

KEY ADVANCES IN THE HISTORY OF STRUCTURAL EQUATION MODELING¹

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Structural equation modeling (SEM) has advanced considerably in the social sciences. The direction of advances has varied by the substantive problems faced by disciplines. For example, path analysis developed to model inheritance in population genetics, and later to model status attainment in sociology. Factor analysis developed in psychology to explore the structure of intelligence, and simultaneous equation models developed in economics to examine supply and demand.

These largely discipline-specific advances came together in the early 1970s to create a multidisciplinary approach to SEM. Later, during the 1980s, responding to criticisms of SEM for failing to meet assumptions implied by maximum likelihood estimation and testing, SEM proponents responded with estimators for data that departed from multivariate normality, and for modeling categorical, ordinal, and limited dependent variables. More recently, advances in SEM have incorporated additional statistical models (growth models, latent class growth models, generalized linear models, and multi-level models) drawn upon artificial intelligence research to attempt to “discover” causal structures, and finally, returned to the question of causality with formal methods for determining causality with non-experimental data.

In this chapter, I trace the key advances in the history of structural equation modeling. I focus on the early history, and try to convey the excitement of major developments in each discipline, culminating with cross-disciplinary integration in the 1970s. I then discuss advances in estimating models from data that depart from the usual assumptions of linearity, normality,

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and continuous distributions. I conclude with brief treatments of more recent advances to provide introductions to advanced chapters in this volume.

EARLY HISTORY: THE DISCIPLINARY ROOTS OF SEM

Sewall Wright's Path Analysis in Genetics and Biology

In 1918, Sewall Wright, a prominent geneticist and biometrician, published the first application of path analysis, which modeled the bone size of rabbits. After computing all possible partial correlations of his measures, he was still dissatisfied with the results, which remained far from a causal explanation. Consequently, Wright developed path analysis to decompose the variation in the size of an individual bone to various hereditary causes (Hill 1995). He subsequently applied path analysis to systems of mating, using data on guinea pigs, which laid the basis for much of subsequent population genetics. For example, in modeling the proportion of white color in spotted guinea pigs, Wright (1920) decomposed the variance into heredity (h), common environment for the litter (e), and other factors such as developmental noise (d). The path coefficient (h) represents the link between genotype and phenotype, and h^2 is the proportion of variance due to heredity, later termed “heritability” in population genetics. Wright also developed models for systems of mating, showing the consequences of continued inbreeding systems, such as continued brother-sister mating, which results in $m = r'_{00}$, where m is the correlation between mates in one generation, and r'_{00} is the correlation between brother and sister of the previous generation (Li 1956). He also derived results for intergenerational consequences of assortative mating. Figure 1 reproduces a path diagram of environment and heredity, which Hill (1995:1500) calls “surely one of the best known diagrams in biological science.”

Wright (1921a, 1934) presented the method of path analysis for estimating causal relations among variables based on the correlation matrix of observed variables, emphasizing path

coefficients (standardized regression coefficients) but also using “path regressions” (unstandardized coefficients). He invented a graphical method of presenting causal relations using path diagrams, consisting of variable labels connected by arrows for direct effects, double-headed arrows for unanalyzed correlations, and the estimated path coefficients listed over single-headed arrows. From path diagrams, Wright could read off total, direct, and indirect effects, and quickly decompose correlations into various causal sources, such as direct effects, indirect effects, common causes, and the like. Among the models Wright estimated by path analysis was what later became known as the MIMIC model. Wright’s estimation method was essentially the method of moments, which follows the intuitive principle of estimating a population moment (or function of moments) using the sample analog moment (or function of moments) (Goldberger 1991). Although he lacked a principle for reconciling multiple ways of expressing a path coefficient in terms of sample moments in overidentified models, he did check to see if they were close and acknowledged the potential gains in efficiency and reduced standard errors from using full information (Goldberger 1972a).

While working for the U.S. Department of Agriculture, Wright (1925) worked on corn and hog correlations, developing a complex, highly-overidentified, recursive system of equations containing observed, unobserved, lagged and un-lagged variables, to describe seasonal data on hog breeding, corn prices, and hog prices. The Department of Agriculture rejected publication of the monograph on the grounds that “an animal husbandman” (Wright’s position at the time) “had no business writing about economics” (Crow 1988). Wright’s research was only published after Henry Wallace read the paper and exploited the influence of his father, then Secretary of Agriculture. Although the recursive model had no explicit demand function, Wright (1925:54) noted in a footnote that a direct negative effect of hog quantity on hog price would be desirable

but the “treatment of such reciprocal relations between variables requires an extension of the theory of path coefficients” (see Goldberger 1972a:983).

