

# Structural Equation Modeling & Mediation

## Introduction to SEM with Lavaan



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# Outline

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Structural Equation Modeling

Mediation

Simple Mediation

Bootstrapping

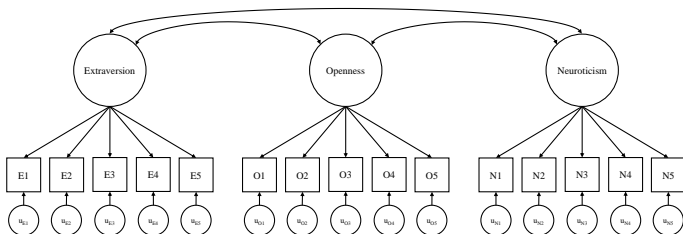


# Full SEM

Structural equation modeling (SEM) simply combines path analysis and CFA.

- SEM allows us to model complicated structural relations among latent variables.

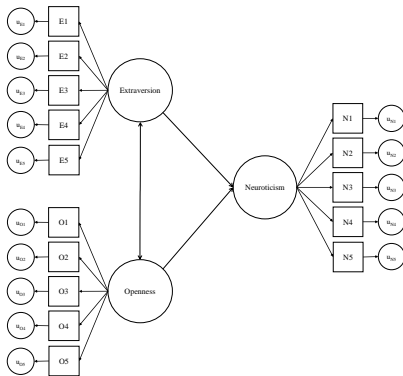
Let's consider a simple, three-factor CFA model.



# CFA → SEM

We first evaluate the validity of the measurement model via CFA.

- We then convert the CFA to an SEM by converting some covariances to latent regression paths.



# Why SEM?

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The beauty of SEM is that we get to model the types of complex relations we can specify via path models while leveraging all the strengths of latent variables.

- When we fit a multiple-group SEM, we are modeling moderation by group.
  - The latent variables give us the ability to evaluate measurement invariance across groups.
  - We'll see more of these ideas in the next lecture.
- Path analysis and SEM lend themselves especially well to mediation analysis and conditional process analysis.



# MEDIATION



# Mediation vs. Moderation

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What do we mean by *mediation* and *moderation*?



# Mediation vs. Moderation

---

What do we mean by *mediation* and *moderation*?

Mediation and moderation are types of hypotheses, not statistical methods or models.

- Mediation tells us *how* one variable influences another.
- Moderation tells us *when* one variable influences another.





# Contextualizing Example

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Say we wish to explore the process underlying exercise habits.

Our first task is to operationalize “exercise habits”

- DV: Hours per week spent in vigorous exercise (*exerciseAmount*).

We may initial ask: what predicts devoting more time to exercise?

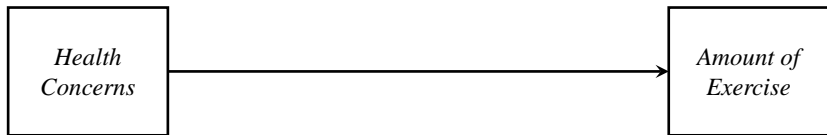
- IV: Concerns about negative health outcomes (*healthConcerns*).



# Focal Effect Only

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The *healthConcerns* → *exerciseAmount* relation is our *focal effect*



- Mediation, moderation, and conditional process analysis all attempt to describe the focal effect in more detail.
- We always begin by hypothesizing a focal effect.

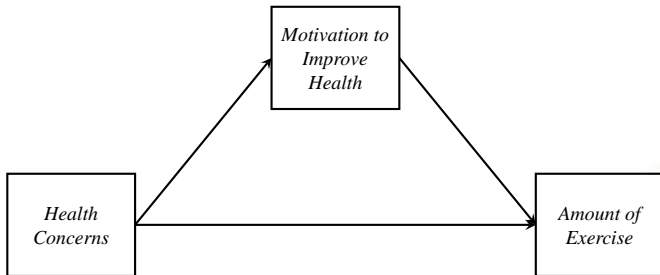


# The Mediation Hypothesis

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A mediation analysis will attempt to describe how health concerns affect amount of exercise.

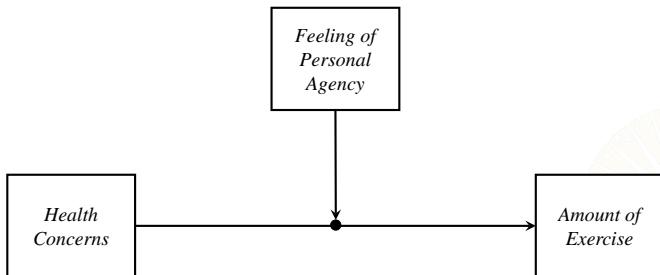
- The *how* is operationalized in terms of intermediary variables.
- Mediator: Motivation to improve health (*motivation*).



# Moderation Hypothesis

A moderation hypothesis will attempt to describe when health concerns affect amount of exercise.

- The *when* is operationalized in terms of interactions between the focal predictor and contextualizing variables
- Moderator: Sense of personal agency relating to physical health (*agency*).



# Conditional Process Analysis

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Conditional process analysis combines the mediation and moderation hypotheses into models of moderated mediation.

- Given a mediation model describing *how* health concerns affect exercise amount, what other variables may modulate the indirect effect.



# Excellent Resource

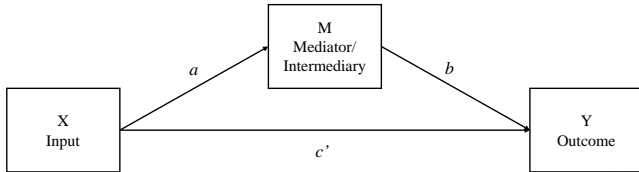
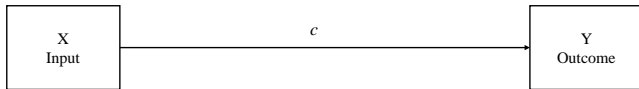
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Plug Hayes' book



# Path Diagrams

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# Necessary Equations

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To get all the pieces of the preceding diagram using OLS regression, we'll need to fit three separate models.

$$Y = i_1 + cX + e_1 \quad (1)$$

$$Y = i_2 + c'X + bM + e_2 \quad (2)$$

$$M = i_3 + aX + e_3 \quad (3)$$

- Equation 1 gives us the total effect ( $c$ ).
- Equation 2 gives us the direct effect ( $c'$ ) and the partialled effect of the mediator on the outcome ( $b$ ).
- Equation 3 gives us the effect of the input on the outcome ( $a$ ).



# Two Measures of Indirect Effect

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Indirect effects can be quantified in two different ways:

$$IE_{diff} = c - c' \quad (4)$$

$$IE_{prod} = a \cdot b \quad (5)$$

$IE_{diff}$  and  $IE_{prod}$  are equivalent in simple mediation.

- Both give us information about the proportion of the total effect that is transmitted through the intermediary variable.
- $IE_{prod}$  provides a more direct representation of the actual pathway we're interested in testing.
- $IE_{diff}$  gets at our desired hypothesis indirectly.



# The Causal Steps Approach

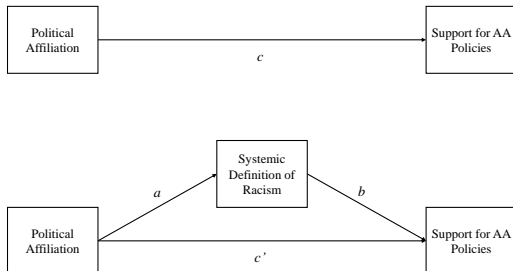
---

Baron and Kenny (1986, p. 1176) describe three/four conditions as being sufficient to demonstrate statistical “mediation.”

1. Variations in levels of the independent variable significantly account for variations in the presumed mediator (i.e., Path  $a$ ).
  - Need a significant  $a$  path.
2. Variations in the mediator significantly account for variations in the dependent variable (i.e., Path  $b$ ).
  - Need a significant  $b$  path.
3. When Paths  $a$  and  $b$  are controlled, a previously significant relation between the independent and dependent variables is no longer significant.
  - Need a significant total effect
  - The direct effect must be “less” than the total effect

# Example Process Model

Consider the following process.



