# PSYCGR01: Statistics

## Moderation and mediation worksheet

Maarten Speekenbrink UCL Experimental Psychology November, 2016

### 1 General Exercises

1. Wuensch et al. (2002) conducted a study on the relation between misanthropy, idealism, and attitudes towards animal rights. Their hypothesis was that when evaluating the ethical status of an action that harms a nonhuman animal, one might weight the benefit to humankind against the cost of the harm done to the animal. To the extent that one does not like humans (i.e., the extent of misanthropy), one will not be likely to think that benefits to humans can justify harm to animals. However, they also thought that this relation would be moderated by people's level of idealism: misanthropy would be less strongly related to support for animal rights among idealists (who tend not to do cost-benefit analysis) than among non-idealists. Using (centered) measures of support for animal rights (AR), misanthropy (MT) and idealism (Id), they estimated the following model

$$AR_i = .30 \times MT_i + .07 \times Id_i - .15 \times (Id \times MT)_i + e_i$$

Assume that all parameters were significant.

- (a) What is the predicted level of support for animal rights for someone with a score of 0 on misanthropy and idealism? What does this prediction reflect?

  Answer: The predicted level is 0. As all variables are centered, the prediction is an average level of support when both idealism and misanthropy are at an average level.
- (b) What is the predicted relation between support for animal rights and misanthropy for someone with a score of -1 on idealism?

Answer:

$$\begin{split} \text{AR}_i &= .30 \times \text{MT}_i + .07 \times (-1) - .15 \times ((-1) \times \text{MT})_i + e_i \\ &= -0.07 + .45 \times \text{MT}_i + e_i \end{split}$$

(c) What is the predicted relation between support for animal rights and misanthropy for someone with a score of 1 on idealism?

Answer:

$$AR_i = 0.07 + .15 \times MT_i + e_i$$

(d) What is the predicted relation between support for animal rights and idealism for someone with a score of -1 on misanthropy?

Answer:

$$AR_i = -.30 + .22 \times Id_i + e_i$$

(e) What is the predicted relation between support for animal rights and idealism for someone with a score of 1 on misanthropy?

Answer:

$$AR_i = .30 - .08 \times Id_i + e_i$$

(f) From these predictions, do you think the researchers' hypothesis is supported? Answer: Yes. The interaction is significant and you can see that the relation between misanthropy and support for animal rights changes as a function of idealism. Note that there is no indication of causality, e.g., from the results, one might just as well say that the relation between support for animal rights and idealism is moderated by misanthropy.

### 2 SPSS Exercises

- 1. Open the dataset Geller.sav. This is the same dataset you analysed in the "Regression models" worksheet. For a description of the variables, you can look there. Recall that the hypothesis of Geller et al. was that the extent to which perceptions of weight and shape determine self-esteem affects the development of eating disorders. With that in mind, it seems plausible that for people who have a high level of self-esteem, the SAWBS measure might be largely irrelevant when predicting eating disorders. People with low self-esteem on the other hand, might develop eating disorders when perceptions of weight and shape play a defining role in their (low) self-esteem. This prediction amounts to an interaction between SAWBS and self-esteem (measured by the RSES), or in other words, that the relation between SAWBS and level of eating disorders is moderated by general self-esteem. Let's assess whether this prediction holds, using the HIQ measure of the level of eating disorders.
  - (a) Estimate the parameters of MODEL C:

$$\text{HIQ}_i = \beta_0 + \beta_1 \text{RSES}_i + \beta_2 \text{SAWBS}_i + \epsilon_i$$

and compute the SSE.

```
> library(foreign)
> geller <- as.data.frame(read.spss("Geller.sav"))
> MODEL_C <- lm(hiq ~ rses + sawbs,data=geller)
> summary(MODEL_C)
```

```
Call:
lm(formula = hiq ~ rses + sawbs, data = geller)
Residuals:
             1Q Median
                              3Q
    Min
                                     Max
-14.3039 -4.6371 0.2091 4.4005 15.2705
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 29.29943 3.74596
                              7.822 1.69e-11 ***
                    0.08999 -5.419 6.01e-07 ***
          -0.48769
rses
           sawbs
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 6.223 on 81 degrees of freedom
Multiple R-squared: 0.5436, Adjusted R-squared: 0.5323
F-statistic: 48.24 on 2 and 81 DF, p-value: 1.599e-14
> SSE_C <- sum(residuals(MODEL_C)^2)</pre>
> SSE_C
[1] 3137.012
```

$$\label{eq:hiq} \text{HIQ}_i = 29.299 - 0.488 \times \text{RSES}_i + 0.088 \times \text{SAWBS}_i + \epsilon_i$$
 and SSE(C) = 3137.012

