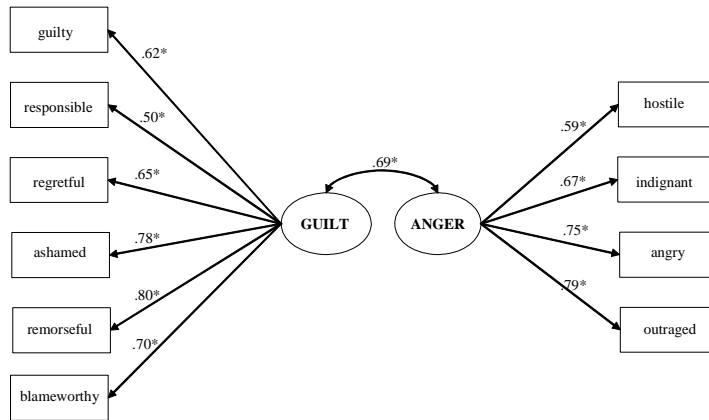


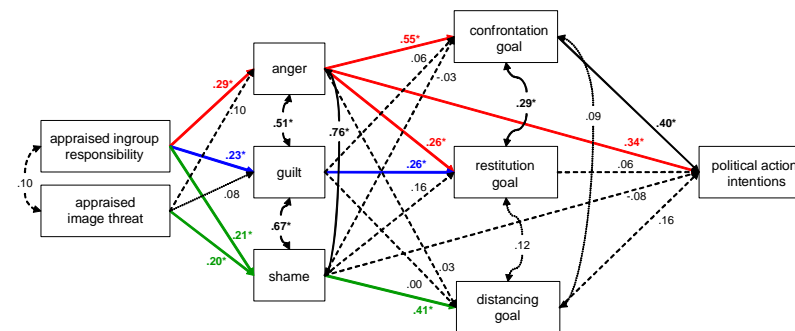
# STATISTICS WORKSHOP



## Confirmatory Factor Analysis (CFA)

&

## Structural Equation Modelling (SEM)



# overview

## PART 1: INTRODUCTION TO CONCEPTS

1. CFA: purpose, key concepts, when to use it
2. SEM: purpose, key concepts, when to use it
3. how CFA and SEM work → *5 minute break*
4. steps in CFA & SEM
  - (1) specifying a model
  - (2) evaluating model fit
  - (3) examining modification indices
  - (4) testing alternative models → *10 minute break*

