Statistical Analysis Using Structural Equation Models

EPsy 8266

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Your paper

- 1. What was the main RQ or aim in the paper you found?
- 2. What model(s) did the authors fit?
- 3. How did they evaluate their models?
- 4. Were the models recursive/nonrecursive?
- 5. What kind of claims/language do the authors use? Was it causal or correlational?

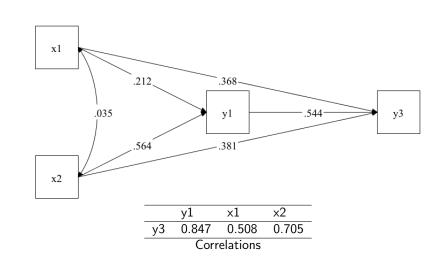
Decomposing and calculating effects

- Direct effect direct influence of variable of interest (X) on another variable (DV) that is unmediated by any other variable, i.e. each single headed arrow represents a direct effect.
- ▶ Indirect effects start from the DV later in the path diagram (on the right). Trace backwards (right to left) against arrows passing intervening variables until you get to X. Each combination of intervening variables is a separate indirect effect.
- Spurious effects (due to common causes) Start from DV. Trace backwards to a variable, Z, that has a direct or indirect effect on both X and DV. Move from Z to X. There are as many spurious effects of X on DV due to Z as many ways you can get from DV to X through Z following the rule above.

Correlated (unanalyzed) effects

- If X is exogenous, find variable Z that is both exogenous and has a direct or indirect effect on DV. Start from variable DV. Trace back to Z. Make the last step through the double headed arrow to X.
- If X is endogenous, find an exogenous variable Z that has a direct or indirect effect on DV and is correlated to another exogenous variable W that has a direct or indirect effect on X. Start from variable DV. Trace back to Z. Travel through the double headed arrow to W. Move from W to X.

Decomposing effects activity



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Count me out of the supposedly "emerging consensus that mediation analysis requires data from designs with time precedence" Hayduk (2018)

Is relationship of between X and Y mediated by M?

The Baron & Kenny approach: a series of if-then statements

- 1. Is there a statistically significant relationship between X and Y? If yes, record the effect as c. Then
- 2. Is there a statistically significant relationship between X and M? If yes, record the effect as a, Then.
- 3. Is there a statistically significant relationship between M and Y? If yes, then.
- 4. Estimate the effect that X and M have simultaneously on Y. Record the M on Y effect as b, record the X on Y effect as c'

If c' is not significantly, then the causal effect of X on Y is fully mediated by M. If |c'| is less than |c|, then the causal effect of X on Y is partially mediated by M.

Sometimes, presented as $\frac{|c-c'|}{c}$, the percent of the total causal effect of X on Y explained by the mediator M.

Approaches to testing indirect effects

- ▶ Baron & Kenny's causal steps approach (Bad)
 - ▶ Do a series of regressions and determine if c' is < c.
 - Not a test of mediation (ab pathway)
- Product of coefficients approach (Sobel test) (Not so good)
 - Determine if ab is significant using a Wald test
 - Significance test of ab is problematic. Assumes normal distributions of SEs.
- SEM estimation of indirect effect (Not great)
 - Determine if ab is significant.

```
lavaan.mod <- "
y1 ~ a*x1 + c*x2 + e*x3;
y3 ~ b*y1 + d*x1 + f*x2;
# indirect effect of x1 on y3
ab := a*b
"</pre>
```

 Use ML estimation to determine significance of indirect effect (still assumes normality assumption)

The bootstrap

- In traditional statistics, standard errors and confidence intervals are based on theoretical sampling distributions of the parameter estimates.
- ► These have certain distributional assumptions and/or rely on asymptotic theory (i.e. when we have a large sample)
- Often we violate these assumptions (i.e., we have non-normal data) or our sample size is really too small for these statistics to be accurate.
- ▶ One thing we can do is to use the bootstrap developed by Efron (1982).
- It creates an empirical sampling distribution of parameter estimates that you can use in an analogous fashion to the theoretical sampling distribution.

How it works

- 1. Let's say your data set is size n. Fit the model and obtain some parameter estimate β .
- 2. Take a sample size of n from your data set with replacement. This is the first bootstrap sample.
- 3. Use this sample, fit your model again and obtain your parameter estimate for β .
- 4. Repeat steps 2 3, a large number of times. Let's call this k times.
- 5. The distribution of your k estimates of β is your empirical sampling distribution. From this you can calculate:
 - Standard error of β This is just the standard deviation of this distribution.
 - 95% confidence interval Locate the 2.5% percentile and the 97.5% percentile.
 - A bias correction is recommended.

