for (i in seq_along(subj_num)) {#i 遍历 subj_num
    if (i == 1) {
        next
        # 由于一个被试不能正确计算贝叶斯因子,所以当 i 等于 1 时,跳过
    }
        # 将 df 数据框中的 subj_idx 列转换为字符串型
    df$subj_idx <- as.character(df$subj_idx)
    # 提取出遍历到的被试编号
    id <- unique(df$subj_idx)[1:i]
    # 从愿数据中筛选被试
    df.selected <- df %>% filter(subj_idx %in% id)
    # 转换为因子型
```

```
df.selected$subj_idx <- as.factor(df.selected$subj_idx)</pre>
  bayesfactors <- ttestBF(</pre>
   x = df.selected$Match_Bad, y = df.selected$Match_Good,
   paired = TRUE
  )# 计算贝叶斯因子
  bf_output[i] <- bayesfactors[1]</pre>
}
bf_output
   [1]
                 NA
                       1.079341
                                   2.095059
                                               3.339921
                                                           6.652698
                                                                      19.097998
## [7]
         31.855000 49.752936 130.961355
                                              58.188628
                                                          93.428773 239.350270
## [13] 127.216734 250.586118 571.531998 1522.706898 2032.527877 2787.948423
```

4 (重复测量) 方差分析的 R 语言实现

4.1 数据的基本信息

[19] 5664.722345 3113.214090

```
subj_num <- unique(df_anova$subj_idx) # 每个被试的编号
n <- length(unique(df_anova$subj_idx)) # 一共有 20 个被试

生成三个列表用来储存两个主效应和交互项

BFs_match <- rep(NA, length(subj_num))

BFs_valence <- rep(NA, length(subj_num))

BFs_int <- rep(NA, length(subj_num))

for (i in seq_along(subj_num)) {
   if (i == 1) {
      next
   }
```

```
df_anova$subj_idx <- as.character(df_anova$subj_idx)</pre>
  id <- unique(df_anova$subj_idx)[1:i]</pre>
  df.selected <- df_anova %>% filter(subj_idx %in% id)
  df.selected$subj_idx <- as.factor(df.selected$subj_idx)</pre>
  df.selected$Matchness <- as.factor(df.selected$Matchness)</pre>
  df.selected$Valence <- as.factor(df.selected$Valence)</pre>
  bayesfactors <- bf <- anovaBF(rt ~ Valence*Matchness + subj_idx,</pre>
    data = data.frame(df.selected),
    whichRandom = "subj idx"
  BFs_match[i] <- bayesfactors[1]</pre>
  BFs_valence[i] <- bayesfactors[2]</pre>
  BFs_int[i] <- bayesfactors[4] / bayesfactors[3]</pre>
}
BFs_match
## [1]
                 NA
                      0.9093042
                                  0.8655600
                                               0.7014032
                                                           1.1991076
                                                                        1.1149060
## [7]
          2.0060562
                      2.6873412
                                  4.2852944 11.2008425
                                                           3.8648072
                                                                        9.7589807
## [13] 13.9544256 18.0472772 27.7213725 33.4998854 40.2743944 95.1521970
## [19] 131.4854521 400.2762456
BFs_valence
## [1]
                 NA
                      0.5367660
                                  0.7057221
                                               2.3770512
                                                           2.0285497
                                                                       5.0915627
  [7]
          6.5876690
                      9.5020914 20.3369769 12.7270155 48.8438452 56.7354946
##
## [13]
        26.1812938 29.6711418 38.2317769 82.3386615 147.4454763 118.9068572
## [19] 247.2357395 122.3666839
BFs_int
                  NA 1.543211e+00 3.694320e+00 5.074814e+00 6.928469e+00
##
  [1]
## [6] 4.006296e+01 7.350215e+01 1.660965e+02 5.892989e+02 2.479563e+02
## [11] 5.720678e+02 1.912574e+03 1.835334e+03 5.449767e+03 1.491577e+04
## [16] 7.541347e+04 7.334727e+04 1.115412e+05 2.082175e+05 6.840184e+04
```