# Causal Steps Example

---

```
## Load some data:
dat1 <- readRDS("../data/adamsKlpsScaleScore.rds")

## Check pre-conditions:
mod1 <- lm(policy ~ polAffil, data = dat1)
mod2 <- lm(policy ~ sysRac, data = dat1)
mod3 <- lm(sysRac ~ polAffil, data = dat1)

## Partial out the mediator's effect:
mod4 <- lm(policy ~ sysRac + polAffil, data = dat1)
```



# Causal Steps Example

```
summary(mod1)
```

Call:

```
lm(formula = policy ~ polAffil, data = dat1)
```

Residuals:

Min	1Q	Median	3Q	Max
-2.7357	-0.8254	0.0643	0.6827	3.2481

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	2.71516	0.35648	7.617	3.32e-11	***
polAffil	0.23675	0.07775	3.045	0.0031	**
---					

Signif. codes:

0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 1.134 on 85 degrees of freedom

Multiple R-squared: 0.09836, Adjusted R-squared: 0.08775

F-statistic: 9.273 on 1 and 85 DF, p-value: 0.003096

# Causal Steps Example

```
summary(mod2)
```

Call:

```
lm(formula = policy ~ sysRac, data = dat1)
```

Residuals:

Min	1Q	Median	3Q	Max
-1.7700	-0.5593	0.0255	0.6277	3.6835

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	0.9295	0.3896	2.386	0.0193 *
sysRac	0.7557	0.1014	7.450	7.14e-11 ***

---

Signif. codes:

0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.9286 on 85 degrees of freedom

Multiple R-squared: 0.395, Adjusted R-squared: 0.3879

F-statistic: 55.5 on 1 and 85 DF, p-value: 7.145e-11

# Causal Steps Example

```
summary(mod3)
```

Call:

```
lm(formula = sysRac ~ polAffil, data = dat1)
```

Residuals:

Min	1Q	Median	3Q	Max
-2.44714	-0.50502	0.05286	0.54498	2.25286

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	2.60605	0.28489	9.147	2.72e-14 ***
polAffil	0.25685	0.06213	4.134	8.34e-05 ***

---

Signif. codes:

0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.906 on 85 degrees of freedom

Multiple R-squared: 0.1674, Adjusted R-squared: 0.1576

F-statistic: 17.09 on 1 and 85 DF, p-value: 8.336e-05

# Causal Steps Example

```
summary(mod4)
```

Call:

```
lm(formula = policy ~ sysRac + polAffil, data = dat1)
```

Residuals:

Min	1Q	Median	3Q	Max
-1.7156	-0.6043	0.0262	0.6474	3.7992

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	0.83266	0.41246	2.019	0.0467 *
sysRac	0.72236	0.11148	6.480	5.93e-09 ***
polAffil	0.05121	0.06998	0.732	0.4663

---

Signif. codes:

0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.9312 on 84 degrees of freedom

Multiple R-squared: 0.3989, Adjusted R-squared: 0.3845

F-statistic: 27.87 on 2 and 84 DF, p-value: 5.211e-10



# Causal Steps Example

---

```
## Extract important parameter estimates:
```

```
a      <- coef(mod3) ["polAffil"]
```

```
b      <- coef(mod4) ["sysRac"]
```

```
c      <- coef(mod1) ["polAffil"]
```

```
cPrime <- coef(mod4) ["polAffil"]
```

```
## Compute indirect effects:
```

```
ieDiff <- unname(c - cPrime)
```

```
ieProd <- unname(a * b)
```

```
ieDiff
```

```
[1] 0.1855374
```

```
ieProd
```

```
[1] 0.1855374
```

# Sobel's Z

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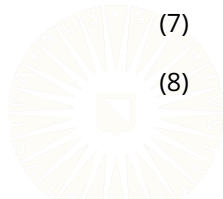
In the previous example, do we have a *significant* indirect effect?

- The direct effect is “substantially” smaller than the total effect, but is the difference statistically significant?
- Sobel (1982) developed an asymptotic standard error for  $IE_{prod}$  that we can use to assess this hypothesis.

$$SE_{sobel} = \sqrt{a^2 \cdot SE_b^2 + b^2 \cdot SE_a^2} \quad (6)$$

$$Z_{sobel} = \frac{ab}{SE_{sobel}} \quad (7)$$

$$95\%CI_{sobel} = ab \pm 1.96 \cdot SE_{sobel} \quad (8)$$



# Sobel Example

```
## SE:
seA <- (mod3 %>% vcov() %>% diag() %>% sqrt())["polAffil"]
seB <- (mod4 %>% vcov() %>% diag() %>% sqrt())["sysRac"]

se <- sqrt(b^2 * seA^2 + a^2 * seB^2) %>% unname()

## z-score:
(z <- ieProd / se)

[1] 3.48501

## p-value:
(p <- 2 * pnorm(z, lower = FALSE))

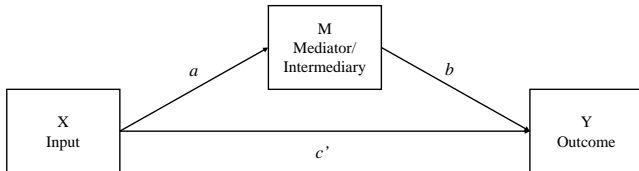
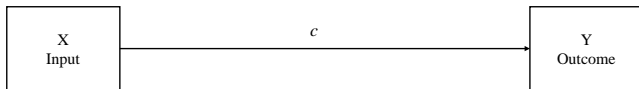
[1] 0.0004921178

## 95% CI:
c(ieProd - 1.96 * se, ieProd + 1.96 * se)

[1] 0.08118957 0.28988525
```

# Recall our Basic Path Diagram

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# Two Measures of Indirect Effect

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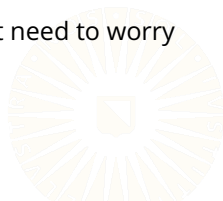
Recall the two definitions of an indirect effect:

$$IE_{diff} = c - c' \quad (9)$$

$$IE_{prod} = a \cdot b \quad (10)$$

It pays to remember a few key points:

- $IE_{diff}$  and  $IE_{prod}$  are equivalent in simple mediation.
- $IE_{diff}$  is only an indirect indication of  $IE_{prod}$ .
- A significant indirect effect can exist without a significant total effect.
- If we only care about the indirect effect, then we don't need to worry about the total effect.



# Two Measures of Indirect Effect

---

Recall the two definitions of an indirect effect:

$$IE_{diff} = c - c' \quad (9)$$

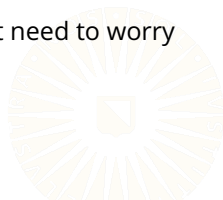
$$IE_{prod} = a \cdot b \quad (10)$$

It pays to remember a few key points:

- $IE_{diff}$  and  $IE_{prod}$  are equivalent in simple mediation.
- $IE_{diff}$  is only an indirect indication of  $IE_{prod}$ .
- A significant indirect effect can exist without a significant total effect.
- If we only care about the indirect effect, then we don't need to worry about the total effect.

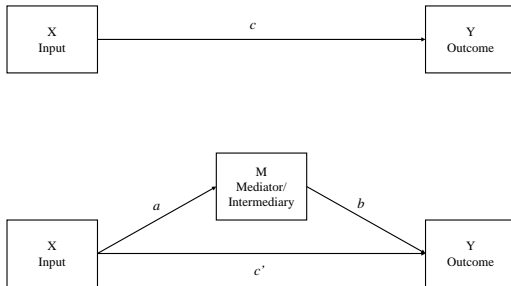
These points imply something interesting:

- We don't need to estimate  $c$ !



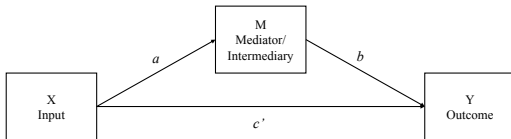
# Simplifying our Path Diagram

Question: If we don't care about directly estimating  $c$ , how can we simplify this diagram?



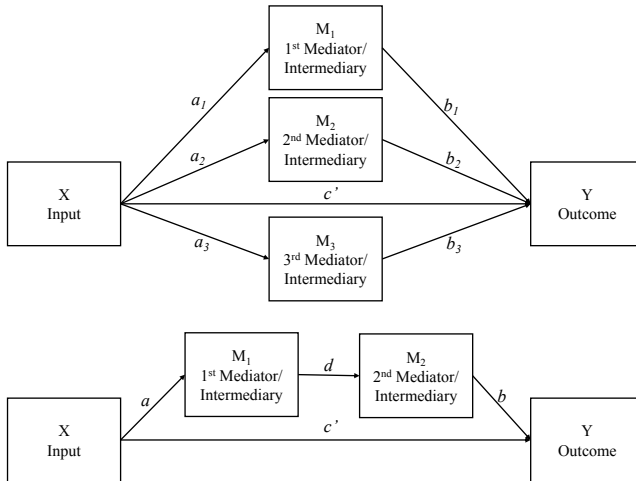
# Simplifying our Path Diagram

Answer: We don't fit the upper model.





