Mediation_analysis_question

2023-04-15

Continous outcome and continous mediator

Call:

Results for continuous outcome and continuous mediator look all right.

```
n = 1000
#C is the covariate
C = rnorm(n)
#X is the treatment variable
X = rnorm(n, mean = 0, sd = 1)
#M is the mediator
M = 0.3*X + 0.5*C + rnorm(n)
#Y is the outcome
Y = M*0.2+0.1*X + 0.3*C
data = data.frame(Y, X, M, C)
#direct effect is 0.1
#indirect effect is 0.06
#total effect is 0.16
out_model = lm(Y ~ X + M + C, data = data)
med_model = lm(M~X + C, data = data)
total_model = lm(Y ~ X + C, data = data)
summary(out_model)
##
## Call:
## lm(formula = Y \sim X + M + C, data = data)
## Residuals:
                      1Q
                             Median
                                            3Q
## -1.393e-14 -6.700e-18 2.730e-17 5.670e-17 3.154e-16
## Coefficients:
##
                Estimate Std. Error
                                        t value Pr(>|t|)
## (Intercept) -2.940e-17 1.570e-17 -1.872e+00 0.0614 .
## X
               1.000e-01 1.629e-17 6.140e+15
                                                  <2e-16 ***
## M
                2.000e-01 1.533e-17 1.305e+16
                                                  <2e-16 ***
## C
               3.000e-01 1.757e-17 1.707e+16
                                                  <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 4.966e-16 on 996 degrees of freedom
## Multiple R-squared: 1, Adjusted R-squared:
## F-statistic: 3.245e+32 on 3 and 996 DF, p-value: < 2.2e-16
summary(med_model)
##
```

```
## lm(formula = M ~ X + C, data = data)
##
## Residuals:
##
               1Q Median
      Min
                              3Q
                                     Max
## -3.2414 -0.7041 0.0172 0.7173 3.5097
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 0.001712
                        0.032450
                                   0.053
                                            0.958
## X
              0.317472
                         0.032117
                                   9.885
                                           <2e-16 ***
## C
              0.537496
                        0.032067 16.761
                                           <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 1.026 on 997 degrees of freedom
## Multiple R-squared: 0.2762, Adjusted R-squared: 0.2748
## F-statistic: 190.2 on 2 and 997 DF, p-value: < 2.2e-16
summary(total_model)
##
## Call:
## lm(formula = Y ~ X + C, data = data)
##
## Residuals:
##
       Min
                 1Q
                    Median
                                  3Q
## -0.64828 -0.14082 0.00344 0.14345 0.70194
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
##
                                   0.053
## (Intercept) 0.0003425 0.0064900
                                            0.958
## X
              0.1634944 0.0064234 25.453
                                            <2e-16 ***
## C
              ## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.2052 on 997 degrees of freedom
## Multiple R-squared: 0.8251, Adjusted R-squared: 0.8247
## F-statistic: 2351 on 2 and 997 DF, p-value: < 2.2e-16
library(mediation)
## Warning: package 'mediation' was built under R version 4.0.2
## Loading required package: MASS
## Warning: package 'MASS' was built under R version 4.0.2
## Loading required package: Matrix
## Warning: package 'Matrix' was built under R version 4.0.2
## Loading required package: mvtnorm
## Loading required package: sandwich
## Warning: package 'sandwich' was built under R version 4.0.2
## mediation: Causal Mediation Analysis
```

```
## Version: 4.5.0
fit_model = mediate(med_model, out_model, treat="X", mediator= "M", boot=T, boot.ci.type = "bca")
## Running nonparametric bootstrap
#result from mediation package looks right
summary(fit model)
## Causal Mediation Analysis
## Nonparametric Bootstrap Confidence Intervals with the BCa Method
##
##
                 Estimate 95% CI Lower 95% CI Upper p-value
                                0.0504
## ACME
                   0.0635
                                               0.08 <2e-16 ***
                                0.1000
                                               0.10 <2e-16 ***
## ADE
                   0.1000
## Total Effect
                   0.1635
                                0.1504
                                               0.18 <2e-16 ***
                                               0.43 <2e-16 ***
## Prop. Mediated
                   0.3884
                                0.3336
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Sample Size Used: 1000
##
##
## Simulations: 1000
```

