

贝叶斯因子序列分析在 R 语言中的实现

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```
rm(list = ls())
```

1 下载和安装需要的 R 语言程序包

```
# install.packages(c("tidyverse", "BayesFactor", "here"))
library(BayesFactor) # 计算 T 检验和方差分析的贝叶斯因子
library(tidyverse)
library(showtext) # 解决中文字体无法显示问题
library(latex2exp) # latex 语法
font_add("song",
```

```

"/System/Library/Fonts/Supplemental/Songti.ttc")
## 从系统增加宋体字
showtext_auto()
library(here)
here()

## [1] "/Users/zhengyuanrui/SBFA_Tutorial"

options(scipen = 9) # 将科学计数法改为在万后 9 位
set.seed(1234)

```

1.1 BayesFacotr 包版本

```

packageVersion("BayesFactor")

## [1] '0.9.12.4.3'

```

2 导入数据

```
df <- readr::read_csv(here("2_Data", "df.sum_jasp.csv"))
```

2.1 数据长宽数据转换

2.1.1 因变量为 RT 的数据整理

```

# 分析因变量为 RT 的使用数据
df.RT <- df %>%
  # 选择被试信息以及 RT_ 开头的列
  dplyr::select(subj_idx, starts_with("RT_")) %>%
  # RT_Bad_Match 到 RT_Good_Nonmatch 列转换为长数据, 列名为 condition, 值名为 rt
  tidyr::pivot_longer(
    cols = RT_Bad_Match:RT_Neutral_Nonmatch,
    names_to = "condition",
    values_to = "rt"
  ) %>%
  # 将 condition 列拆分为三列, DV_Name 为因变量名称
  # Valence 是道德信息, Matchness 是匹配信息
  tidyr::separate(col = condition,
    into = c("DV_Name", "Valence", "Matchness"),

```

```

      sep = "_") %>%
# 类型为 character 的转换为因子类型，便于后续分析
dplyr::mutate_if(is.character, as.factor)

head(df.RT)

```

```

## # A tibble: 6 x 5
##   subj_idx DV_Name Valence Matchness   rt
##   <fct>    <fct>   <fct>   <fct>   <dbl>
## 1 v010001 RT      Bad     Match    775.
## 2 v010001 RT      Bad     Nonmatch  793.
## 3 v010001 RT      Good    Match    716.
## 4 v010001 RT      Good    Nonmatch  817.
## 5 v010001 RT      Neutral Match    747.
## 6 v010001 RT      Neutral Nonmatch  786.

```

2.1.2 因变量为 ACC 的数据整理

```

df.ACC <- df %>%
  dplyr::select(subj_idx, starts_with("ACC_")) %>%
  tidyr::pivot_longer(
    cols = ACC_Bad_Match:ACC_Neutral_Nonmatch,
    names_to = "condition",
    values_to = "ACC"
  ) %>%
  tidyr::separate(col = condition,
    into = c("DV_Name", "Valence", "Matchness"),
    sep = "_") %>%
  dplyr::mutate_if(is.character, as.factor)

head(df.ACC)

```

```

## # A tibble: 6 x 5
##   subj_idx DV_Name Valence Matchness  ACC
##   <fct>    <fct>   <fct>   <fct>   <dbl>
## 1 v010001 ACC      Bad     Match    0.723
## 2 v010001 ACC      Bad     Nonmatch  0.730
## 3 v010001 ACC      Good    Match    0.896
## 4 v010001 ACC      Good    Nonmatch  0.8
## 5 v010001 ACC      Neutral Match    0.838
## 6 v010001 ACC      Neutral Nonmatch  0.794

```

2.1.3 因变量为 dPrime 的数据整理

```
df.dPrime <- df %>%
  dplyr::select(subj_idx, starts_with("dPrime_")) %>%
  tidyr::pivot_longer(
    cols = dPrime_Bad:dPrime_Neutral,
    names_to = "condition",
    values_to = "dPrime"
  ) %>%
  tidyr::separate(col = condition,
    into = c("DV_Name", "Valence"),
    sep = "_" ) %>%
  dplyr::mutate_if(is.character, as.factor)

head(df.dPrime)
```

```
## # A tibble: 6 x 4
##   subj_idx DV_Name Valence dPrime
##   <fct>    <fct>   <fct>   <dbl>
## 1 v010001 dPrime   Bad      1.21
## 2 v010001 dPrime   Good     2.10
## 3 v010001 dPrime   Neutral  1.81
## 4 v010003 dPrime   Bad      1.02
## 5 v010003 dPrime   Good     2.05
## 6 v010003 dPrime   Neutral  1.04
```

3 正确的 BF 计算