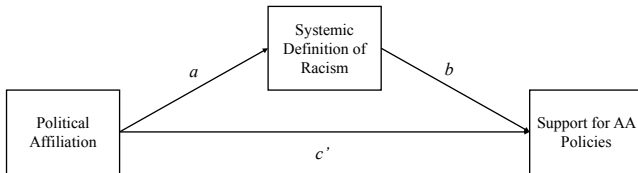
# Why Path Analysis?



# Example

---

Let's revisit the above example using path analysis in **lavaan**.



# Example

---

```
## Load the lavaan package:  
library(lavaan)  
  
## Specify the basic path model:  
mod1 <- "  
policy ~ sysRac + polAffil  
sysRac ~ polAffil  
"  
  
## Estimate the model:  
out1 <- sem(mod1, data = dat1)
```



# Example

```
## Look at the results:  
partSummary(out1, c(5, 6))
```

## Regressions:

	Estimate	Std.Err	z-value	P(> z )
policy ~				
sysRac	0.722	0.110	6.595	0.000
polAffil	0.051	0.069	0.745	0.456
sysRac ~				
polAffil	0.257	0.061	4.182	0.000

## Variances:

	Estimate	Std.Err	z-value	P(> z )
.policy	0.837	0.127	6.595	0.000
.sysRac	0.802	0.122	6.595	0.000

# Example

---

```
## Include the indirect effect:
mod2 <- "
policy ~ b*sysRac + polAffil
sysRac ~ a*polAffil

ab := a*b # Define a parameter for the indirect effect
"

## Estimate the model:
out2 <- sem(mod2, data = dat1)
```



# Example

```
## Look at the results:
```

```
partSummary(out2, 5:7)
```

Regressions:

		Estimate	Std.Err	z-value	P(> z )
policy ~					
sysRac	(b)	0.722	0.110	6.595	0.000
polAffil		0.051	0.069	0.745	0.456
sysRac ~					
polAffil	(a)	0.257	0.061	4.182	0.000

Variances:

	Estimate	Std.Err	z-value	P(> z )
.policy	0.837	0.127	6.595	0.000
.sysRac	0.802	0.122	6.595	0.000

Defined Parameters:

	Estimate	Std.Err	z-value	P(> z )
ab	0.186	0.053	3.532	0.000

# Example

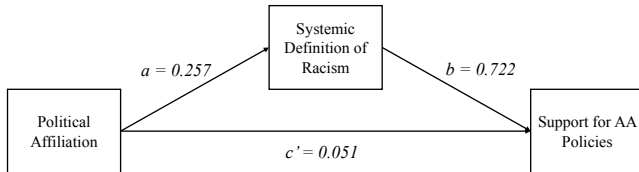
*## We can also get CIs:*

```
parameterEstimates(out2, zstat = FALSE, pvalue = FALSE, ci = TRUE)
```

	lhs	op	rhs	label	est	se	ci.lower	ci.upper
1	policy	~	sysRac	b	0.722	0.110	0.508	0.937
2	policy	~	polAffil		0.051	0.069	-0.084	0.186
3	sysRac	~	polAffil	a	0.257	0.061	0.136	0.377
4	policy	~~	policy		0.837	0.127	0.588	1.086
5	sysRac	~~	sysRac		0.802	0.122	0.564	1.040
6	polAffil	~~	polAffil		2.444	0.000	2.444	2.444
7	ab	:=	a*b	ab	0.186	0.053	0.083	0.289

# Results

---





# We're not there yet...

---

Path analysis allows us to directly model complex (and simple) relations, but the preceding example still suffers from a considerable limitation.

- The significance test for the indirect effect is still conducted with the Sobel Z approach.

Path analysis (or full SEM) doesn't magically get around distributional problems associated with Sobel's Z test.

- To get a robust significance test of the indirect effect, we need to use *bootstrapping*.



# Bootstrapping

---

Bootstrapping was introduced by Efron (1979) as a tool for non-parametric inference.

- Traditional inference requires that we assume a parametric sampling distribution for our focal parameter.
- We need to make such an assumption to compute the standard errors we require for inferences.
- If we cannot safely make these assumptions, we can use bootstrapping.



# Bootstrapping

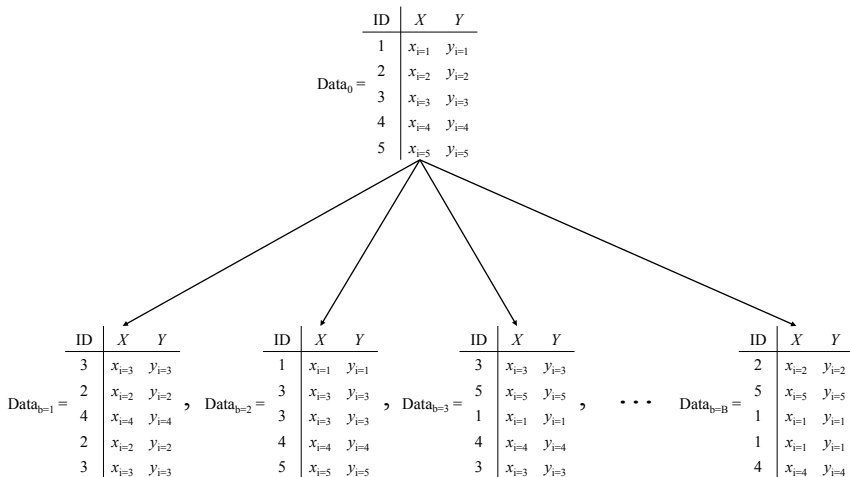
---

Assume our observed data  $Data_0$  represent the population and:

1. Sample rows of  $Data_0$ , with replacement, to create  $B$  new samples  $\{Data_b\}$ .
2. Calculate our focal statistic on each of the  $B$  bootstrap samples.
3. Make inferences based on the empirical distribution of the  $B$  estimates calculated in Step 2



# Bootstrapping



# Example

---

Suppose I'm on the lookout for a retirement location. Since I want to relax in my old-age, I'm concerned with ensuring a low probability of dragon attacks, so I have a few salient considerations:

- Shooting for a location with no dragons, whatsoever, is a fools errand (since dragons are, obviously, ubiquitous).
- I merely require a location that has at least two times as many dragon-free days as other kinds.



# Example

---

I've been watching several candidate locales over the course of my (long and illustrious) career, and I'm particularly hopeful about one quiet hamlet in the Patagonian highlands.

- To ensure that my required degree of dragon-freeness is met, I'll use the *Dragon Risk Index* (DRI):

$$DRI = \text{Median} \left( \frac{\text{Dragon-Free Days}}{\text{Dragonned Days}} \right)$$



# Example

---

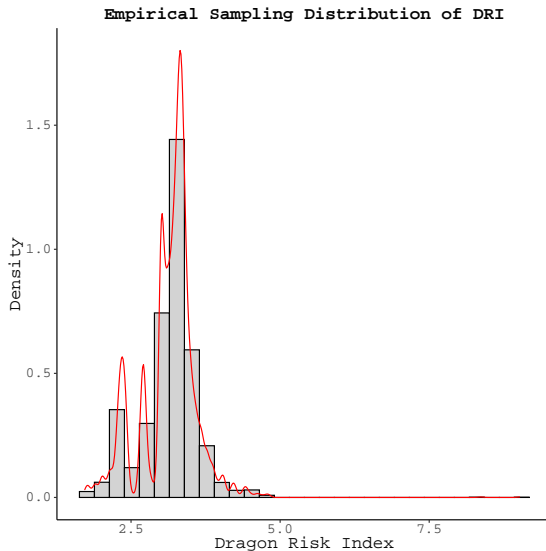
```
## Read in the observed data:
rawData <- readRDS("../data/daysData.rds")

## Compute the observed test statistic:
obsDRI <- median(rawData$goodDays / rawData$badDays)
obsDRI

[1] 3.24476

## Draw the bootstrap samples:
set.seed(235711)
nSams <- 5000
bootDRI <- rep(NA, nSams)
for(b in 1:nSams) {
  bootSam <- rawData[sample(1:nrow(rawData), replace = TRUE), ]
  bootDRI[b] <- median(bootSam$goodDays / bootSam$badDays)
}
```

# Example





# Example

---

To see if I can be confident in the dragon-freeness of my potential home, I'll summarize the preceding distribution with a (one-tailed) percentile confidence interval:

```
bootLB <- sort(bootDRI)[0.05 * nSams]
bootUB <- Inf

## The bootstrapped Percentile CI:
c(bootLB, bootUB)

[1] 2.288555      Inf
```



# Bootstrapped Inference for Indirect Effects

---

We can apply the same procedure to testing the indirect effect.