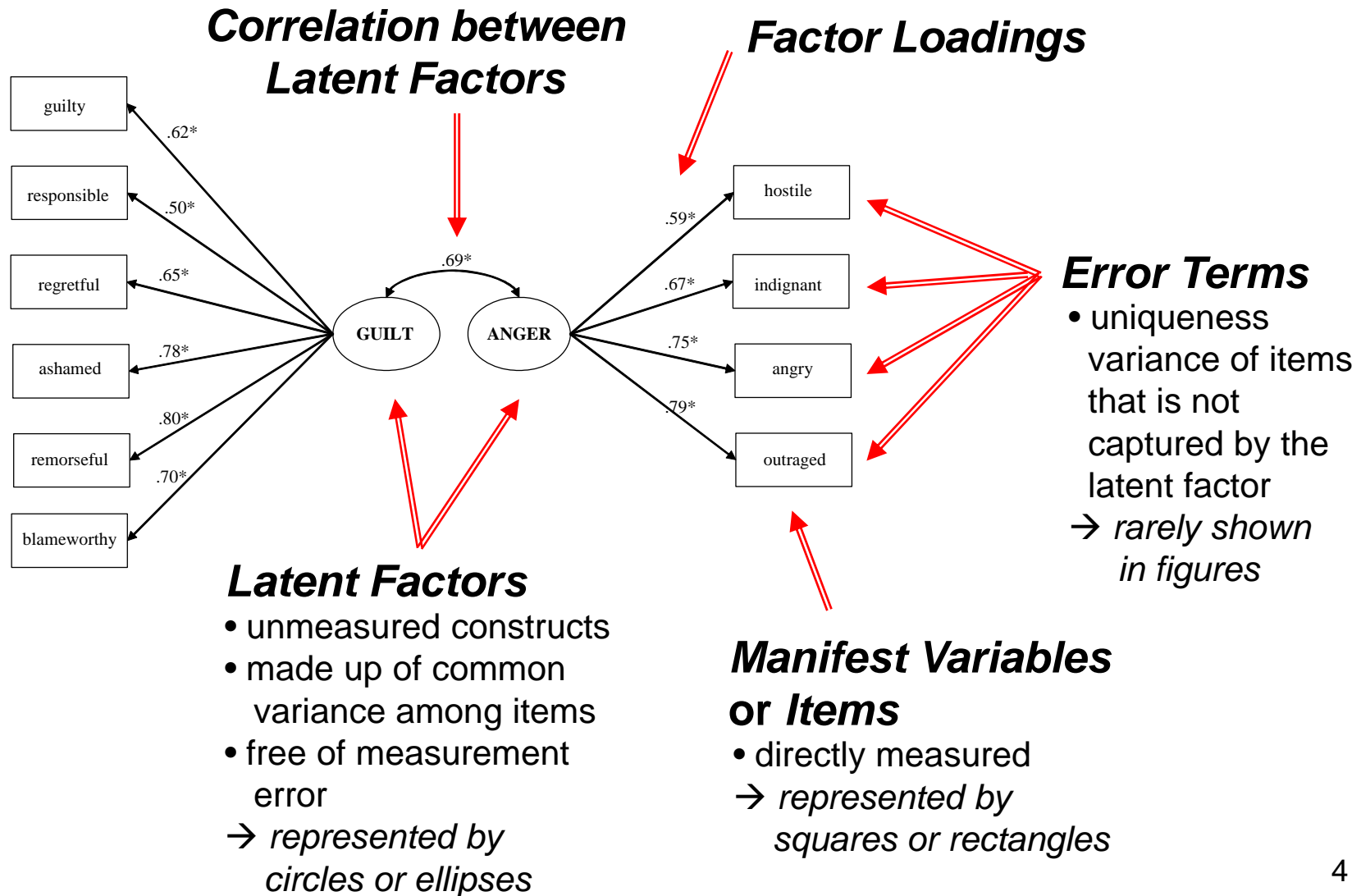
## PART 2: TESTING MODELS IN AMOS

5. how to specify a model
6. how to test a CFA model
7. how to interpret output

# purpose of CFA

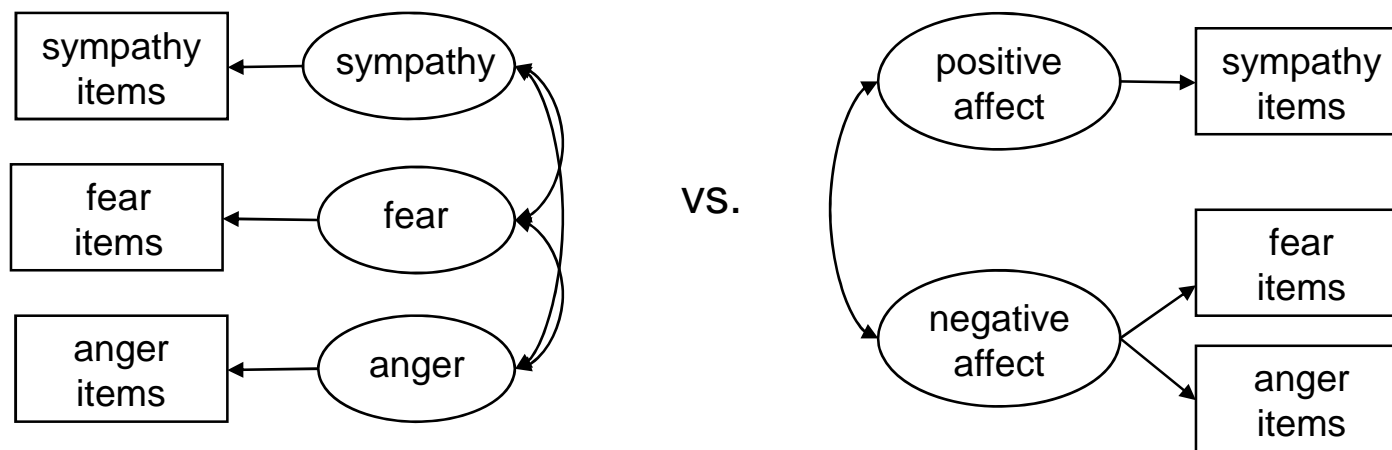
- applied to single set of variables to test hypotheses about the relative independence of subsets of variables
- similar aims to ***exploratory factor analysis (EFA)***:
  1. identify underlying constructs or factors that account for associations between subsets of variables
  2. identify how strongly each item is associated with one or more factors (factor loadings)
- key difference between EFA and CFA:
  - \* ***EFA is data-driven***
    - *SPSS calculates all possible factor loadings – factor structure is interpreted post-hoc based on the results*
  - \* ***CFA is theory-driven***
    - *program calculates only those factor loadings that we have hypothesized (all others are constrained to be “0”)*
    - *factor structure is specified a priori*