(b) Estimate the parameters of MODEL A1:

$$\mbox{HIQ}_i = \beta_0 + \beta_1 \mbox{RSES}_i + \beta_2 \mbox{SAWBS}_i + \beta_3 (\mbox{RSES} \times \mbox{SAWBS})_i + \epsilon_i$$
 and compute the SSE.

```
> MODEL_A1 <- lm(hiq ~ rses + sawbs + rses:sawbs,data=geller)
> summary(MODEL_A1)

Call:
lm(formula = hiq ~ rses + sawbs + rses:sawbs, data = geller)

Residuals:
    Min    1Q    Median    3Q    Max
-14.2729   -4.4008    0.1306    4.4596    15.1809

Coefficients:
```

```
Estimate Std. Error t value Pr(>|t|)

(Intercept) 31.110904    5.451364    5.707   1.87e-07 ***

rses         -0.536468    0.139465   -3.847   0.000239 ***

sawbs         0.062231    0.057902    1.075   0.285711

rses:sawbs         0.000734    0.001598    0.459   0.647181
---

Signif. codes:    0 '***'   0.001 '**'   0.05 '.'   0.1 ' ' 1

Residual standard error:   6.254 on 80 degrees of freedom

Multiple R-squared:   0.5448,Adjusted R-squared:   0.5277

F-statistic:   31.91 on 3 and 80 DF, p-value:   1.144e-13

> SSE_A1 <- sum(residuals(MODEL_A1)^2)
> SSE_A1

[1] 3128.757
```

```
\text{HIQ}_i = 31.111 - 0.536 \times \text{RSES}_i + 0.062 \times \text{SAWBS}_i + 0.001(\text{RSES} \times \text{SAWBS})_i + e_i and \text{SSE}(\text{A1}) = 3128.757.
```

(c) What is the null-hypothesis you test when comparing MODEL A1 to MODEL C? Test this null-hypothesis.

```
> anova(MODEL_C,MODEL_A1)
Analysis of Variance Table

Model 1: hiq ~ rses + sawbs
Model 2: hiq ~ rses + sawbs + rses:sawbs
   Res.Df   RSS Df Sum of Sq   F Pr(>F)
1   81 3137.0
2   80 3128.8   1   8.2546 0.2111 0.6472
```

Answer:  $H_0: \beta_3 = 0$ , F = 0.211,  $P(F_{1,80} \ge 0.221) = .647$ , PRE = 0.00263, so there is no reason to reject the null hypothesis that there is no interaction.

(d) Compute the correlations between the three predictors in MODEL A1. Assess the level of multicollinearity.

```
> ## an alternative way is to get the predictors of
> ## MODEL_A1 via the model.matrix function:
> preds <- model.matrix(MODEL_A1)</pre>
> head(preds)
  (Intercept)
                           sawbs rses:sawbs
                  rses
            1 32.22844 56.93904 1835.0564
1
2
            1 32.39728 117.16003 3795.6668
3
            1 44.93232 48.46506 2177.6473
            1 38.63428 66.88492 2584.0507
4
5
            1 35.48696 88.12559 3127.3093
6
            1 29.77724 -22.88933
                                 -681.5811
> ## Note that the model.matrix has an (intercept) predictor
> ## which always equals 1. This is an artifical predictor
> ## such that the slope of this predictor equals the intercept
> ## (a slope for a predictor which always equals 1 is exactly
> ## like an intercept!)
> cor(preds[,2:4])
                            sawbs rses:sawbs
                  rses
            1.00000000 -0.3782735 -0.04348224
rses
sawbs
           -0.37827350 1.0000000 0.90549922
rses:sawbs -0.04348224 0.9054992 1.00000000
> ## a better way to assess multicollinearity
> library(car)
> vif(MODEL_A1)
               sawbs rses:sawbs
      rses
  2.775567 15.384581 13.208161
> # or tolerance
> 1/vif(MODEL_A1)
                sawbs rses:sawbs
      rses
0.36028671 0.06500014 0.07571076
```

Answer: SAWBS and RSES have a moderate (but significant) negative correlation of r = -.378. The main problem is the correlation between SAWBS and the interaction term, which is r = .905.