This whole resampling procedure is known as **non-parametric bootstrapping**.

Method 1: The non-parametric bootstrap

This toy example is using the mtcars data set (?mtcars). Let's say we want to predict mile per gallon given the weight of the car. We want to use the bootstrap on the slope of weight.

```
step1 <- coef(lm(mpg ~ wt, data = mtcars))[[2]]
step1
## [1] -5.344472</pre>
```

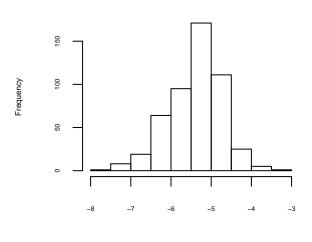
Our estimate is that for 1000 lbs, we expect that miles per gallon to decrease by 5.34.

Method 1: The R code

```
k <- 500
n <- nrow(mtcars)
beta.weight <- rep(NA, k)
for(i in 1:k){
  obs <- sample(x = n, size = n, replace = T)
  tmp <- mtcars[obs,]
  beta.weight[i] <- coef(lm(mpg ~ wt, data = tmp))[[2]]
}</pre>
```

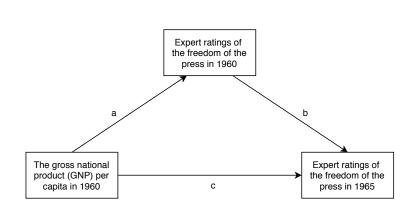
Method 1: Plotting the empirical distribution

Empirical Sampling Distribution of the Slope of Weight



Method 1: Using the empirical distribution

```
# Calculate a 95% confidence interval
quantile(beta.weight, probs = c(.025, .975))
## 2.5% 97.5%
## -6.796561 -4.189005
```



Bootstrapping the indirect effect

```
model <- '
  v1 ~ a*x1
  y5 \sim c*x1 + b*y1
  # indirect effects
  ab := a*b
  totl := c + a*b
fit <- sem(model, data = PoliticalDemocracy)
summary(fit)
# define the function to extract the indirect effect
boot.fun <- function(x) {</pre>
  parameterEstimates(x)[parameterEstimates(x)$label %in% c("ab"), "est"]
# run the bootstrap
# - by default this is nonparameteric
ab.dist <- bootstrapLavaan(fit, R = 5000, FUN = boot.fun)
ab.dist <- data.frame(ab.dist)
names(ab.dist) <- c("ab")</pre>
hist(ab.dist$ab)
quantile(ab.dist$ab, probs = c(.025, .975))
```

Bootstrapping a covariance matrix

- ▶ If you have just a covariance matrix because the data are not available, then you can use a parametric bootstrap, aka a Monte Carlo bootstrap.
 - To do this, you generate a sample of size n from a multivariate normal distribution with a covariance matrix equal to the covariance matrix you've obtained.
 - 2. This can be done in R using MASS::mvrnorm
 - 3. We we aren't resampling k, we are generating k sets of new data with our observed covariance matrix.
 - 4. Note, this relies on multivariate normality! But we don't have the data, so maybe it's multivariate normal?
 - 5. This is very useful for obtaining standard errors when we don't have a theoretical distribution. So, with latent variables, this can be useful getting SEs for standardized loadings and indirect paths.
 - 6. We want a large k, the larger the better. Computing is fast and cheap nowadays!
 - 7. The larger the k, the more smoother the distribution will be.

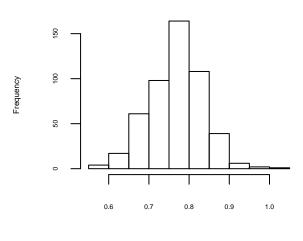
Method 2: The parametric bootstrap from the HZ covariance

In this example, we are trying to get a 95% CI around the standardized coefficient for visual on x1.

```
library("lavaan")
library("MASS")
hs.cov <- cov(HolzingerSwineford1939[,7:ncol(HolzingerSwineford1939)])
k <- 500
n < -301
HS.model \leftarrow 'visual = x1 + x2 + x3
               textual = x4 + x5 + x6
               speed = x7 + x8 + x9
est.par <- rep(NA, k)
for(i in 1:k){
  tmp <- mvrnorm(n, mu = rep(0, ncol(hs.cov)), Sigma = hs.cov)</pre>
  tmp <- as.data.frame(tmp)</pre>
  fit.tmp <- cfa(HS.model, tmp)</pre>
  est.par[i] <- standardizedSolution(fit.tmp)[1,4]</pre>
```