- The problem with Sobel's Z is exactly the type of issue for which bootstrapping was designed
  - We don't know a reasonable finite-sample sampling distribution for the *ab* parameter.
- Bootstrapping will allow us to construct an empirical sampling distribution for *ab* and construct confidence intervals for inference.



# Bootstrapped Inference for Indirect Effects

---

## PROCEDURE:

1. Resample our observed data with replacement
2. Fit our hypothesized path model to each bootstrap sample
3. Store the value of  $ab$  that we get each time
4. Summarize the empirical distribution of  $ab$  to make inferences



# Example

---

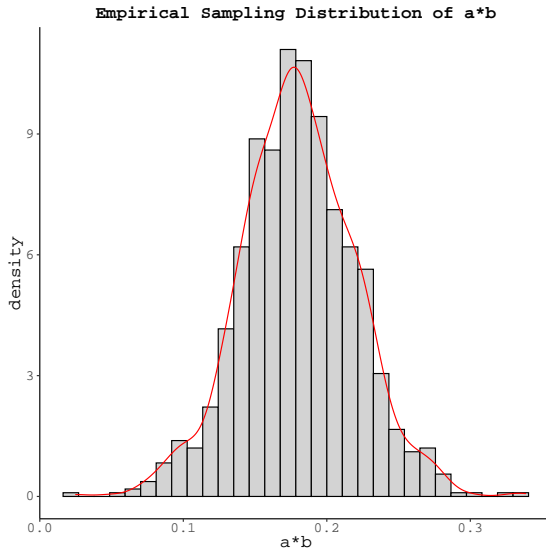
```
nSams <- 1000
abVec <- rep(NA, nSams)
for(i in 1:nSams) {
  ## Resample the data:
  bootSam <- dat1[sample(1:nrow(dat1), replace = TRUE), ]

  ## Fit the path model:
  bootOut <- sem(mod2, data = bootSam)

  ## Store the estimated indirect effect:
  abVec[i] <- coef(bootOut)[c("a", "b")] %>% prod()
}
```



# Example



# Example

---

```
## Calculate the percentile CI:  
lb <- sort(abVec)[0.025 * nSams]  
ub <- sort(abVec)[0.975 * nSams]  
c(lb, ub)  
  
[1] 0.09804422 0.26334737
```



# Example

```
## Much more parsimoniously:
```

```
bootOut2 <- sem(mod2, data = dat1, se = "boot", bootstrap = 1000)
```

```
parameterEstimates(bootOut2, zstat = FALSE, pvalue = FALSE)
```

	lhs	op	rhs	label	est	se	ci.lower	ci.upper
1	policy	~	sysRac	b	0.722	0.105	0.519	0.926
2	policy	~	polAffil		0.051	0.078	-0.095	0.209
3	sysRac	~	polAffil	a	0.257	0.060	0.129	0.375
4	policy	~~	policy		0.837	0.163	0.530	1.158
5	sysRac	~~	sysRac		0.802	0.128	0.557	1.063
6	polAffil	~~	polAffil		2.444	0.000	2.444	2.444
7	ab	:=	a*b	ab	0.186	0.041	0.102	0.265

# Monte Carlo Method/Parametric Bootstrap

We can also use a *parametric bootstrap* of the individual parameters  $a$  and  $b$  to get a somewhat robust test of the indirect effect.

- Assuming normal sampling distributions for  $a$  and  $b$  is not, generally, problematic
- We can save ourselves a lot of computational effort by assuming normality for  $a$  and  $b$ , then:
  1. Fit the hypothesized path model to the raw data
  2. Extract  $a$ ,  $b$ , and  $ACOV(\{a, b\})$  from the fitted path model
  3. Parameterize a bivariate normal distribution  $N(a, b | \mu, \Sigma)$  with  $\mu = \{a, b\}$  and  $\Sigma = ACOV(\{a, b\})$
  4. Draw simulated values  $\{\tilde{a}, \tilde{b}\}$  from  $N(a, b | \mu, \Sigma)$
  5. Compute the simulated indirect effect  $\tilde{ab} = \tilde{a} \cdot \tilde{b}$  and store it
  6. Summarize the empirical distribution of  $\tilde{ab}$  for inference.





# Example

---

```
## Load package to draw the Monte Carlo samples
library(mvtnorm)

## Specify the model (note the parameter labels):
mod3 <- "
policy ~ polAffil + b*sysRac
sysRac ~ a*polAffil
"

## Fit the model:
out4 <- sem(mod3, data = dat1)

## Extract the important estimates:
a <- coef(out4)["a"]
b <- coef(out4)["b"]
acov <- vcov(out4)[c("a", "b"), c("a", "b")]
```

## Aside Regarding Asymptotic Covariances

The *asymptotic covariance matrix* (ACOV) is (-1 times) the inverse of the Fisher information matrix of the model parameters.

- The *ACOV* contains the expected covariance among the ML estimates of the model parameters.
- The diagonal elements of the matrix (i.e., the asymptotic variances) are the square of the usual ML SE estimates.

```
round(vcov(out4), 4)
```

	plcy~A b	a	plcy~~	syR~~R	
policy~polAffil	0.005				
b	-0.003	0.012			
a	0.000	0.000	0.004		
policy~~policy	0.000	0.000	0.000	0.016	
sysRac~~sysRac	0.000	0.000	0.000	0.000	0.015

# Back to the Example

---

```
## Look at ACOV(a,b):
```

```
round(acov, 4)
```

```
      a      b
```

```
a 0.0038 0.000
```

```
b 0.0000 0.012
```

```
## Draw the Monte Carlo samples:
```

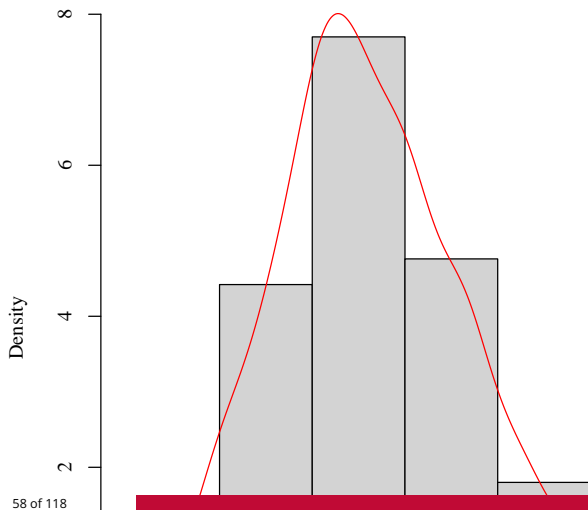
```
samMat <- rmvnorm(nSams, c(a, b), acov)
```

```
abVec <- samMat[, "a"] * samMat[, "b"]
```



# Back to the Example

Monte Carlo Distribution of  $a*b$



# Kris Preacher's Website

---

If you don't want to program the Monte Carlo approach yourself (although each of you easily can), you should consider the very handy Web App on Professor Kris Preacher's website

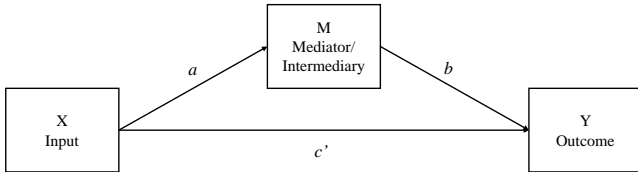
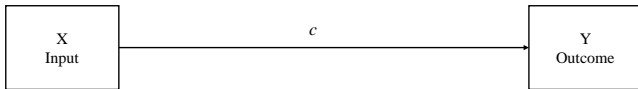
<http://www.quantpsy.org>.

- Kris' website has a vast array of hugely helpful resources for anyone doing mediation or moderation analysis.
- You should definitely check it out!