(e) There has been a long-standing debate on the merits of centering predictors in interactive regression models. Let's see what happens when we center the predictors first. Compute two new predictors as

```
\begin{split} & \text{RSESO}_i = \text{RSES}_i - \overline{\text{RSES}} \\ & \text{SAWBSO}_i = \text{SAWBS}_i - \overline{\text{SAWBS}} \end{split}
```

and estimate the MODEL A2:

$$\text{HIQ}_i = \beta_0 + \beta_1 \text{RSESO}_i + \beta_2 \text{SAWBSO}_i + \beta_3 (\text{RSESO} \times \text{SAWBSO})_i + \epsilon_i$$

```
> geller$rses0 <- scale(geller$rses,scale=FALSE)</pre>
> geller$sawbs0 <- scale(geller$sawbs,scale=FALSE)</pre>
> MODEL_A2 <- lm(hiq ~ rses0*sawbs0,data=geller)
> summary(MODEL_A2)
Call:
lm(formula = hiq ~ rses0 * sawbs0, data = geller)
Residuals:
     Min
               1Q
                   Median
                                 30
                                         Max
-14.2729 -4.4008 0.1306
                            4.4596
                                    15.1809
Coefficients:
              Estimate Std. Error t value Pr(>|t|)
(Intercept) 16.904612 0.719333 23.500 < 2e-16 ***
rses0
             -0.494188
                         0.091531 -5.399 6.68e-07 ***
                                   5.522 4.03e-07 ***
sawbs0
              0.088656
                         0.016055
rses0:sawbs0 0.000734
                         0.001598
                                    0.459
                                             0.647
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
Residual standard error: 6.254 on 80 degrees of freedom
Multiple R-squared: 0.5448, Adjusted R-squared: 0.5277
F-statistic: 31.91 on 3 and 80 DF, p-value: 1.144e-13
```

#### Answer:

$$\mathtt{HIQ}_i = 16.905 - 0.494 \times \mathtt{RSESO}_i + 0.089 \times \mathtt{SAWBSO}_i + 0.001 \times (\mathtt{RSESO} \times \mathtt{SAWBSO})_i + \epsilon_i$$

Note that the parameter value for the interaction stays the same, but those of the individual predictors change. By centering, the predictors will have a value of 0 when they are at their mean values, and the slopes of the predictors in an interaction model are the slopes when the other predictors are 0. For example, the slope of the centered predictor  $\mathtt{RSESO}_i$  is the slope when  $\mathtt{SAWBSO}_0 = 0$ , so at  $\mathtt{SAWBS}$ .

(f) Compare the results of MODEL A2 (significance of individual predictors and overall model fit) to those obtained for MODEL A1. Are there differences? If so, can you explain these?

Answer: There is no difference for the interaction effect, but now the individual effect of SAWBS is significant. By centering, we are now estimating the "simple"

effect of SAWBS when RSES is at its mean. This is different from the effect of SAWBS when RSES = 0. Note that the  $R^2$  of this model is the same as for the uncentered model. So this model will make the same predictions (and is "in a deep sense" the same) as the uncentered model.

(g) Compute the correlations between the three predictors in MODEL A2 and assess the level of multicollinearity. Are there differences with model A1 in this respect?

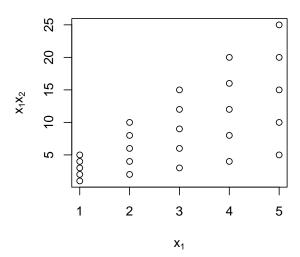
```
> cor(model.matrix(MODEL_A2)[,2:4])
                  rses0
                            sawbs0 rses0:sawbs0
rses0
              1.0000000 -0.3782735
                                      0.2102434
sawbs0
             -0.3782735 1.0000000
                                     -0.1841580
rses0:sawbs0 0.2102434 -0.1841580
                                      1.0000000
> # tolerance
> 1/vif(MODEL_A2)
                   sawbs0 rses0:sawbs0
       rses0
   0.8364523
                0.8454558 0.9430226
```