In 1928, Sewall Wright’s father, economist Phillip Wright, published a study of the tariff, which included an appendix—coauthored by the two Wrights—that applied path analysis to reciprocal relations between variables. They began by writing out what later became known as the reduced form equations:

$$P = p_1 \frac{\sigma_P}{\sigma_D} D + p_2 \frac{\sigma_P}{\sigma_S} S \quad (1)$$

$$Q = q_1 \frac{\sigma_Q}{\sigma_D} D + q_2 \frac{\sigma_Q}{\sigma_S} S \quad (2)$$

where D and S indicate shifts in demand and supply curves after transforming P and Q to trend ratios, and σ ’s represent standard deviations of variables. Wright and Wright (1928) noted that the two equations contain four unknowns. They then suggested that if two external variables, A and B, could be found (based on external knowledge of markets) such that A were correlated with D but not S, and B were correlated with S but not D, the principles of path analysis would yield solutions for the four unknowns. The Wrights had arrived at a more general treatment than offered by Henry Schultz that year or “indeed in 1938” (Goldberger 1972a:984). Wright (1934) later developed more general solutions, noting that a mere single external variable is sufficient if the supply and demand situations were uncorrelated—that is, the disturbances of the two equations were orthogonal—and also that, in very complex models with many more shifts than external variables, one could solve for parameters by assuming plausible values for some of the unknowns (see Epstein 1987). His solution to the simultaneous equation problem would be rediscovered by Rejersøl (1945, cited in Morgan 1990), who used the term, “instrumental variable estimates” (which he attributed to Frisch).

Wright's (1934) path analysis was ignored not only in biology but statistics as well, perhaps because it contained elements "that were objectionable" to the two dominant schools of statistics (Shipley 2000:70):

The Phenomenalist school of Pearson disliked Wright's notion that one *should* distinguish "causes" from correlations. The Realist school of Fisher disliked Wright's notion that one *could* study causes by looking at correlations. Professional statisticians therefore ignored it. And biologists were drawn to Fisher's methods, which included inferential statistics, were rooted in experimental design, and were easy to grasp (Shipley 2000).

Causal Models and Path Analysis in Sociology

Perhaps the earliest reference to path analysis by a sociologist appeared in an appendix to Blalock's (1961a:191-93) classic treatment of causal inferences in nonexperimental research, where he briefly discussed "some related approaches," and concludes that path analysis "might readily be extended so as to be highly useful in the less precise nonexperimental sciences."²

Blalock had spearheaded a voluminous literature in sociology on causal models in the 1960s by elaborating on Simon's (1954) method of making causal inferences from correlational data.

Building on Yule's (1932) original definition of spurious correlation, Simon (1954) began by noting that a zero partial correlation ($r_{xy.z} = 0$) between independent variable, x , and dependent variable, y , holding z constant implies a spurious correlation between x and y when z (a confounder) is causally-prior to x and y , but an intervening variable between x and y when z is causally-prior to x and causally-subsequent to y . Simon (1954:41) correctly asserted that the determination of whether a partial correlation is or is not spurious "can only be reached if a priori assumptions are made that certain other causal relations do not hold among the variables"—

² In an essay titled "What if?" Duncan (1992) wondered whether, if he had sooner convinced Blalock that Sewall Wright's path analysis could solve Simon's questions, Blalock's appendix would have been more complete and Duncan would have lost motivation to write his later 1966 article?

namely exclusionary restrictions on coefficients and uncorrelated errors in equations. He then went on to expound on these conditions in all possible three-variable models. Blalock (1961b, 1962) extended this method to a five-variable model and then to an exhaustive exposition of the four-variable case. Later, Boudon (1965) showed that these results could be shown for path coefficients, drawing from Wright (1934), but renaming them “dependence coefficients.”

But it was Duncan’s (1966) classic expository article, “Path analysis: Sociological Examples,” and his monumental monograph with Peter Blau, *The American Occupational Structure* (Blau and Duncan 1967) that launched the path analysis movement in sociology, and later, in political science. Duncan (1966) used published correlations to apply path analysis to recursive models of class values, population density, occupational prestige, and synthetic cohorts. Subsequently, Costner (1969) and Blalock (1969) used path analysis to address multiple-indicator causal models, and in particular, to find that tetrad-difference restrictions on observed correlations provide a way of testing models—a result that emerged from factor analysis in psychology. These papers emphasized population models and glossed over sample-population distinctions, efficient estimation, and formal hypothesis testing.

Substantively, Duncan and Hodge (1963) had earlier published a four-variable status attainment model on data from Chicago, in which son’s education intervened between the effects of father’s occupation on son’s occupation in 1940 and 1950. They estimated the models using equation-by-equation multiple regression. Blau and Duncan’s (1967) monograph expanded on this model by using national data, distinguishing between hierarchies of occupations and rewards from the process by which individual characteristics sort people into those occupations, and examining whether the American stratification system approximated an open opportunity structure more than a rigid class hierarchy. The study produced an explosion of productive

research using path analysis to model status attainment, most notably the Wisconsin status attainment model (e.g., Sewell and Hauser 1975).