Binary outcome and continous mediator

Results for binary outcome and continuous mediator are not on the right magnitude. The estimated total effect is only 0.01 where the true total effect is 0.16.

```
n = 1000
#C is the covariate
C = rnorm(n)
#X is the treatment variable
X = rnorm(n, mean = 0, sd = 1)
#M is the mediator
M = 0.3*X + 0.5* C + rnorm(n)
score = -3+M*0.2+0.1*X + 0.3* C
P = exp(score)/(1+exp(score))
#Y is the outcome
Y = rbinom(P, size = 1, prob = P)
sum(Y)
```

```
## [1] 52
#direct effect is 0.1
#indirect effect is 0.06
#total effect is 0.16
data = data.frame(Y, X, M, C)
out_model = glm(Y ~ X + M + C, data = data, family = binomial(link = 'logit'))
med_model = lm(M~X + C, data = data)
total_model = glm(Y ~ X + C, data = data, family = binomial(link = 'logit'))
summary(out_model)
```

##

```
## Call:
## glm(formula = Y ~ X + M + C, family = binomial(link = "logit"),
      data = data)
##
## Deviance Residuals:
      Min
                1Q
                    Median
                                  3Q
                                          Max
## -0.9215 -0.3663 -0.2807 -0.2130
##
## Coefficients:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -3.14178
                          0.17324 -18.136 < 2e-16 ***
                          0.14893
                                   0.412
               0.06142
                                             0.680
## X
## M
                          0.14738
               0.17276
                                    1.172
                                             0.241
## C
               0.64572
                          0.16569
                                    3.897 9.74e-05 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 408.73 on 999 degrees of freedom
## Residual deviance: 383.39 on 996 degrees of freedom
## AIC: 391.39
##
## Number of Fisher Scoring iterations: 6
summary(med model)
##
## Call:
## lm(formula = M ~ X + C, data = data)
##
## Residuals:
      Min
               1Q Median
##
                               3Q
                                      Max
## -2.6010 -0.6304 -0.0154 0.6435 3.6997
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
                          0.03074 -0.944
## (Intercept) -0.02902
                                             0.345
## X
               0.30238
                          0.03015 10.028
                                            <2e-16 ***
## C
               0.48442
                          0.03080 15.729
                                            <2e-16 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.972 on 997 degrees of freedom
## Multiple R-squared: 0.2629, Adjusted R-squared: 0.2614
## F-statistic: 177.8 on 2 and 997 DF, p-value: < 2.2e-16
summary(total_model)
## Call:
## glm(formula = Y ~ X + C, family = binomial(link = "logit"), data = data)
## Deviance Residuals:
##
      Min
                1Q
                     Median
                                  3Q
                                          Max
```

```
## -0.8803 -0.3624 -0.2856 -0.2205
                                        2.9068
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -3.1299
                            0.1714 -18.264 < 2e-16 ***
                 0.1102
                            0.1421
                                     0.776
                                              0.438
## X
## C
                 0.7169
                            0.1534
                                     4.673 2.97e-06 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 408.73 on 999 degrees of freedom
##
## Residual deviance: 384.76 on 997 degrees of freedom
## AIC: 390.76
##
## Number of Fisher Scoring iterations: 6
library(mediation)
fit_model = mediate(med_model, out_model, treat="X", mediator= "M", boot=T, boot.ci.type = "bca")
## Running nonparametric bootstrap
#result from mediation package is not on the right magnitude.
summary(fit model)
##
## Causal Mediation Analysis
## Nonparametric Bootstrap Confidence Intervals with the BCa Method
##
##
                            Estimate 95% CI Lower 95% CI Upper p-value
## ACME (control)
                             0.00256
                                         -0.00180
                                                                   0.22
                                                           0.01
## ACME (treated)
                             0.00269
                                         -0.00173
                                                           0.01
                                                                   0.22
## ADE (control)
                             0.00302
                                         -0.01112
                                                           0.02
                                                                   0.69
## ADE (treated)
                             0.00315
                                         -0.01212
                                                          0.02
                                                                   0.69
## Total Effect
                             0.00571
                                         -0.01003
                                                           0.02
                                                                   0.51
## Prop. Mediated (control)
                                                        4081.85
                                                                  0.58
                             0.44776
                                          4.84618
## Prop. Mediated (treated)
                             0.47146
                                          4.48338
                                                        3848.42
                                                                   0.58
                                                          0.01
## ACME (average)
                             0.00263
                                         -0.00180
                                                                  0.22
## ADE (average)
                             0.00309
                                         -0.01160
                                                           0.02
                                                                  0.69
                                                                  0.58
## Prop. Mediated (average) 0.45961
                                          4.65026
                                                       3965.13
## Sample Size Used: 1000
##
##
## Simulations: 1000
```