# key terms in CFA



# when to use CFA

- CFA involves *a priori* hypotheses, and so is becoming preferred over EFA (at least in social psychology)
- you might use CFA instead of EFA if you have clear hypotheses (based on previous theory and/or research) about the factors underlying a set of items
- CFA is also useful if you want to compare different factor structures that are theoretically viable



# purpose of SEM

- **used to test hypothesized relationships between variables**
  - \* assumes linear relationships
  - \* assumes multivariate normality
  - \* variables can be continuous or categorical (not in AMOS though)
  - \* can be used with correlational or experimental data
- **theory-driven approach**
  - \* researcher specifies order of association between variables
  - \* researcher specifies which relationships should be tested (all other relationships constrained to be “0”)
- **comprehensive approach**
  - \* can subsume standard techniques of regression and ANOVA

# advantages of SEM

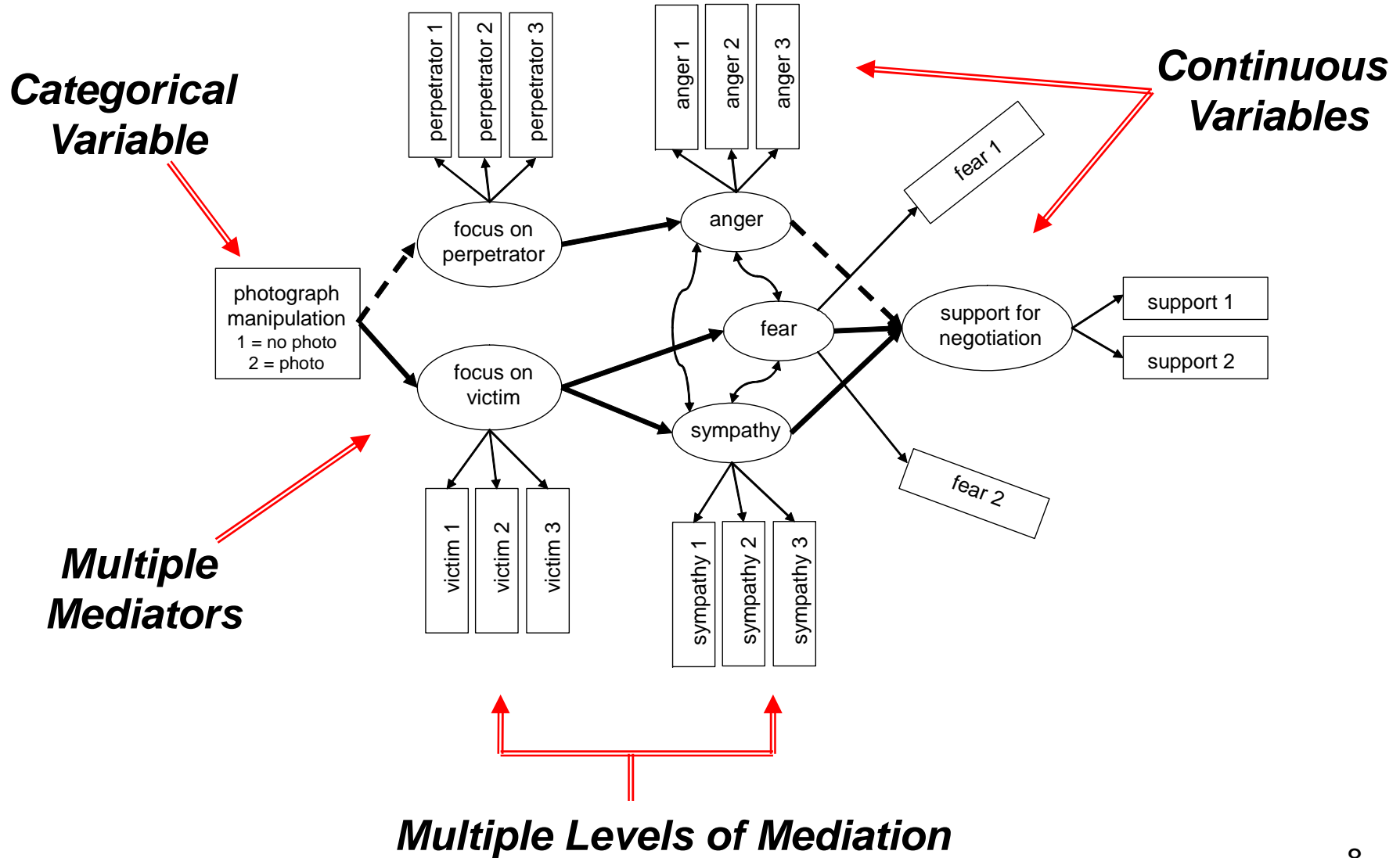
## 1. can provide more accurate estimate of relationships

