补充材料

潘晚坷 温秀娟 金海洋

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1 数据模拟

我们借助一个假想的心理学实验展示如何模拟虚拟数据。在模拟实验中,40 名抑郁症患者和40 名健康对照组被试观看30 张积极和30 张中性图片,期间我们采集了他们的脑电数据。因变量是晚期正电位(late positive potentials, LPP)的波幅。简单来说,这是一个2 (组别 group: 抑郁症组 depression、对照组 control) × 2 (图片类型type: 积极 positive、中性 neutral)的混合实验设计,其中组别为被试间因素,图片类型为被试内因素。该假想实验的数据是使用 faux 工具包生成 (DeBruine 2023),下面是模拟这个实验所预设的参数。

```
subj_n <- 80  # 总被试量: 抑郁患者 30 人, 健康对照组被试 30 人 trial_n <- 30  # 每张图片呈现的次数
```

固定效应

1 数据模拟 2

```
b0 <- 0.5
          # 截距 (所有条件的均值)
           #图片类型的固定效应 (主效应)
b1 <- 6.5
b2 <- 0.1 # 组别的固定效应 (主效应)
          # 图片类型与组别的交互作用
b3 <- 0.1
# 随机效应
u0s <- 2 # 被试的随机截距
u1s <- 2 # 被试的随机斜率 (图片类型)
#误差项
sigma <- 2
# 生成假定实验的条件的数据矩阵
df_simu <- add_random(subj = subj_n) %>%
 #添加被试的组别信息(被试间)
 add_between("subj", group = c("depression", "control")) %>%
 #添加图片类型的信息(被试内)
 add_within("subj", type = c("netural", "positive")) %>%
 # 每种图片呈现 30 次
 add_random(trial = trial_n) %>%
 # 图片类型的编码: 中性 =-0.5; 正性 =0.5
 add_contrast("type", "anova", colnames = "type_code") %>%
 # 被试组别的编码: 抑郁症组 =-0.5; 控制组 =0.5
 add_contrast("group", "anova",colnames = "group_code") %>%
 #添加基于被试的随机截距和斜率 (图片类型)
 add_ranef("subj", u0s = u0s, u1s = u1s, .cors=0.5) %>%
 #添加观察值的误差项
 add_ranef(sigma = sigma) %>%
 # 最后根据设置的固定效应和随机效应参数值, 生成因变量。
 mutate(LPP = (b0+u0s) +
                           # 截距
         (b1+u1s) * type_code + # 图片材料的斜率
        b2 * group_code + # 组别的斜率
        b3 * type_code * group_code + # 交互作用
                       # 误差项
         sigma)
```

2 方差分析 3

```
df_simu <- df_simu %>%
  select(subj, group, type, LPP) # 去除冗余的信息
# 保存生成的数据
save(df_simu, file = "simulated_data.rdata")
# 查看生成的数据
head(df_simu,10)
## # A tibble: 10 x 4
                                 LPP
##
     subj
                       type
            group
##
     <chr> <fct>
                       <fct>
                               <dbl>
## 1 subj01 depression netural -1.64
   2 subj01 depression netural -4.00
##
   3 subj01 depression netural -1.53
## 4 subj01 depression netural -3.22
## 5 subj01 depression netural -4.34
## 6 subj01 depression netural 1.49
## 7 subj01 depression netural -3.46
## 8 subj01 depression netural 2.93
## 9 subj01 depression netural -3.27
## 10 subj01 depression netural -1.44
```

2 方差分析

我们使用 bruceR 包进行混合实验设计方差分析 (Bao 2023)。以下为示例代码与结果。其中,图片类型的主效应显著 $(F(1,78)=983.78,p<0.01,\eta_p^2=.93)$,而组别的主效应、组别与图片类型的交互作用不显著 (p>0.1)。

```
df_simu = df_simu |> mutate(
组别 = factor(group, labels = c(" 抑郁组"," 控制组")),
图片类型 = factor(type, labels = c(" 中性图片"," 积极图片"))
```

