Analyzing RCTs: A Cookbook

Inge Christoffer Olsen 2019-08-09

Contents

1	\mathbf{Pre}	face	5	
	1.1	Structure of the book	5	
	1.2	Acknowledgements	5	
2				
	2.1	Trial Flow	7	
	2.2	Simulated dataset	7	
3	Continuous endpoints 9			
	3.1		9	
	3.2		9	
4	Dichotomous endpoints 11			
	4.1	Single follow-up	11	
5	Applications 15			
	5.1	Example one	15	
		Example two	15	
G	Fin	ol Words	17	

4 CONTENTS

Preface

Placeholder

- 1.1 Structure of the book
- 1.2 Acknowledgements

Introduction

Placeholder

- 2.1 Trial Flow
- 2.2 Simulated dataset

Continuous endpoints

Placeholder

- 3.1 Single follow-up
- 3.1.1 Stata code
- 3.1.2 R code
- 3.1.3 Reporting
- 3.2 Repeated follow-up
- 3.2.1 Simple model
- 3.2.2 Model with treatment-time interaction
- 3.2.3 Model with treatment-time interaction and baseline information

Dichotomous endpoints

4.1 Single follow-up

For a single follow-up assessment of a dichotomous endpoint, the main method I use is a standard logistic regression. Then we can adjust for stratification factors in the randomisation in addition to other pre-specified covariates, both categorical and continuous. In the simulated example, we define that the primary outcome is the dichotomous categorical outcome at time 3. Note that usually the baseline status of all patients are negative for the outcome, so adjusting for baseline is not necessary.

4.1.1 Stata code

```
use "stata/rct", clear
tabulate catout trt
logistic catout i.trt i.site covar if time==3, coef
## . use "stata/rct", cle(all strata combined)
## . tabulate catout trt
##
## Categorica |
                      Treatment
## 1 outcome | Placebo Active |

        Negative |
        95
        131 |

        Positive |
        105
        61 |

##
                                              226
                                              166
## -----+----
       Total | 200 192 |
                                              392
##
```

```
##
## . logistic catout i.trt i.site covar if time==3, coef
## Logistic regression
                                           Number of obs =
                                                                   98
                                           LR chi2(5)
##
                                                                48.59
##
                                           Prob > chi2
                                                                0.0000
## Log likelihood = -36.862204
                                           Pseudo R2
                                                                0.3973
##
## -----
##
                  Coef. Std. Err. z P>|z|
                                                   [95% Conf. Interval]
       catout
##
         trt |
      Active | -2.890301 .7850252 -3.68 0.000
##
                                                   -4.428922 -1.351679
##
##
         site |

    2
    |
    .7783404
    .8580245
    0.91
    0.364

    3
    |
    1.423791
    .7786531
    1.83
    0.067

    4
    |
    .0253234
    .8082887
    0.03
    0.975

##
                                                   -.9033566
                                                              2.460037
##
                                                   -.1023412
                                                              2.949923
##
                                                   -1.558893
                                                              1.60954
##
        covar | 1.001078 .2329461
                                     4.30 0.000
##
                                                    .5445124
                                                             1.457644
##
        _cons | -2.463577 .8925892 -2.76 0.006
                                                   -4.21302 -.7141344
## -----
##
## .
```

Note that the use the coef option to get the log odds ratio estimates.

4.1.2 R code

```
rct <- read_dta("stata/rct.dta") %>%
 modify_at(c("trt","catout"), haven::as_factor, levels = "labels") %>%
 modify_at(c("site","time"), haven::as_factor)
rct %>%
 filter(time==3) %>%
  glm(catout ~ trt + site + covar , data=., family = binomial) %>%
  summary
##
## Call:
## glm(formula = catout ~ trt + site + covar, family = binomial,
      data = .)
##
## Deviance Residuals:
## Min 1Q Median
                                 3Q
                                        Max
## -1.9659 -0.5335 0.1943 0.5095 2.8873
```

```
##
## Coefficients:
          Estimate Std. Error z value Pr(>|z|)
## trtActive -2.89030 0.78502 -3.682 0.000232 ***
## site2
          0.77834 0.85802 0.907 0.364337
## site3
           1.42379 0.77865 1.829 0.067470 .
## site4
           ## covar 1.00108 0.23295 4.297 1.73e-05 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
     Null deviance: 122.318 on 97 degrees of freedom
## Residual deviance: 73.724 on 92 degrees of freedom
## AIC: 85.724
## Number of Fisher Scoring iterations: 6
```

Applications

Some significant applications are demonstrated in this chapter.

- 5.1 Example one
- 5.2 Example two

Final Words

We have finished a nice book.