- \* usually has two components
  - (1) measurement = items loading on to latent factors
  - (2) structural = relationships between factors
- \* use of latent factors partials out items' error variance (uniqueness)

## 2. can test complex sets of relationships between variables

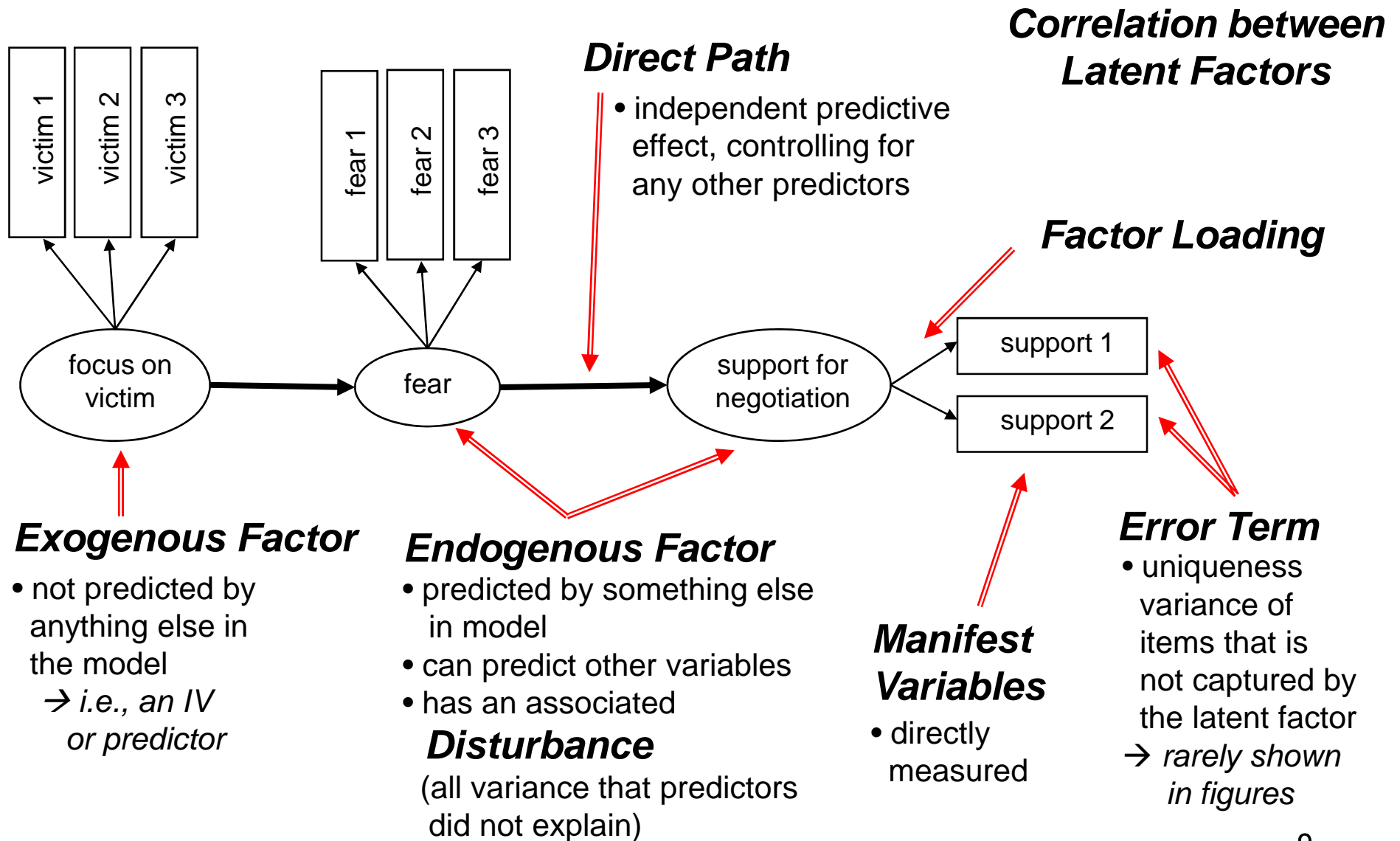
- \* **regression** can test:
  - (a) multiple predictors of 1 outcome OR
  - (b) 1 mediator between 1 predictor and 1 outcome
- \* **SEM** can test:
  - (a) multiple predictors of multiple outcomes AND
  - (b) multiple mediators between multiple predictors and multiple outcomes