# Simple Mediation

---



# Simple Mediation is Too Simple

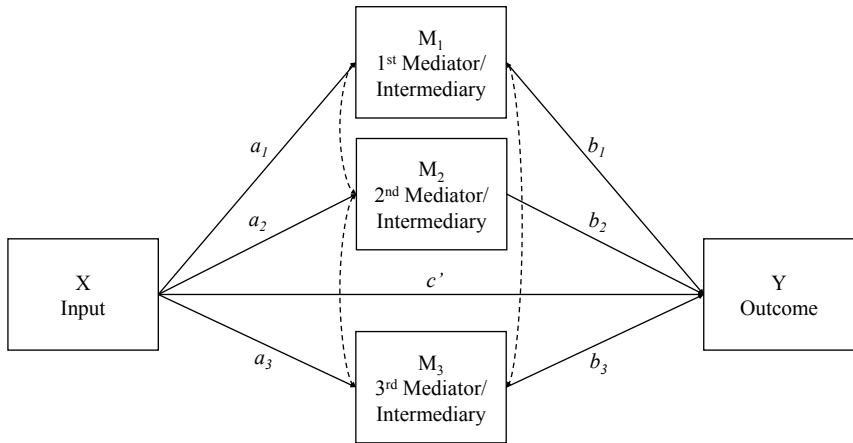
---

We can justify multiple mediator models by asking: “What mediates the effects in a simple mediation model?”

- Mediation of the direct effect leads to *parallel multiple mediator models*.
- Mediation of the *a* or *b* paths produces *serial multiple mediator models*.



# Parallel Multiple Mediation





# Parallel Multiple Mediation

---

To get all of the information in the preceding diagram, we need to fit four equations:

$$\begin{aligned}Y &= i_Y + b_1M_1 + b_2M_2 + b_3M_3 + c'X + e_Y \\M_1 &= i_{M1} + a_1X + e_{M1} \\M_2 &= i_{M2} + a_2X + e_{M2} \\M_3 &= i_{M3} + a_3X + e_{M3}\end{aligned}$$

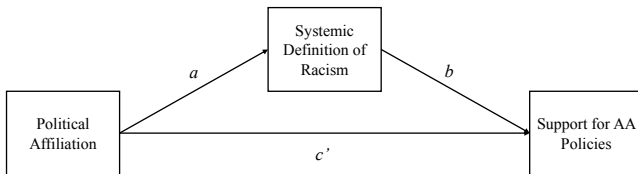
In general, a parallel mediator model with  $K$  mediator variables will require  $K + 1$  separate equations.

Path modeling can make this task much simpler.

- Also allows us to explicitly estimate the correlations between parallel mediators.

# Parallel Multiple Mediation

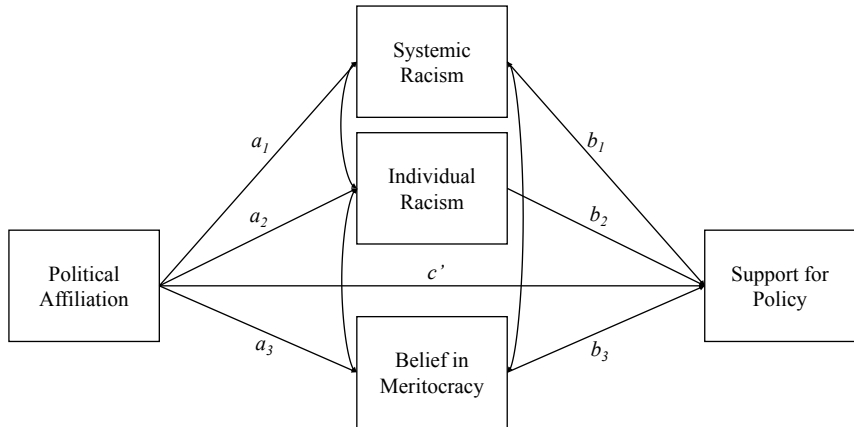
Let's reconsider the example from last week:



Question: What might be mediating the residual direct effect?

# Parallel Multiple Mediation

Potential Answer:



# A Quick Note on Inference

---

In simple mediation:

- We have one indirect effect:  $ab$ .
- The total effect is equal to the direct effect plus the indirect effect  
 $c = c' + ab$

In parallel multiple mediation:

- We have  $K$  *specific indirect effects*, where  $K$  is the number of mediators:  
 $a_1b_1, a_2b_2, \dots, a_Kb_K$ .
- The *Total Indirect Effect* is equal to the sum of all the specific indirect effects:  $IE_{tot} = \sum_{k=1}^K a_kb_k$ .
- The *Total Effect* is equal to the direct effect plus the total indirect effect:  
 $c = c' + IE_{tot}$

Inference for the specific indirect effects is basically the same as it is for the sole indirect effect in simple mediation.

- Caveat: Each specific indirect effect must be interpreted as conditional on all other mediators in the model.

# Example

---



# Example

```
library(lavaan)

## Read in the data
dataDir <- "../data/"
fileName <- "adamsKlpsScaleScore.rds"
dat1 <- readRDS(paste0(dataDir, fileName))

nBoot <- 2500 # Number of bootstrap samples
bootType <- "bca.simple" # Type of CI

## Parallel Multiple Mediator Model:
mod1.1 <- "
policy ~ b1*sysRac + b2*indRac + b3*merit + cp*polAffil
sysRac ~ a1*polAffil
indRac ~ a2*polAffil
merit ~ a3*polAffil

sysRac ~~ indRac + merit
indRac ~~ merit

ab1 := a1*b1
ab2 := a2*b2
ab3 := a3*b3
```

# Example

---

```
parameterEstimates(out1.1, boot.ci.type = bootType)[ , -c(1 : 3)]
```

```
Error in parameterEstimates(out1.1, boot.ci.type = bootType): object 'out1.1'  
not found
```



# Comparing Specific Indirect Effects

---

When we have multiple specific indirect effects in a single model, we can test if they are statistically different from one another.

Question: How might we go about doing such a test (assuming we're using path modeling)?





# Comparing Specific Indirect Effects

---

When we have multiple specific indirect effects in a single model, we can test if they are statistically different from one another.

Question: How might we go about doing such a test (assuming we're using path modeling)?

Answer: There are, at least, two reasonable methods:

1. Use nested model  $\Delta\chi^2$  tests
2. Define a new parameter corresponding to the null hypothesis and use bootstrapping



# Example

---



# Example

```
## Test differences in specific indirect effects:
```

```
mod1.2 <- "
```

```
policy ~ b1*sysRac + b2*indRac + b3*merit + cp*polAffil
```

```
sysRac ~ a1*polAffil
```

```
indRac ~ a2*polAffil
```

```
merit ~ a3*polAffil
```

```
sysRac ~~ indRac + merit
```

```
indRac ~~ merit
```

```
ab1 := a1*b1
```

```
ab2 := a2*b2
```

```
ab3 := a3*b3
```

```
totalIE := ab1 + ab2 + ab3
```

```
ab1 == ab2
```

```
"
```

```
out1.2 <-
```

```
sem(mod1.2, data = dat1, se = "boot", bootstrap = nBoot)
```

```
Error in lavaan::lavaan(model = mod1.2, data = dat1, se = "boot", bootstrap
```

```
= nBoot, : lower FPPOR: missing observed variables in dataset: merit
```

# Example

---



# Example

```
## Test differences in specific indirect effects:
```

```
mod1.3 <- "
```

```
policy ~ b1*sysRac + b2*indRac + b3*merit + cp*polAffil
```

```
sysRac ~ a1*polAffil
```

```
indRac ~ a2*polAffil
```

```
merit ~ a3*polAffil
```

```
sysRac ~~ indRac + merit
```

```
indRac ~~ merit
```

```
ab1 := a1*b1
```

```
ab2 := a2*b2
```

```
ab3 := a3*b3
```

```
totalIE := ab1 + ab2 + ab3
```

```
test1 := ab2 - ab1
```

```
"
```

```
out1.3 <-
```

```
sem(mod1.3, data = dat1, se = "boot", bootstrap = nBoot)
```

```
Error in lavaan::lavaan(model = mod1.3, data = dat1, se = "boot", bootstrap
```

```
= nBoot, : lavaan FPPOR: missing observed variables in dataset: merit
```