2 方差分析 4

```
# Two-way mixed ANOVA test
df_simu |> bruceR::MANOVA(
 subID = "subj",
 between = "组别",
 within = "图片类型",
 dv = "LPP",
 digits = 2
 # file = " 重复测量方差分析结果.doc"
)
##
##
      * Data are aggregated to mean (across items/trials)
##
      if there are >=2 observations per subject and cell.
##
      You may use Linear Mixed Model to analyze the data,
      e.g., with subjects and items as level-2 clusters.
##
## ===== ANOVA (Mixed Design) ======
##
## Descriptives:
##
##
   "组别" "图片类型" Mean
                           S.D. n
##
   抑郁组
            中性图片 -2.71 (2.12) 40
##
   抑郁组 积极图片 3.78 (2.56) 40
##
   控制组
          中性图片 -2.43 (1.97) 40
          积极图片 4.07 (2.62) 40
##
   控制组
##
## Total sample size: N = 80
##
## ANOVA Table:
## Dependent variable(s):
                            LPP
## Between-subjects factor(s): 组别
## Within-subjects factor(s): 图片类型
## Covariate(s):
```

```
##
##
                          MS MSE df1 df2
                                             F
                                                      p <sup>2</sup>p [90% CI of <sup>2</sup>p] <sup>2</sup>G
##
## 组别
                        3.19 9.18
                                    1 78
                                             0.35
                                                   .557
                                                                  .00 [.00, .06] .00
## 图片类型
                    1689.20 1.72
                                    1
                                       78 983.78 <.001 ***
                                                                  .93 [.90, .94] .67
## 组别 * 图片类型
                        0.00 1.72
                                    1
                                       78
                                             0.00
                                                   .969
                                                                  .00 [.00, .00] .00
##
## MSE = mean square error (the residual variance of the linear model)
## ^{2}p = partial eta-squared = SS / (SS + SSE) = F * df1 / (F * df1 + df2)
## ^{2}p = partial omega-squared = (F - 1) * df1 / (F * df1 + df2 + 1)
## <sup>2</sup>G = generalized eta-squared (see Olejnik & Algina, 2003)
## Cohen' s f^2 = {}^2p / (1 - {}^2p)
##
## Levene's Test for Homogeneity of Variance:
##
                 Levene's F df1 df2
##
##
## DV: 中性图片
                       0.772
## DV: 积极图片
                       0.027
                               1 78
                                     .869
##
##
## Mauchly's Test of Sphericity:
## The repeated measures have only two levels. The assumption of sphericity is always met.
```

3 不收敛 MCMC 链演示

我们使用 bayesplot 和 posterior 包模拟和绘制 MCMC 链 (Gabry et al. 2019; Vehtari et al. 2021)。图 1 为 MCMC 链不收敛的示例。其中,四条链存在明显的分离,并且第四条链并没有达到稳定分布。 $\hat{R}=2.73$ 的结果大于 1.1,说明该 MCMC 链的收敛结果 很差。

```
# 模拟生成 4 条不收敛的 MCMC 链, 每条链包含 4000 个样本
n_chains <- 4
chain_length <- 4000
```

```
# 生成三种链,一种收敛的链 good_chains,两种不收敛的链 bad_chains0 和 bad_chains1
good_chains <- rbeta(n = chain_length*n_chains, shape1 = 2, shape2 = 5)
good_chains <- matrix(good_chains, nrow = n_chains)
bad_chains <- matrix(
    rnorm(chain_length*n_chains, mean = sort(good_chains), sd = 0.05),
    nrow = n_chains)

chains <- array(0, dim = c(chain_length, n_chains,1))
chains[,,1] = bad_chains
dimnames(chains) <- list(
    Iteration = NULL,
    Chain = paste0("chain:",1:n_chains),
    Parameter = c("bad_chains")
)

# 绘制轨迹图
mcmc_trace(chains)
```

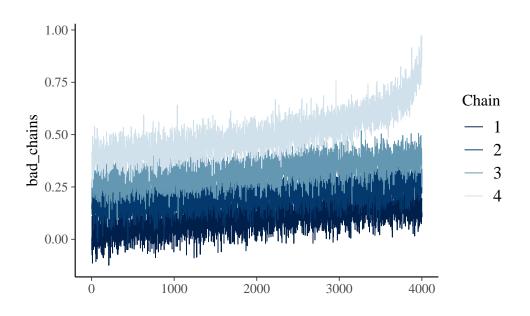


图 1: 不收敛 MCMC 链演示图

计算 rhat

rhat(extract_variable_matrix(chains, "bad_chains")) # 2.73128

[1] 2.728107

参考文献

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