# what SEM can do





# key terms in SEM



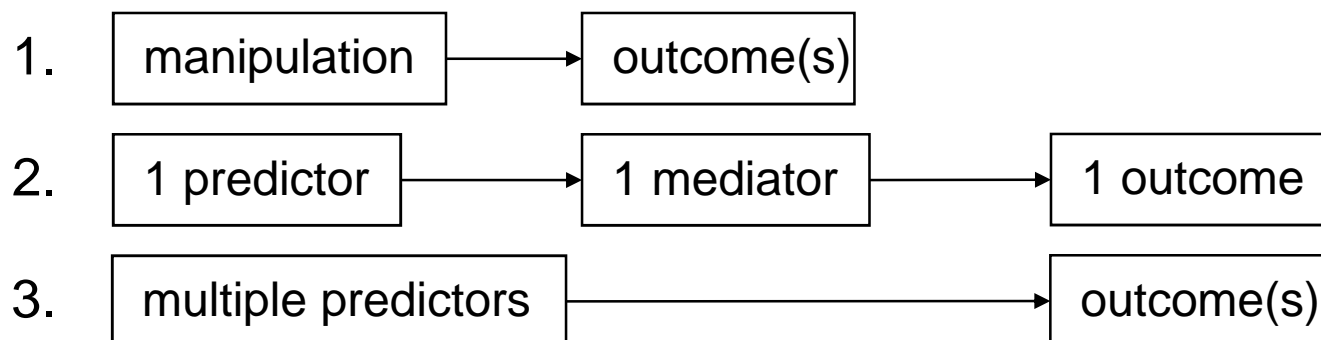
# when to use SEM

- ***SEM can be used to...***

- \* control for measurement error
  - \* test a complex set of relationships (i.e., multiple mediators and/or moderators) between a large number of variables (i.e., > 4 or 5)
- especially useful in longitudinal research

- ***SEM should not be used to...***

- \* make causal claims from correlational data
- \* test very simple models that regression or ANOVA can do:



# relationship between CFA & SEM

- both CFA and SEM use structural equation modeling procedures
  - \* specific mathematical model
  - \* theory-driven approach
- CFAs can be conducted alone – focus solely on underlying factor structure
- SEM usually includes CFA (measurement part of the model)
- key difference between the two:
  - \* **CFA** specifies correlational associations between factors in the model (i.e., bidirectional)
  - \* **SEM** specifies “causal” associations between factors in the model (i.e., unidirectional) as well as correlations

# how CFA and SEM work – (i)

- we supply two things to the software package
  - \* data set
  - \* model: statement of relationships between variables
- the software first calculates a ***variance-covariance matrix***
  - \* observed variances and covariances among variables
- it then estimates ***parameters*** in the model
  - \* parameters indicate the nature and size of relationships between variables in the population (correlations or direct paths)
  - \* we can never know the true value of a parameter, but statistics help estimate it
  - \* parameters are ***fixed*** (i.e., set to be “0”) or ***free*** (to be estimated from the data)

# how CFA and SEM work – (ii)

- based on the parameter estimates, the software computes an ***estimated variance-covariance matrix***
- it then compares the ***estimated*** and ***actual*** variance-covariance matrices
- in the end, the software produces two things:
  - (a) information regarding the similarity between the estimated and actual variance-covariance matrices  
→ *how well the model “fits” the data*
  - (b) parameter estimates  
→ *nature of relationships between variables*

\*\*\*\*\* 5 minute break \*\*\*\*\*

# summary of steps in CFA/SEM

## 1. specify a model

- A. decide on the order of association between variables
- B. decide whether each parameter should be free or constrained
- C. consider the size of your sample