Answer: The correlation between SAWBS and RSES has not changed, but the correlation between SAWBS and the interaction term is much smaller. To see why this is, let's take a simple example of two predictors,  $X_1$  and  $X_2$ , each with values from 1 to 5. We can compute the interaction  $X_1 \times X_2$  as in the table below

	$X_1$						
$X_2$	1	2	3	4	5		
1	1	2	3	4	5		
2	2	4	6	8	10		
3	3	6	9	12	15		
4	4	8	12	16	20		
5	5	10	15	20	25		

If we plot the values of  $X_1$  against the interaction  $X_1X_2$  we get the following

#### uncentered

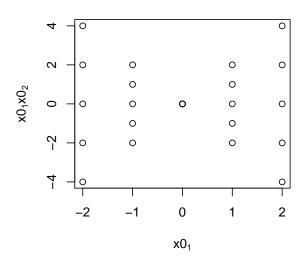


Which shows that there is a positive relation between  $X_1$  and the interaction  $X_1X_2$ . If we first center  $X_1$  and  $X_2$  (let's call these centered variables  $X0_1$  and  $X0_2$  respectively) and then compute the interaction, we get the values in the table below:

	$X0_1$							
$X0_2$	-2	-1	0	1	2			
-2	4	2	0	-2	-4			
-1	2	1	0	-1	-2			
0	0	0	0	0	0			
1	-2	-1	0	1	2			
2	-4	-2	0	2	4			

If we plot the values of  $X0_1$  against the interaction  $X0_1 \times X0_2$  we get the following

#### centered



in which there is no evidence of a positive (or negative) correlation. The main reason why centering can reduce multicollinearity is that centering makes sure that for each predictor there are both positive and negative values. When we then compute a product of these variables, two negative values become positive in the product, while one negative value and one positive value become negative in the product. So, if a centered predictor has a negative value, the product (interaction) predictor can be both positive and negative (depending on the value of the other predictor).

- 2. This exercise is also based on the Geller.sav dataset. Let's attempt to investigate the researchers' hypothesis more directly, which we can view as one of mediation. Namely, it is assumed that perceptions of weight and shape influence self-esteem, and the extent to which they do determines the level of eating disorders. In other words, the effect of weight and shape perceptions on eating disorders is assumed to be mediated by self-esteem.
  - (a) First, calculate an aggregated measure of weight and shape perceptions as the mean of these two values (for each participant) and call this measure WSPercep.

```
> geller$wspercep <- (geller$wtpercep + geller$shpercep)/2
> # alternatively, you can use
> # geller£wspercep <- rowMeans(geller[,c("wtpercep","shpercep")])</pre>
```

(b) Estimate the parameters of MODEL 1:

$$\text{HIQ}_i = \beta_0 + \beta_1 \text{WSPercep}_i + \epsilon_i$$

```
> MODEL_1 <- lm(hiq ~ wspercep,data=geller)
> summary(MDOEL_1)
Error in summary(MDOEL_1): object 'MDOEL_1' not found
```

$$\mathtt{HIQ}_i = 41.69 - 6.73 \times \mathtt{WSPercep}_i + e_i$$

(c) Estimate the parameters of MODEL 2:

$$RSES_i = \beta_0 + \beta_1 WSPercep_i + \epsilon_i$$

```
> MODEL_2 <- lm(rses ~ wspercep,data=geller)</pre>
> summary(MODEL_2)
Call:
lm(formula = rses ~ wspercep, data = geller)
Residuals:
             1Q Median
    Min
                              3Q
                                       Max
-17.4672 -4.2555 0.2929 4.1053 19.2571
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 19.7249 3.6141 5.458 5.01e-07 ***
                      0.9524 4.618 1.42e-05 ***
         4.3987
wspercep
___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 7.349 on 82 degrees of freedom
Multiple R-squared: 0.2064, Adjusted R-squared: 0.1967
F-statistic: 21.33 on 1 and 82 DF, p-value: 1.417e-05
```