# Example

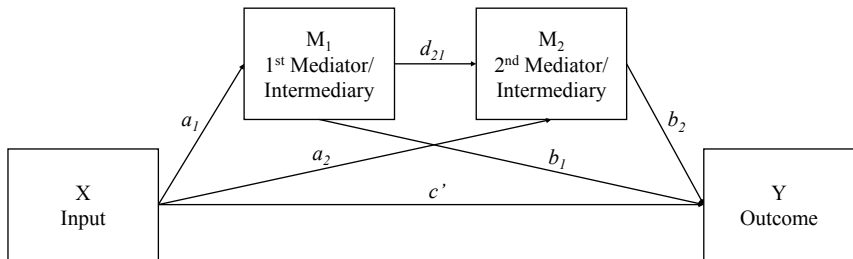
---

```
parameterEstimates(out1.3, boot.ci.type = bootType)[ , -c(1 : 3)]
```

```
Error in parameterEstimates(out1.3, boot.ci.type = bootType): object 'out1.3'  
not found
```



# Serial Multiple Mediation



# Serial Multiple Mediation

---

To get all of the information in the preceding diagram, we need to fit three equations:

$$\begin{aligned}Y &= i_Y + b_1M_1 + b_2M_2 + c'X + e_Y \\M_2 &= i_{M2} + d_{21}M_1 + a_2X + e_{M2} \\M_1 &= i_{M1} + a_1X + e_{M1}\end{aligned}$$

As with parallel mediator models, a serial mediator model with  $K$  mediator variables will required  $K + 1$  separate equations.

Again, path modeling can make this task much simpler.

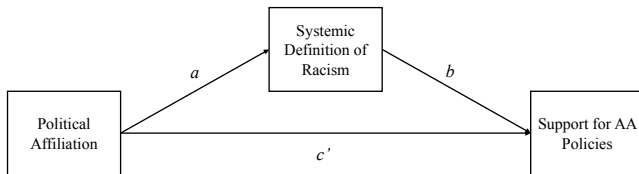
- Also allows us to fit more parsimonious, restricted models.



# Serial Multiple Mediation

---

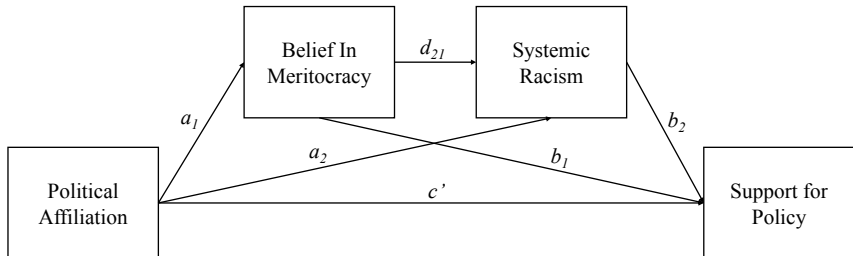
Okay, back to our simple mediation example:



Question: What could be mediating the  $a$  path?

# Serial Multiple Mediation

Potential Answer:



# A Quick Note on Inference

---

Parallel multiple mediation operates much like a number of combined simple mediation models, serial multiple mediation is not so straight-forward.

In serial multiple mediation:

- Every possible path from  $X$  to  $Y$  that passes through, at least, one mediator is a specific indirect effect.
  - With the saturated two-mediator model shown above, we have:
$$IE_{spec} = \{a_1b_1, a_2b_2, a_1d_{21}b_2\}$$
- The *Total Indirect Effect* is, again, equal to the sum of all the specific indirect effects:  $IE_{tot} = \sum_{k=1}^{|\{IE_{spec}\}|} IE_{spec,k}$ .
- The *Total Effect* is equal to the direct effect plus the total indirect effect:  $c = c' + IE_{tot}$

# A Quick Note on Inference

---

Inference for the specific indirect effects is basically the same as it is for the sole indirect effect in simple mediation, when using path modeling or bootstrapping.

- Caveat: Normal-theory, Sobel-Type, standard errors for the specific indirect effects that involve more than two constituent paths can be very complex.
  - This isn't really a problem since you should always use bootstrapping, anyway!



# Example

---



# Example

```
## Serial Multiple Mediator Model:
```

```
mod2.1 <- "  
policy ~ b1*merit + b2*sysRac + cp*polAffil  
sysRac ~ d21*merit + a2*polAffil  
merit ~ a1*polAffil
```

```
ab1 := a1*b1  
ab2 := a2*b2  
fullIE := a1*d21*b2  
totalIE := ab1 + ab2 + fullIE  
"
```

```
out2.1 <-  
  sem(mod2.1, data = dat1, se = "boot", bootstrap = nBoot)
```

```
Error in lavaan::lavaan(model = mod2.1, data = dat1, se = "boot", bootstrap  
= nBoot, : lavaan ERROR: missing observed variables in dataset: merit
```

```
summary(out2.1)
```

```
Error in h(simpleError(msg, call)): error in evaluating the argument  
'object' in selecting a method for function 'summary': object 'out2.1' not  
found
```

# Example

---

```
parameterEstimates(out2.1, boot.ci.type = bootType)[ , -c(1 : 3)]
```

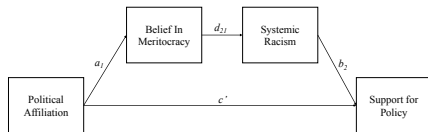
```
Error in parameterEstimates(out2.1, boot.ci.type = bootType): object 'out2.1'  
not found
```



# Restricted Models

In this example, the  $a_2$  and  $b_1$  paths are non-significant as are the simple specific indirect effects  $a_1b_1$  and  $a_2b_2$ .

- There is a school of thinking that would prescribe constraining the  $a_2$  and  $b_1$  paths to zero as in:



- This model will ascribe a larger effect size to  $a_1d_{21}b_2$  since it must convey all of the indirect influence of  $X$  on  $Y$ .
  - We should first fit a saturated model, but subsequently culling non-significant paths can, sometimes, be appropriate.



# Example

---

```
mod2.2 <- "  
policy ~ cp*polAffil + b2*sysRac  
merit ~ a1*polAffil  
sysRac ~ d21*merit  
  
fullIE := a1*d21*b2  
"  
  
out2.2 <-  
  sem(mod2.2, data = dat1, se = "boot", bootstrap = nBoot)  
  
Error in lavaan::lavaan(model = mod2.2, data = dat1, se = "boot", bootstrap  
= nBoot, : lavaan ERROR: missing observed variables in dataset: merit  
  
summary(out2.2)  
  
Error in h(simpleError(msg, call)): error in evaluating the argument  
'object' in selecting a method for function 'summary': object 'out2.2' not  
found
```

# Example

---

```
parameterEstimates(out2.2, bootstrap = bootType)[ , -c(1 : 3)]
```

```
Error in parameterEstimates(out2.2, bootstrap = bootType): unused argument  
(bootstrap = bootType)
```



# Example

---

As in parallel multiple mediation, we can test for differences in the specific indirect effects of a serial multiple mediator model:



# Example

```
## Test Differences between Indirect Effects
## in Serial Multiple Mediator Model (Method 1):
mod2.3 <- "
policy ~ cp*polAffil + b1*merit + b2*sysRac
merit ~ a1*polAffil
sysRac ~ a2*polAffil + d21*merit

ab1 := a1*b1
ab2 := a2*b2
fullIE := a1*d21*b2
totalIE := ab1 + ab2 + fullIE

fullIE == ab1
fullIE == ab2
"

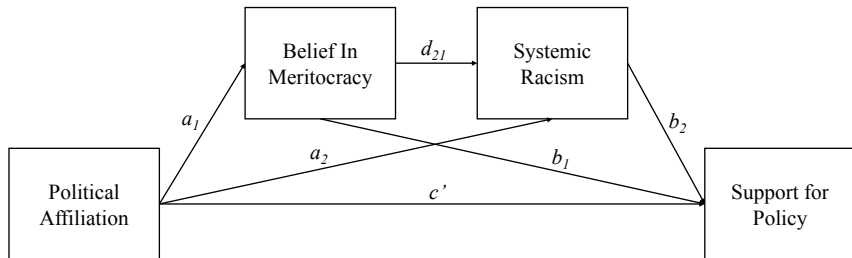
out2.3 <-
  sem(mod2.3, data = dat1, se = "boot", bootstrap = nBoot)

Error in lavaan::lavaan(model = mod2.3, data = dat1, se = "boot", bootstrap
= nBoot, : lavaan ERROR: missing observed variables in dataset: merit

summary(out2.3)
```

# Serial Multiple Mediation

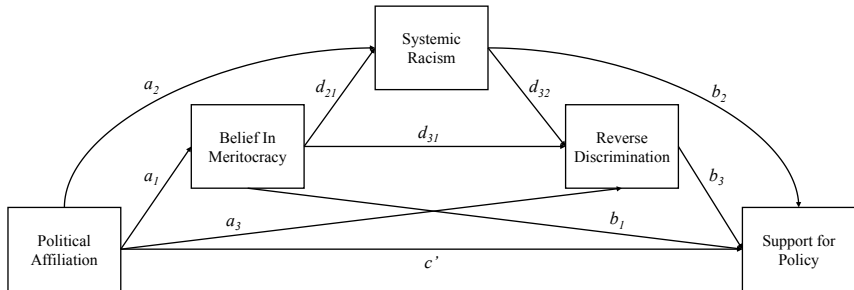
Okay, so we've supported an interesting hypothesis with the following model but why stop there?