## 2. evaluate model fit

- A.  $\chi^2$  test
- B. absolute and incremental fit indices
- C. residual indices

## 3. examine modification indices

- A. WALD test (*only in Lisrel & EQS, not AMOS*)
- B. Lagrange Multiplier test (*in EQS; Modification Indices in AMOS*)

## 4. test alternative models

- A. different orders of association (SEM only)
- B. nested models

# step 1: specifying a model

## A. decide on the order of association between variables

- which variables are exogenous (predictors) and which are endogenous (mediators or outcomes)?
    - \* if cross-sectional design, you have to decide
  - how to decide the order of association:
    - \* does previous theory and/or research suggest (or dictate) a particular order of association?
    - \* what is your research question?
  - can compare models with different orders of association...
  - ...**BUT** it's hard to data-fish in SEM
- ***should have clear idea of model(s) you want to test***  
→ ***should be able to defend your choice(s)***



# step 1: specifying a model

## B. decide whether each parameter should be free or constrained

- **constrained** = parameter set to be “0”  
(i.e., left out of the model)
- **free** = parameter allowed to be estimated  
(i.e., included in the model)
- need at least one constrained parameter for software to be able to estimate var-cov matrix and assess model fit
- more free parameters = model fits data better  
(fewer constraints = fewer places where “mis-fit” can occur)
- BUT more free parameters = less parsimony & fewer d.f.

→ *have to balance these two issues*

# step 1: specifying a model

**B. decide whether each parameter should be free or constrained**

## HOW TO DECIDE WHICH PARAMETERS TO INCLUDE:

- \* which relationships do you hypothesize to be important?
- \* which relationships do you have to control for?
- \* which variables actually have an association?  
(i.e., correlation > .20)
- \* which relationships do you want to show to be “0”?

→ ***need to consider theoretical, empirical, and rhetorical questions***

# step 1: specifying a model

## C. consider the size of your sample

- **issue #1: statistical stability of model**
  - \* if too few participants, mathematical basis of analysis is unsound, and output should not be trusted
- **issue #2: statistical power**
  - \* if too few participants, may not be able to detect small effects

### SO...HOW MANY PARTICIPANTS DO I NEED?

- \* ideally, 10+ participants for every estimated parameter
  - *includes factor loadings, direct effects, and correlations*
- \* if between 5 and 10 participants per estimated parameter
  - *may compromise statistical power*
- \* if < 5 participants per estimated parameter
  - *will compromise statistical stability of model*

## step 2: evaluating model fit

- model fit refers to how similar the *estimated* variance-covariance matrix is to the *actual* variance-covariance matrix  
→ *more similarity between the two matrices = good fit*
- good fit means that the hypothesized model provides a good account for the actual relationships in the dataset
- good fit does **NOT** mean that the model is “correct”  
→ only that it is plausible, and so cannot be rejected
- good fit does **NOT** mean that the model explains a large percentage of variance in the endogenous variables

# step 2: evaluating model fit

FIT DEPENDS ON SPECIFIED MODEL, BUT ALSO ON D.F.

- \* fewer degrees of freedom = fewer constraints = better chance of good fit
  - \* fewer constraints can be due to simplicity of model (i.e., fewer variables)
  - \* fewer constraints can be due to more estimated parameters (not fixed at “0”)
  - \* therefore, models that are simple and/or have more free parameters have a better chance of fitting the data well
- ***good fit is not necessarily impressive – need to look at model complexity and the # of fixed parameters***

# step 2: evaluating model fit

## A. $\chi^2$ test

- tests degree of similarity between the estimated variance-covariance matrix and actual variance-covariance matrix
- really a “badness-of-fit” index:  
large  $\chi^2$  value and small  $p$  value means that there *is* a significant difference between estimated and actual matrices
- \* rejecting the null hypothesis = model **does not** fit well
- \* accepting the null hypothesis = model **does fit** well

→ *want a small and non-significant  $\chi^2$  value*

# step 2: evaluating model fit

## A. $\chi^2$ test

### DRAWBACKS OF $\chi^2$ TEST:

- \* very sensitive to sample size  
(larger  $N$  = more chance of finding significant differences)
- \* assumption of multivariate normality is often violated

→ ***MUST report  $\chi^2$ , whether it's good or bad***

→ *if  $\chi^2$  test looks bad, you have two options:*

(1) can calculate the  $\chi^2$  / degrees of freedom ratio:

- \* divide  $\chi^2$  value by degrees of freedom
- \* if  $< 2$ , indicates good fit

(2) if other fit indices suggest good fit, downplay  $\chi^2$

# step 2: evaluating model fit

## B. absolute and incremental fit indices

- represent how much of the variance in the covariance-matrix **has** been accounted for by the model
- **NOT** testing a null hypothesis
- software calculates the following indices in this category:
  - \* normed fit index (NFI)
  - \* non-normed fit index (NNFI)
  - \* incremental fit index (IFI)
  - \* comparative fit index (CFI)
  - \* goodness-of-fit index (GFI)
  - \* adjusted goodness-of-fit index (AGFI)
- range from 0 to 1, with **higher** values indicating better fit
- general standard for good fit = .95 or higher (when  $N < 250$ )
  - \* some debate about whether this is too strict

→ **should report: NFI, IFI, CFI, GFI**



# step 2: evaluating model fit

## C. residual indices

- represent the **discrepancies** (“residuals”) between estimated and observed covariances
- **NOT** testing a null hypothesis
- software calculates the following indices in this category:
  - \* SRMR = standardized root mean squared residual
  - \* RMSEA = root mean square error of approximation
- range from 0 to 1, with **lower** values indicating better fit
- general standards for acceptable fit:
  - \* SRMR = .08 or lower
  - \* RMSEA = .06 or lower

→ **should report both SRMR (use and RMSEA**

→ *RMSEA can be sensitive to Type 1 errors (if  $N < 250$ ) and outliers*

# step 2: evaluating model fit

## COMPARING THE FIT OF DIFFERENT MODELS

- all three sets of fit indices assess absolute, rather than relative, fit
- **NEVER** compare incremental (CFI, GFI, etc.) or residual fit indices (SRMR, RMSEA) between models
  - *there is no way to test whether the difference in fit indices is statistically reliable/significant*
- **CAN** use  $\chi^2$  test to compare fit of 2 models in **one** case: when two models are nested within each other
  - *more on this in Step 4 (testing alternative models)*

# what if your model has bad fit?

- **bad fit** = model does not account for all relationships in data  
→ 1 or more fixed parameter needs to be freed
  - **what is wrong?**
    - \* CFA: an item should load onto other factors
    - \* SEM: two possible problems...
      - (1) measurement: an item should load onto other factors
      - (2) structural: a relationship between factors should be added
  - **what you need to do:**
    - \* add paths to model (modification indices can help – see Step 3)
  - **BUT remember:**
    - \* testing repeated models increases Type 1 errors
    - \* post-hoc modification moves away from *a priori* approach
- ***keep the model-tweaking to a minimum***  
→ ***make sure that your changes make theoretical sense*** 27

# step 3: modification indices

## A. WALD test → EQS only; not available in AMOS

- tests whether you can **drop** any paths you have estimated  
→ to help improve parsimony of model and free up d.f.
- using a  $\chi^2$  distribution, WALD tests whether **dropping** each parameter would significantly **worsen** the overall fit of the model
  - \* modified model = fewer estimated parameters than original
- fewer estimated parameters = worse fit (b/c more constraints)
  - \* so, the modified model (with fewer estimated parameters) will NEVER fit better than the original
  - \* the best the modified model can do is to fit just as well as the original
- \* **significant**  $\chi^2$  = dropping parameter WOULD worsen model fit  
→ **keep parameter**
- \* **non-sig.**  $\chi^2$  = dropping parameter WOULD NOT worsen fit  
→ **can drop parameter**

# step 3: modification indices

## B. LaGrange Multiplier test → “Modification Indices” in Amos

- tests whether you need to **add** any of the paths you left out  
→ to better account for relationships in the data
- using a  $\chi^2$  distribution, LM tests whether **adding** each parameter would significantly **improve** overall model fit
  - \* modified model = more estimated parameters than original
- more estimated parameters = better fit (b/c fewer constraints)
  - \* modified model (with more estimated parameters) will NEVER fit worse than the original
  - \* the worst the modified model can do is to fit just as well as the original
- \* **significant**  $\chi^2$  = adding the parameter WOULD improve model fit  
→ **add parameter**
- \* **non-sig.**  $\chi^2$  = adding the parameter WOULD NOT sig. improve fit  
→ **parameter not needed**

# step 4: testing alternative models

## A. different orders of association

- good fit does NOT necessarily mean that your model wins
  - \* does not discount other models that are as plausible
- a different order of association may be theoretically viable

*hypothesized:*



*plausible alternative:*



# step 4: testing alternative models

## A. different orders of association

### WHAT TO DO?

#### → test the alternative model

- \* examine fit indices: does it have good absolute fit?
- \* if alternative model has the same # of degrees of freedom as the original, cannot directly compare fit

→ **if** alternative model meets the absolute thresholds for good fit, you **HAVE** to say that both are viable alternatives...