```
bayesfactors <- BayesFactor::generalTestBF(
  rt ~ Valence*Matchness*subj_idx - subj_idx:Valence:Matchness,
  data = data.frame(df.RT),
  whichRandom = "subj_idx",
  neverExclude = "subj_idx",
  whichModels = "all")
```

```
bayesfactors
```

```
## Bayes factor analysis
## -----
## [1] Valence + subj_idx + Valence:subj_idx + Matchness:subj_idx
```

```
## [2] Matchness + subj_idx + Valence:subj_idx + Matchness:subj_idx : 2
## [3] Valence:Matchness + subj_idx + Valence:subj_idx + Matchness:subj_idx : 5
## [4] Valence + Matchness + subj_idx + Valence:subj_idx + Matchness:subj_idx : 5
## [5] Valence + Valence:Matchness + subj_idx + Valence:subj_idx + Matchness:subj_idx : 3
## [6] Matchness + Valence:Matchness + subj_idx + Valence:subj_idx + Matchness:subj_idx : 1
## [7] Valence + Matchness + Valence:Matchness + subj_idx + Valence:subj_idx + Matchness:subj_idx : 1
## [8] subj_idx + Valence:subj_idx + Matchness:subj_idx : 8
##
## Against denominator:
##   Intercept only
## ---
## Bayes factor type: BFlinearModel, JZS

# 感兴趣的效应都要先除零模型（仅包括随机效应的模型）
null <- bayesfactors[8]

# 全模型
full <- bayesfactors[7]
BF_full.n <- full/null # 全模型与 null 对比
BF_excinx.n <- bayesfactors[4]/null
BF_inx <- BF_full.n/BF_excinx.n
BF_inx

## Bayes factor analysis
## -----
## [1] Valence + Matchness + Valence:Matchness + subj_idx + Valence:subj_idx + Matchness:subj_idx : 2
##
## Against denominator:
##   rt ~ Valence + Matchness + subj_idx + Valence:subj_idx + Matchness:subj_idx
## ---
## Bayes factor type: BFlinearModel, JZS

BF_m.n <- bayesfactors[2]/null
BF_excinx.n/BF_m.n

## Bayes factor analysis
## -----
## [1] Valence + Matchness + subj_idx + Valence:subj_idx + Matchness:subj_idx : 2661.48 ±2.67%
##
## Against denominator:
##   rt ~ Matchness + subj_idx + Valence:subj_idx + Matchness:subj_idx
## ---
## Bayes factor type: BFlinearModel, JZS
```

```

BF_v.n <- bayesfactors[1]/null
BF_excinx.n/BF_v.n

## Bayes factor analysis
## -----
## [1] Valence + Matchness + subj_idx + Valence:subj_idx + Matchness:subj_idx : 515.3729 ±2.59%
##
## Against denominator:
##   rt ~ Valence + subj_idx + Valence:subj_idx + Matchness:subj_idx
## ---
## Bayes factor type: BFlinearModel, JZS

ttestBF(df$RT_Good_Match,
        df$RT_Neutral_Match,
        paired = TRUE,
        nullInterval = c(Inf, 0))[2]

## t is large; approximation invoked.
## t is large; approximation invoked.

## Bayes factor analysis
## -----
## [1] Alt., r=0.707 !(0<d<Inf) : 14411.55 ±NA%
##
## Against denominator:
##   Null, mu = 0
## ---
## Bayes factor type: BFoneSample, JZS

```

4 查看数据的被试信息

```

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

## [1] 20

```

5 配对样本 T 检验的 R 语言实现

5.1 假设: good_match 条件优于 Neutral_match 条件

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(1, 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$subj_idx)[1:i]
  # 从原数据中筛选被试
  df.selected <- df %>% filter(subj_idx %in% id)
  # 转换为因子型
  df.selected$subj_idx <- as.factor(df.selected$subj_idx)
  # 计算贝叶斯因子
  bayesfactors <- ttestBF(df.selected$RT_Good_Match,
                          df.selected$RT_Neutral_Match,
                          paired = TRUE,
                          nullInterval = c(0, Inf))
  bf_output[i] <- bayesfactors[2]
}
```