Question: What might mediated the  $b_2$  path?

# Serial Multiple Mediation

Potential Answer:



# Serial Multiple Mediation

---

Question: How many equations do we need to get the information in the preceding diagram?



# Serial Multiple Mediation

Question: How many equations do we need to get the information in the preceding diagram?

$$\text{Policy} = i_Y + b_1\text{Merit} + b_2\text{SysRac} + b_3\text{RevDisc} + c'\text{PolAff} + e_Y$$

$$\text{RevDisc} = i_{M3} + d_{31}\text{Merit} + d_{32}\text{SysRac} + a_3\text{PolAff} + e_{M3}$$

$$\text{SysRac} = i_{M2} + d_{21}\text{Merit} + a_2\text{PolAff} + e_{M2}$$

$$\text{Merit} = i_{M1} + a_1\text{PolAff} + e_{M1}$$

Which produces the following set of specific indirect effects:

- $a_1b_1$
- $a_2b_2$
- $a_3b_3$
- $a_1d_{31}b_3$
- $a_1d_{21}b_2$
- $a_2d_{32}b_3$
- $a_1d_{21}d_{32}b_3$



# Example

---



# Example

```
## Serial Multiple Mediator Model with 3 Mediators:
mod3.1 <- "
policy ~ b1*merit + b2*sysRac + b3*revDisc + cp*polAffil
revDisc ~ d31*merit + d32*sysRac + a3*polAffil
sysRac ~ d21*merit + a2*polAffil
merit ~ a1*polAffil

ab1 := a1*b1
ab2 := a2*b2
ab3 := a3*b3

partIE1 := a1*d31*b3
partIE2 := a1*d21*b2
partIE3 := a2*d32*b3

fullIE := a1*d21*d32*b3

totalIE := ab1 + ab2 + ab3 + partIE1 + partIE2 + partIE3 + fullIE
"

out3.1 <-
  sem(mod3.1, data = dat1, se = "boot", bootstrap = nBoot)
```

# Example

---

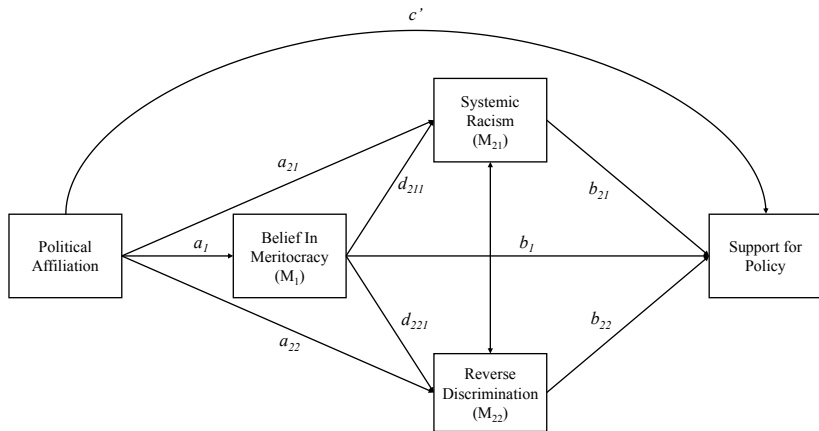
```
parameterEstimates(out3.1, boot.ci.type = bootType)[ , -c(1 : 3)]
```

```
Error in parameterEstimates(out3.1, boot.ci.type = bootType): object 'out3.1'  
not found
```



# Hybrid Multiple Mediation

We can also combine parallel and serial mediation models:



# Example

---



# Example

```
## Hybrid Multiple Mediator Model:
```

```
mod4.1 <- "
```

```
policy ~ b1*merit + b21*sysRac + b22*revDisc + cp*polAffil
```

```
sysRac ~ d211*merit + a21*polAffil
```

```
revDisc ~ d221*merit + a22*polAffil
```

```
merit ~ a1*polAffil
```

```
sysRac ~~ revDisc
```

```
ab1 := a1*b1
```

```
ab21 := a21*b21
```

```
ab22 := a22*b22
```

```
fullIE21 := a1*d211*b21
```

```
fullIE22 := a1*d221*b22
```

```
totalIE := ab1 + ab21 + ab22 + fullIE21 + fullIE22
```

```
"
```

```
out4.1 <-
```

```
sem(mod4.1, data = dat1, se = "boot", bootstrap = nBoot)
```

```
Error in lava::lavsem(model = mod4.1, data = dat1, se = "boot", bootstrap = nBoot)
```

# Example

---

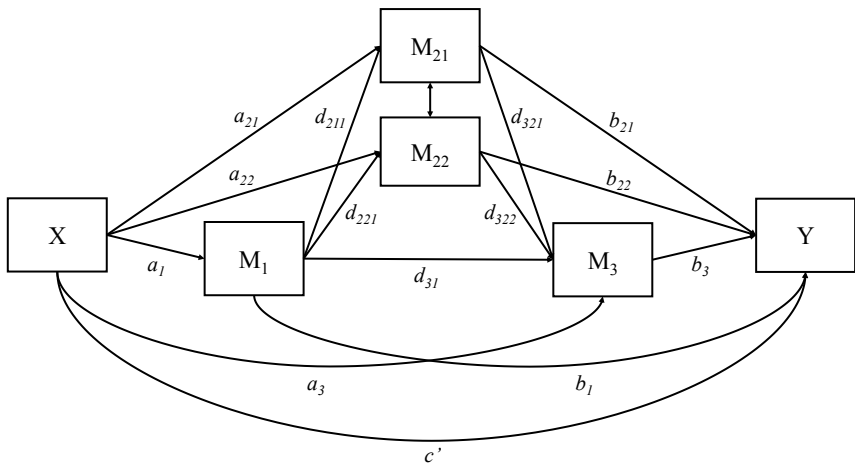
```
parameterEstimates(out4.1, boot.ci.type = bootType)[ , -c(1 : 3)]
```

```
Error in parameterEstimates(out4.1, boot.ci.type = bootType): object 'out4.1'  
not found
```



# Practice

List all of the specific indirect effects present in this model:

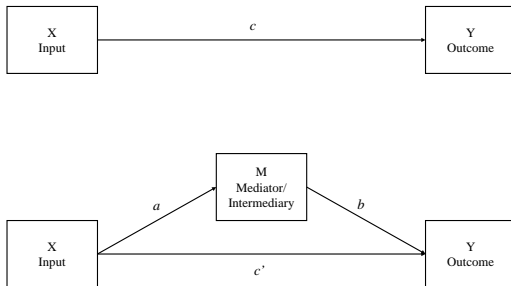




# Boring Model

---

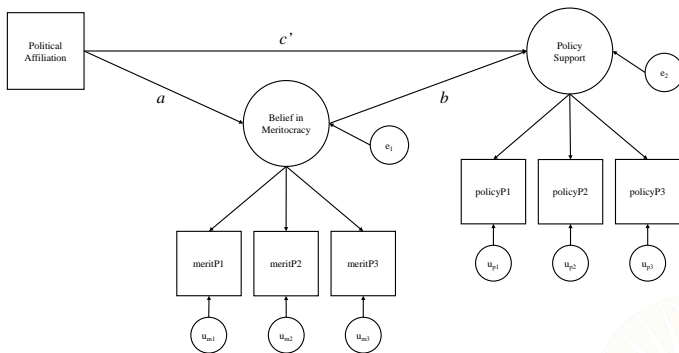
So far, all of our models have been similar to:



But there is no reason that we need to restrict ourselves to mucking about with observed variables.

# Better Model

We can (and should) test for indirect effects using *latent variable models* such as:



Measurement error can be a big problem for mediation analysis, so latent variable modeling is highly recommended.