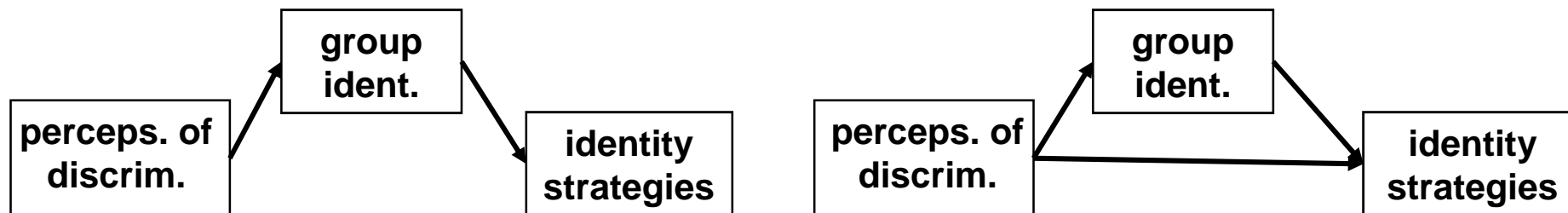
#### → ...but you can still compare relative fit:

- \* model with lower AIC value is *relatively* better fitting
- \* though cannot test reliability or magnitude of this difference

# step 4: testing alternative models

## B. nested models

- nested model = logical subset of another model
- obtained by changing number of parameters (adding or dropping paths)
- model with fewer parameters is nested within the model with more parameters



→ models include the same variables, but have different d.f.



# step 4: testing alternative models

## B. nested models

### WHAT TO DO?

compare  $\chi^2$  values to see which model fits better:

1. subtract smaller  $\chi^2$  value from larger one
2. subtract smaller degrees of freedom from larger one
3. look up corresponding  $p$  value in  $\chi^2$  table

\* ***non-significant***  $\chi^2$  = no difference in fit of the two models  
→ more parsimonious nested model fits as well as larger one  
→ larger model offers no advantage, so nested model is better

\* ***significant***  $\chi^2$  = sig. difference in the fit of the two models  
→ nested model fits significantly worse than larger model  
→ larger model is better

# variations of SEM

- **problems that may prevent you from using SEM:**
  1. your sample is too small
  2. the model does not fit well because of measurement component (e.g., identification items want to load onto well-being factor)
- but you may still be interested in testing a complex model
  - can use a variation of “proper” SEM
- **variation #1: use measured variables instead of latent factors**
  - \* create scales as you would for regression or ANOVA
  - \* specify model using measured variables, instead of building factors
  - \* no longer accounts for measurement error, but can still test complex m's
- **variation #2: use mix of measured variables and latent factors**
  - \* maybe only a subset of your measures are problematic
  - \* or, factors don't make sense for all variables (e.g., categorical)
  - \* can test a model with some measured variables and some factors
  - \* account for only some measurement error, but can still test complex m's

# more advanced possibilities...

## *(1) compare models between groups*

- does a pattern of relationships holds for different groups?
- what to do: specify the model in the two groups, constrain them to be equal, and examine the fit of this constrained model
- **good fit** = no significant differences between the groups
- **bad fit** = there is at least one significant difference
  - *can then establish which parameter estimates are different*

## *(2) compare two paths in the same model*

- is one association/relationship stronger than another?
- what to do: constrain the relevant parameters to be equal, and test the overall fit of the model
- **good fit** = no significant differences between the parameters
- **bad fit** = there is a significant difference between the parameters

\*\*\*\*\* 10 minute break \*\*\*\*\*

Kline, R. B. (2004). *Principles and practices of structural equation modeling (2<sup>nd</sup> Ed)*. New York: Guilford.

Maruyama, G. M. (1998). *Basics of structural equation modeling*. Thousand Oaks, CA: Sage.

Bollen, K. A., & Long, J. S. (1993). *Testing structural equation models*. Thousand Oaks, CA: Sage.