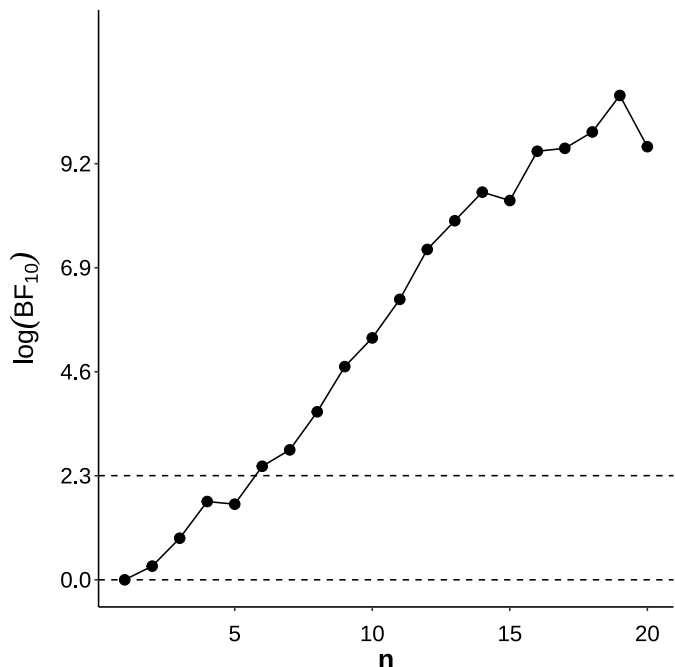
5.2 结果

```
tibble(bf_output) %>%
  dplyr::mutate(n = 1:nrow(.)) %>%
  dplyr::rename("Bayes Factor" = "bf_output") %>%
  dplyr::mutate(logBF = log(`Bayes Factor`)) %>%
  ggplot(aes(x = n, y = logBF)) +
  geom_point(size = 3) +
  geom_line() +
  geom_hline(aes(yintercept = log(10)), linetype = "dashed") +
  geom_hline(aes(yintercept = log(1)), linetype = "dashed") +
  scale_y_continuous(
    limits = c(0, 12),
    breaks = c(round(log(1), 1),
```

```

round(log(10), 1),
round(log(100), 1),
round(log(1000), 1),
round(log(10000), 1))) +
labs(y = TeX("$\\log(BF_{10})$")) +
theme(
  panel.background = element_blank(),
  plot.margin = unit(c(1, 1, 1, 1), "cm"),
  plot.background = element_rect(fill = "white", color = NA),
  plot.title = element_text(size = 22, face = "bold",
    hjust = 0.5,
    margin = margin(b = 15)),
  axis.line = element_line(color = "black", size = .5),
  axis.title = element_text(size = 18, color = "black",
    face = "bold"),
  axis.text = element_text(size = 15, color = "black"),
  axis.text.x = element_text(margin = margin(t = 10)),
  axis.title.y = element_text(margin = margin(r = 10)),
  axis.ticks = element_line(size = .5),
  panel.grid = element_blank(),
  legend.position = c(0.20, 0.8),
  legend.background = element_rect(color = "black"),
  legend.text = element_text(size = 15),
  legend.margin = margin(t = 5, l = 5, r = 5, b = 5),
  legend.key = element_rect(color = NA, fill = NA))

```



6 （重复测量）方差分析的 R 语言实现

6.1 数据的基本信息

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

## [1] 20
```

6.2 检验交互项

生成三个向量用来储存两个主效应和交互项

```
BFs_match <- rep(1, length(subj_num))

BFs_valence <- rep(1, length(subj_num))

BFs_int <- rep(1, length(subj_num))

for (i in seq_along(subj_num)) {
  if (i == 1) {
    next
  }
  df.RT$subj_idx <- as.character(df.RT$subj_idx)
  id <- unique(df.RT$subj_idx)[1:i]
  df.selected <- df.RT %>% dplyr::filter(subj_idx %in% id)
  df.selected$subj_idx <- as.factor(df.selected$subj_idx)
  df.selected$Matchness <- as.factor(df.selected$Matchness)
  df.selected$Valence <- as.factor(df.selected$Valence)
  bayesfactors <- BayesFactor::generalTestBF(
    rt ~ Valence*Matchness*subj_idx - subj_idx:Valence:Matchness,
    data = data.frame(df.selected),
    whichRandom = "subj_idx",
    neverExclude = "subj_idx",
    whichModels = "all", progress = FALSE)

  null <- bayesfactors[8]
  full <- bayesfactors[7] # 全模型
  BF_full.n <- full/null # 全模型与 null 对比
  BF_excinx.n <- bayesfactors[4]/null
  BF_m.n <- bayesfactors[2]/null
```

```
BF_v.n <- bayesfactors[1]/null
BFs_match[i] <- BF_excinx.n/BF_v.n# 计算 Matchness 主效应的 BF
BFs_valence[i] <- BF_excinx.n/BF_m.n# 计算 Valence 的 BF
BFs_int[i] <- BF_full.n/BF_excinx.n# 计算交互项的 BF
}

aov_output <- tibble::tibble(BFs_int, BFs_valence, BFs_match)# 整合为数据框
head(aov_output)# 查看数据
```

```
## # A tibble: 6 x 3
##   BFs_int BFs_valence BFs_match
##   <dbl>      <dbl>      <dbl>
## 1      1          1          1
## 2    3.82      0.558      0.700
## 3   24.6      0.645      0.708
## 4  143.        1.35      0.642
## 5   84.1      1.57      0.946
## 6 1801.        2.67      0.953
```

6.3 结果

6.3.1 长宽数据、log 变换