Answer:

$$RSES_i = 19.72 + 4.40 \times WSPercep_i + e_i$$

(d) Estimate the parameters of MODEL 3:

$$\mathtt{HIQ}_i = \beta_0 + \beta_1 \mathtt{WSPercep}_i + \beta_2 \mathtt{RSES}_i + \epsilon_i$$

```
> MODEL_3 <- lm(hiq ~ wspercep + rses,data=geller)
> summary(MODEL_3)
Call:
lm(formula = hiq ~ wspercep + rses, data = geller)
Residuals:
     Min
                   Median
                                 30
                                         Max
               1Q
                   0.4411
-11.7633 -4.1639
                            4.3885
                                    16.3544
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) 50.6493
                        3.6454 13.894 < 2e-16 ***
wspercep
            -4.7309
                        0.9236 -5.122 2.01e-06 ***
rses
             -0.4540
                        0.0954 -4.759 8.37e-06 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 6.349 on 81 degrees of freedom
Multiple R-squared: 0.525, Adjusted R-squared: 0.5132
F-statistic: 44.76 on 2 and 81 DF, p-value: 8.085e-14
```

$$\mathtt{HIQ}_i = 50.65 - 4.73 \times \mathtt{WSPercep}_i - 0.45 \times \mathtt{RSES}_i + e_i$$

- (e) Assess the mediation hypothesis by assessing the significance of the parameters in the three models estimated above (i.e. following the causal steps procedure). Answer: The regression coefficient (slope) for WSPercep in the first model (with HIQ as dependent variable) is significant, as is the coefficient for WSPercep in the second model (with RSES as dependent variable). In the third model, both regression coefficients are significant. However, the significance of the WSPercep effect is reduced, from F = 52.90 in the first model to F = 26.24 in the third model. So, while the effect of weight and shape perceptions is not completely mediated by self-esteem, there is evidence of partial mediation.
- (f) Use the Sobel-Aroian test to test the significance of the indirect effect of WSPercep on HIQ.

```
> mySobelTest <- function(a,b,se_a,se_b) {
+     Z <- (a*b)/sqrt((a^2 * se_b^2 + b^2 * se_a^2 + se_a^2*se_b^2))
+     # two-sided p-value
+     p <- 2*(1-pnorm(abs(Z)))
+     cat("Sobel-Aroian Test \n")
+     cat(paste("Z = ",Z,", p = ",p,"\n",sep=""))</pre>
```

```
+ }
>
> a <- summary(MODEL_2)$coefficients["wspercep","Estimate"]
> se_a <- summary(MODEL_2)$coefficients["wspercep","Std. Error"]
> b <- summary(MODEL_3)$coefficients["rses","Estimate"]
> se_b <- summary(MODEL_3)$coefficients["rses","Std. Error"]
> mySobelTest(a,b,se_a,se_b)

Sobel-Aroian Test
Z = -3.2772581185632, p = 0.00104820497747915
```

$$Z = \frac{a \times b}{\sqrt{a^2 \times s_b^2 + b^2 \times s_a^2 + s_a^2 \times s_b^2}}$$

$$= \frac{4.40 \times (-0.45)}{\sqrt{4.40^2 \times (0.095)^2 + (-0.45)^2 \times (0.952)^2 + (0.952)^2 \times (0.095)^2}}$$

$$= -3.28$$

Using the standard-normal distribution, the probability of obtaining this value or a smaller one is  $P(Z \le -3.28) = .0005$ . To perform a two-sided test (which is what we'd normally want to do), we need  $P(|Z| \ge 3.28) = P(Z \le -3.28) + P(Z \ge 3.28) = 2 \times .0005 = .001$ , which is smaller than  $\alpha = .05$ , so the mediated effect of WSPercep on HIQ, via RSES, is significant.

(g) Now, let's look at another possibly mediated effect. Previous research (e.g., Miotto et al., 2002) has found a significant relation between social desirability and level of eating disorders. Let's assess whether this effect is mediated by depression. Use both the causal steps and Sobel-Aroian procedures to assess whether the effect of SocDesir on HIQ is mediated by BDI.

```
socdesir -1.6216 0.4695 -3.454 0.000876 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 8.554 on 82 degrees of freedom
Multiple R-squared: 0.127, Adjusted R-squared: 0.1164
F-statistic: 11.93 on 1 and 82 DF, p-value: 0.0008762
> summary(lm(bdi ~ socdesir,data=geller))
Call:
lm(formula = bdi ~ socdesir, data = geller)
Residuals:
    Min
              1Q
                 Median
                               3Q
                                       Max
-15.8848 -4.8417 0.3759 4.1713 17.4800
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 16.4399 2.0193 8.141 3.67e-12 ***
socdesir
          -1.1880 0.3753 -3.166 0.00217 **
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
Residual standard error: 6.838 on 82 degrees of freedom
Multiple R-squared: 0.1089, Adjusted R-squared: 0.09803
F-statistic: 10.02 on 1 and 82 DF, p-value: 0.002173
> summary(lm(hiq ~ socdesir + bdi,data=geller))
Call:
lm(formula = hiq ~ socdesir + bdi, data = geller)
Residuals:
    Min
              1Q
                 Median
                               30
-13.4835 -6.3022 0.3493 5.0465 17.0102
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
                       2.7068 4.563 1.77e-05 ***
(Intercept) 12.3511
                       0.3963 -1.802 0.0752 .
socdesir
            -0.7142
             0.7638
bdi
                       0.1101 6.939 8.85e-10 ***
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