Method 2: Plotting the empirical distribution

Standardized factor loading of Visual on X1

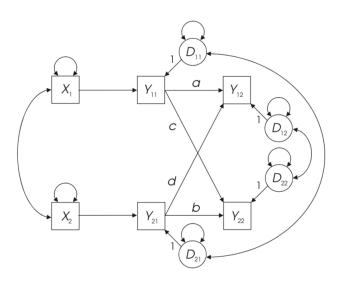


Std factor loading

Method 2: Using the empirical distribution

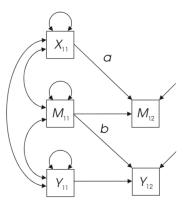
```
# Point estimate
mean(est.par)
## [1] 0.7693568
# Calculate a 95% confidence interval
quantile(est.par, probs = c(.025, .975))
## 2.5% 97.5%
## 0.6356613 0.8898625
# Truth?
fit.true <- cfa(HS.model, HolzingerSwineford1939)</pre>
standardizedSolution(fit.true)[1,]
       lhs op rhs est.std se z pvalue
## 1 visual = ^{\circ} x1 0.772 0.055 14.041
```

Panel model

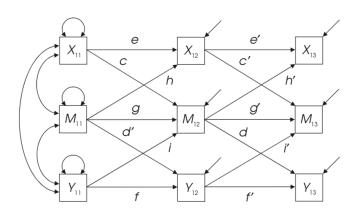


Panel mediation model 1

(a) Half longitudinal mediation



Panel mediation model 2



Non-recursive models

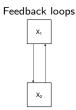
- ▶ Models with reciprocal paths (feedback loops)
- Models with correlated errors

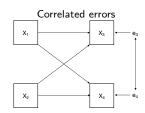
When models include one of these, then the assumption of predictors being uncorrelated with errors and/or errors being uncorrelated is broken

OLS regression will not yield unbiased estimators

Estimation for Path Analysis

A path analysis model is recursive if it doesn't contain:





- ▶ If the path analysis model has feedback loops and/or has correlated errors then a method that estimates the paths simultaneously must be used. For example maximum likelihood for the multivariate vector of all observed variables
- Note that the simultaneous ML method can also be used when the model is recursive. This is simpler than performing several different regressions because it is done all in one step.

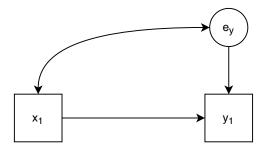
Reciprocal paths

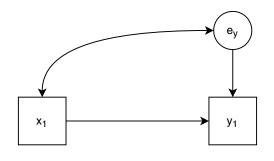


- 1. One variable causes another and it in turn causes the first and so on.
- 2. Equilibrium must be assumed, i.e. direct effects must remain stable throughout an infinite number of reverberations.
- 3. Total effect is calculated as the infinite sum

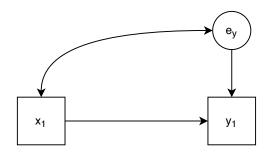
Identifiability

- ▶ Necessary and sufficient conditions for the identifiability of models with reciprocal paths are complex (Bollen, 1989).
- ► Each endogenous variable in a feedback loop must have its own unique predictor (instrumental variable).
- ▶ Some people have argued that reciprocal paths can be used to test competing models of causation by identifying the primary cause.



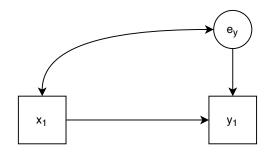


$$Y_1 = \beta X_1 + e_y$$



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Will β be biased?



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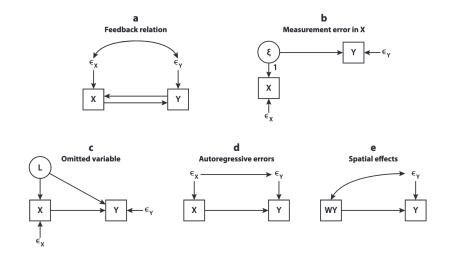
The problem is that $Cov(X, e_y) \neq 0$

Correlated errors

By default, we assume manifest variables are related to one another solely through there relationship through the factor, right?

- Implies that there are other predictor variables (not included in the model) that would cause the two endogenous variables to be correlated
- Good way to allow association between endogenous variables when there is no reason to believe the association is causal
- The model is identified as long as reciprocal direct paths between the variables would be identified.

Bollen, 2012 - Sources of correlated errors



What is an instrumental variable

For a variable Z to be an instrumental variable the following **must** hold

- 1. $Cov(Z, e_y) = 0$
- 2. $Cov(Z, X_1) \neq 0$

This allows the model to no longer be underidentified.

Rigdon's (1995) graphical rules for just identified