```
dat_plot <- aov_output %>% dplyr::mutate(n = 1:nrow(.)) %>%
  tidyr::pivot_longer(BFs_int:BFs_match, names_to = "Effect",
    values_to = "Bayes Factor") %>%
  dplyr::mutate(`logBF` = log(`Bayes Factor`)) %>%
  dplyr::mutate(dplyr::across(where(is.double),
    ~round(.x, digits = 2)))
head(dat_plot, 10)
```

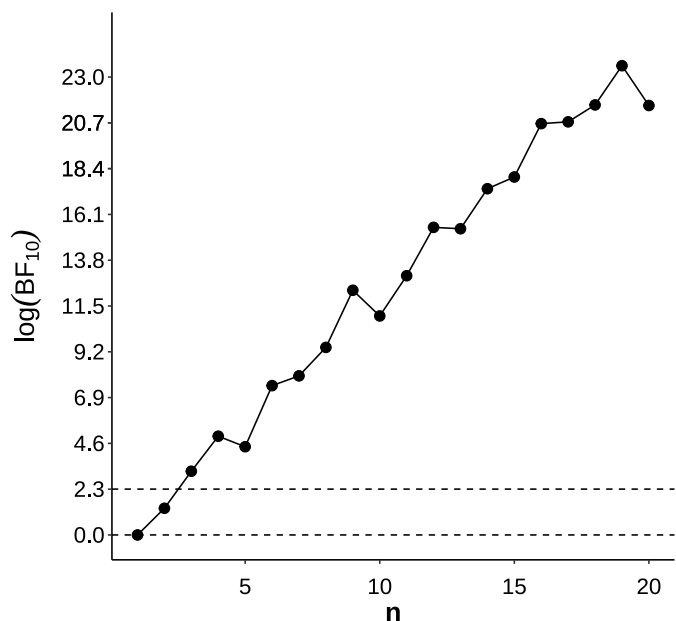
```
## # A tibble: 10 x 4
##       n Effect      `Bayes Factor` logBF
##   <int> <chr>          <dbl> <dbl>
## 1     1 1 BFs_int          1      0
## 2     1 1 BFs_valence      1      0
## 3     1 1 BFs_match        1      0
## 4     2 2 BFs_int        3.82  1.34
## 5     2 2 BFs_valence     0.56 -0.58
## 6     2 2 BFs_match       0.7  -0.36
## 7     3 3 BFs_int       24.6   3.2
```

```
## 8      3 BFs_valence      0.65 -0.44
## 9      3 BFs_match       0.71 -0.35
## 10     4 BFs_int         143.    4.96

dat_plot %>% dplyr::filter(Effect == "BFs_int") %>%
  ggplot(aes(x = n, y = logBF)) +
  geom_point(size = 3) +
  geom_line() +
  geom_hline(aes(yintercept = log(1)), linetype = "dashed")+
  geom_hline(aes(yintercept = log(10)), linetype = "dashed")+
  labs(y = TeX("$\\log(BF_{10})$")) +
  ggtitle("交互作用的贝叶斯因子数值变化趋势") +
  scale_y_continuous(
    limits = c(0, 25),
    breaks = c(0, round(log(10), 1),
      round(log(100), 1),
      round(log(1000), 1),
      round(log(10000), 1),
      round(log(100000), 1),
      round(log(1000000), 1),
      round(log(10000000), 1),
      round(log(100000000), 1),
      round(log(1000000000), 1),
      round(log(10000000000), 1),
      round(log(100000000000), 1),
      round(log(1000000000000), 1)
    )
  ) +
  theme(
    panel.background = element_blank(),
    plot.margin = unit(c(1, 1, 1, 1), "cm"),
    plot.background = element_rect(fill = "white", color = NA),
    plot.title = element_text(size = 22,
      family = "song",
      face = "bold",
      hjust = 0.5,
      margin = margin(b = 15)),
    axis.line = element_line(color = "black", size = .5),
    axis.title = element_text(size = 18, color = "black",
      face = "bold"),
    axis.text = element_text(size = 15, color = "black"),
    axis.text.x = element_text(margin = margin(t = 10)),
```

```
axis.title.y = element_text(margin = margin(r = 10)),  
axis.ticks = element_line(size = .5),  
panel.grid = element_blank(),  
legend.position = c(0.20, 0.8),  
legend.background = element_rect(color = "black"),  
legend.text = element_text(size = 15),  
legend.margin = margin(t = 5, l = 5, r = 5, b = 5),  
legend.key = element_rect(color = NA, fill = NA))
```

交互作用的贝叶斯因子数值变化趋势



```
# ggsave("RT_inx.png", width = 10, height = 7, dpi = 300)
```