```
Residual standard error: 6.816 on 81 degrees of freedom
Multiple R-squared: 0.4525, Adjusted R-squared: 0.439
F-statistic: 33.47 on 2 and 81 DF, p-value: 2.543e-11
> # sobel-aroian test
> # As an example of "good" R use
> # let's write a simpler R function for this...
> simpleSobelTest <- function(source, mediator, dependent) {</pre>
   # fit model 2
  mod <- lm(mediator ~ source)
    a <- summary(mod)$coefficients["source", "Estimate"]</pre>
  se_a <- summary(mod)$coefficients["source", "Std. Error"]</pre>
+
   # fit model 3
+
  mod <- lm(dependent ~ source + mediator)</pre>
  b <- summary(mod)$coefficients["mediator", "Estimate"]</pre>
  se_b <- summary(mod)$coefficients["mediator", "Std. Error"]</pre>
  Z \leftarrow (a*b)/sqrt((a^2 * se_b^2 + b^2 * se_a^2 + se_a^2*se_b^2))
   # two-sided p-value
  p \leftarrow 2*(1-pnorm(abs(Z)))
  result <- list(
+
     "Z" = Z,
      "p-value" = p
+
+
  cat("Sobel-Aroian Test \n\n")
   cat("Path:",deparse(substitute(source)),"->",
        deparse(substitute(mediator)),"->",
+
        deparse(substitute(dependent)),"\n")
    cat("a * b = ",a*b,"\n\n")
+
    # note: the signif function "rounds" things by a specified
            number of significant (non-zero) digits
    cat(paste("Z = ", signif(Z,3), ", p = ", signif(p,3), "\n", sep=""))
    return(result)
+ }
>
> result <- simpleSobelTest(source=geller$socdesir,</pre>
                               mediator=geller$bdi,
                               dependent=geller$hiq)
Sobel-Aroian Test
Path: geller$socdesir -> geller$bdi -> geller$hiq
a * b = -0.9073957
Z = -2.86, p = 0.0043
```

> result

\$Z

[1] -2.855555

\$`p-value`

[1] 0.004296164

Answer: In model

$$\mathtt{HIQ}_i = 24.91 - 1.62 \times \mathtt{SocDesir}_i + e_i$$

the effect of SocDesir is significant, F(1,82) = 11.93, p < .001. In model

$$\mathtt{BDI}_i = 16.44 - 1.19 \times \mathtt{SocDesir}_i + e_i$$

the effect of SocDesir is significant, F(1,82) = 10.02, p = .002. In model

$$\mathtt{HIQ}_i = 12.35 + 0.76 \times \mathtt{BDI}_i - 0.71 \times \mathtt{SocDesir}_i + e_i$$

the effect of BDI is significant, F(1,81) = 48.15, p < .001, but the effect of SocDesir is not, F(1,81) = 3.25, p = .075. So, according to the causal steps procedure, there is evidence that the effect of social desirability on eating disorders is *completely* mediated by depression. Using the Sobel-Aroian test, we can assess whether this mediated effect itself is significant:

$$Z = \frac{(-1.19) \times (0.76)}{\sqrt{(-1.19)^2 \times (0.11)^2 + (0.76)^2 \times (0.375)^2 + (0.375)^2 \times (0.11)^2}}$$
  
= -2.85

As  $P(Z \le -2.85) = .0022$ , a two sided tests gives  $P(|Z| > 2.85) = 2 \times .0022 = .0044$ . So the indirect effect of social desirability on eating disorders (mediated by depression) is significant.