# Example

---



# Example

```
library(lavaan)
dataDir <- "../data/"
dat1 <- readRDS(paste0(dataDir, "adamsKlpsData.rds"))

## Specify the CFA model:
mod1.1 <- "
merit =~ meritP1 + meritP2 + meritP3
policy =~ policyP1 + policyP2 + policyP3
"

## Fit the CFA and check model:
out1.1 <- cfa(mod1.1, data = dat1, std.lv = TRUE)

Error in lavaan::lavaan(model = mod1.1, data = dat1, std.lv = TRUE,
model.type = "cfa", : lavaan ERROR: some latent variable names collide
with observed
variable names:  merit policy

## Check model fit:
round(fitMeasures(out1.1)[c("chisq", "df", "pvalue", "cfi",
                           "tli", "rmsea", "srmr")], 4)

Error in h(simpleError(msg, call)): error in evaluating the argument
'object' in
```

# Interpretation of Indirect Effects

---

Although indirect effects are composed parameters, they have direct interpretations, independent of the interpretations of their constituent paths:

- The  $X \rightarrow M \rightarrow Y$  indirect effect  $ab$  is interpreted as:
  - The expected change in  $Y$  for a unit change in  $X$  that is transmitted indirectly through  $M$ , or...
  - For a unit change in  $X$ ,  $Y$  is expected to change by  $ab$  units, indirectly through  $M$ , or...
  - Participants who differ by one unit on  $X$  are expected to differ by  $ab$  units on  $Y$  as a result of the effect of  $X$  on  $M$  which, in turn, affects  $Y$ .
- The interpretation/scaling of the indirect effect is entirely defined by the input  $X$  and outcome  $Y$ 
  - The scaling of the intermediary variable  $M$  does not affect the interpretation of the indirect effect.

# Partially Standardized Indirect Effect

---

$$ab_{ps} = \frac{ab}{SD_Y}$$

$$c'_{ps} = \frac{c'}{SD_Y}$$

$$c_{ps} = \frac{c}{SD_Y} = ab_{ps} + c'_{ps}$$

- Simple
- Removes binding to the scale of  $Y$
- Still scale-bound by  $X$
- Not clear what constitutes a “large” effect



# Completely Standardized Indirect Effect

---

$$ab_{cs} = \frac{SD_X ab}{SD_Y}$$

$$c'_{cs} = \frac{SD_X c'}{SD_Y}$$

$$c_{cs} = \frac{SD_X c}{SD_Y} = ab_{cs} + c'_{cs}$$

- Simple
- Removes all scale binding
- Not clear what constitutes a “large” effect



# Ratio of the Indirect Effect to the Total Effect

---

$$P_M = \frac{ab}{c} = \frac{ab}{c' + ab}$$

- Very simple
- Not bounded by 0 and 1
- Explodes toward  $\pm\infty$  as  $c \rightarrow 0$
- Very unstable
  - High between-sample variability
  - Requires  $N \geq 500$





# Ratio of the Indirect Effect to the Direct Effect

---

$$R_M = \frac{ab}{c'} = \frac{P_M}{1 - P_M}$$

- Very simple
- Not bounded by 0 and 1
- Explodes toward  $\pm\infty$  as  $c' \rightarrow 0$
- Very unstable
  - High between-sample variability
  - Requires  $N \geq 2000$



# Proportion of Variance in Y Explained by the Indirect Effect

---

Developed by Fairchild, MacKinnon, Taborga, and Taylor (2009).

- Given a non-zero total effect, represents the proportion of variance in Y accounted for by the indirect effect.

$$R^2_{med} = r^2_{MY} - (R^2_{Y.MX} - r^2_{XY})$$

- Mostly sensible interpretation
- Predicated on the assumption that  $\beta_{YX} \neq 0$
- $|ab| > |c| \Rightarrow R^2_{med} < 0$ 
  - Not a strict proportion



# Kappa Squared

---

Developed by Preacher and Kelley (2011).

- Gives the proportion of the *maximum possible* indirect effect represented by *ab*.

$$\kappa^2 = \frac{ab}{\max(ab)}$$

- Bounded by 0 and 1
- Values closer to 1.0 indicate a bigger effect
- A bit of a pain to calculate.



# Computing $\max(ab)$

$$a \in \left\{ \frac{\sigma_{YM}\sigma_{YX} \pm \sqrt{\sigma_M^2\sigma_Y^2 - \sigma_{YM}^2}\sqrt{\sigma_X^2\sigma_Y^2 - \sigma_{YX}^2}}{\sigma_X^2\sigma_Y^2} \right\} = [a_{low}, a_{high}],$$

$$b \in \left\{ \pm \frac{\sqrt{\sigma_X^2\sigma_Y^2 - \sigma_{YX}^2}}{\sqrt{\sigma_X^2\sigma_M^2 - \sigma_{MX}^2}} \right\} = [b_{low}, b_{high}],$$

$$\max(a) = \begin{cases} a_{high}, & \text{if } \hat{a} > 0 \\ a_{low}, & \text{if } \hat{a} < 0 \end{cases}, \quad \max(b) = \begin{cases} b_{high}, & \text{if } \hat{b} > 0 \\ b_{low}, & \text{if } \hat{b} < 0 \end{cases},$$

$$\max(ab) = \max(a)\max(b)$$

# Example

---



# Example

```
## Specify the model:
mod2 <- "
policy ~ b*sysRac + cp*polAffil
sysRac ~ a*polAffil

ab := a*b
"

## Estimate the model:
out2 <- sem(mod2, data = dat1)
##
## Extract/compute the necessary quantities:
ab <- prod(coef(out2)[c("a", "b")])
ab

[1] 0.1015958

cPrime <- coef(out2)["cp"]
##
sdY <- sd(dat1$policy)
sdX <- sd(dat1$polAffil)
##
r2MY <- with(dat1, cor(policy, sysRac))^2
```

# Compute $\kappa^2$

---

```
## Subset the data:
tmpData <- dat1[ , c("polAffil", "sysRac", "policy")]
colnames(tmpData) <- c("x", "m", "y")
##
## Extract pertinent variance/covariance elements:
cov1 <- cov(tmpData)

sYM <- cov1["x", "m"]
sYX <- cov1["y", "x"]
sMX <- cov1["m", "x"]
s2X <- cov1["x", "x"]
s2M <- cov1["m", "m"]
s2Y <- cov1["y", "y"]
```

# Compute $\kappa^2$

```
## Possible range of a:
aMarg <- sqrt(s2M * s2Y - sYM^2) * sqrt(s2X * s2Y - sYX^2)
aInt <- c(
  (sYM * sYX - aMarg) / (s2X * s2Y),
  (sYM * sYX + aMarg) / (s2X * s2Y)
)
aInt

[1] -0.4378558  0.5793099

##
## Possible range of b:
bMarg <- sqrt(s2X * s2Y - sYX^2) / sqrt(s2X * s2M - sMX^2)
bInt <- c(-1 * bMarg, bMarg)
bInt

[1] -1.289996  1.289996

##
## max(a):
aMax <- ifelse(coef(out2)["a"] < 0,
               aInt[1],
               aInt[2])
```



# Practice

---

Suppose:

1.  $\Sigma$  is given by:

	x	m	y
x	1.5		
m	0.3	1.4	
y	0.6	0.45	1.55

2. The estimated paths are:

- $a = 0.2$
- $b = 0.246$
- $ab = 0.049$

Compute  $\kappa^2$  for the estimated  $ab$ .



# References

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- Baron, R. M., & Kenny, D. A. (1986). The moderator–mediator variable distinction in social psychological research: Conceptual, strategic, and statistical considerations. *Journal of Personality and Social Psychology*, 51(6), 1173.
- Efron, B. (1979). Bootstrap methods: Another look at the jackknife. *The Annals of Statistics*, 7(1), 1–26. doi: 10.1214/aos/1176344552
- Fairchild, A. J., MacKinnon, D. P., Taborga, M. P., & Taylor, A. B. (2009). R-squared effect-size measures for mediation analysis. *Behavior Research Methods*, 41(2), 486–498.
- Preacher, K. J., & Kelley, K. (2011). Effect size measures for mediation models: Quantitative strategies for communicating indirect effects. *Psychological Methods*, 16(2), 93.
- Sobel, M. E. (1982). Asymptotic confidence intervals for indirect effects in structural equation models. *Sociological Methodology*, 13(1982), 290–312.