Factor Analysis in Psychology

In psychology, interest in SEM grew from factor analysis, which is a statistical method for analyzing a correlation matrix of observed variables to identify a small number of factors, components, or latent variables, that contain the essential information in the original variables. Thus, the primary goal is to attain “scientific parsimony or economy of description” (Harmon 1960:4). The method was originally developed to model psychological theories of ability and behavior. Spearman (1904) is often credited as the founding father of factor analysis, although earlier, Pearson (1901a) published a paper on fitting planes by orthogonal least squares—the foundation for principal component analysis—which was later applied to the analysis of correlation matrices by Hotelling (1933). Spearman’s work on factor analysis derived explicitly from his work on intelligence testing. He specified a two-factor theory of intelligence, in which all mental processes involved a general factor, g , plus specific factor, s . The general factor enters all activities—some more than others—while the specific factors were unique to the task at hand (the specific mental activity). Spearman claimed that the disparate items from intelligence tests would reveal two factors: a general factor and an item-specific factor. Moreover, Spearman (1927) showed that four variables cannot be described by a single factor unless their intercorrelations satisfy the conditions of two vanishing tetrad differences:³

$$r_{12}r_{34} - r_{14}r_{23} = 0 \quad (3)$$

$$r_{13}r_{24} - r_{14}r_{23} = 0 \quad (4)$$

³ A third tetrad difference, $r_{12}r_{34} - r_{13}r_{24} = 0$, is a function of the first two and will always be true if the first two are true.

Criticisms of the two-factor theory of intelligence on theoretical as well as empirical grounds—tetrads often failed to vanish or, equivalently, correlation matrices failed to be of unit-rank, even after considering sampling error—led to interest in multiple factor analysis, in which group factors were identified after extracting a general factor (e.g., Holzinger 1941). Thurstone—who founded the Psychometric Society—noted that a vanishing tetrad difference implied a vanishing second-order determinant of the matrix of observables, and extended this to the vanishing of higher-order determinants as a condition for more than one factor. He then generalized the result: the number of common factors is determined by the rank of the matrix of observables (see Harmon 1960). In addition, Thurstone developed the centroid method of factoring a correlation matrix (as a pragmatic compromise to the computationally-burdensome principle axis method). Moreover, he developed a definition of simple structure for factor analysis based on five principles (the most important of which are to minimize negative loadings and maximize zero loadings) to facilitate interpretation and insure that loadings were invariant to the inclusion of other items. This spawned interest in various methods of rotation of the initial solution, such as Kaiser's (1958) Varimax orthogonal rotation. Thurstone's original hand rotation was oblique, allowing factors to be correlated, but it was not until Jennrich and Sampson (1966) developed a computational method of achieving an oblique rotation and Jennrich and Clarkson (1980) worked out the standard errors of rotated loadings, that oblique solutions became practical (see Browne 2000).

The problem of rotating factor solutions is avoided when confirmatory factor analysis is used. Here, the number of factors and the pattern of loadings—including restrictions on loadings—are specified in advance, transforming the problem into one of identification of a model's parameters from observed moments—the same issue that arises in simultaneous

equation models.⁴ The factor model specifies $y = \Lambda\eta + \varepsilon$, where y is a vector of p observables, η is a vector of m latent factors, where ($m < p$), Λ is a $p \times m$ matrix of loadings, and ε is a vector of p error terms representing “unique” variance in y . Identification is typically achieved by specifying zero-restrictions on elements of Λ to create, for example, sets of congeneric tests, in which items load solely on single factors (e.g., Jöreskog 1971a). The zero loadings create tetrad difference overidentifying restrictions on observed covariances, as noted above. The covariance structure then becomes:

$$\Sigma = \Lambda\Psi\Lambda + \Theta \quad (5)$$

where $\Sigma = E(y y')$, $\Psi = E(\eta \eta')$, and $\Theta = E(\varepsilon \varepsilon')$. A maximum likelihood approach to factor analysis was developed by Lawley (1940), and fully elaborated by Anderson and Rubin (1956). But, according to Browne (2000:663), computational procedures were not available until “nested algorithms involving eigenvalues and eigenvectors and imposing inequality constraints on unique variance estimates were discovered independently by Jöreskog (1967) and by Jennrich and Robinson (1969).” If S , the covariance matrix of observables follows a Wishart distribution, the log-likelihood function of the model is

$$\log L = -\frac{1}{2} n \left[\log |\Sigma| + tr(S \Sigma^{-1}) \right]. \quad (6)$$

Jöreskog (1967) and his colleagues developed computer software programs for confirmatory factor analysis estimated by maximum likelihood.

Simultaneous Equation and Errors-in-Variables Models in Economics

The structural equation approach in econometrics is usually attributed to Haavelmo (1943) and the Cowles Commission (1952), most notably Koopmans (1945). But, as Morgan (1990) points

⁴ Anderson (1991) notes that economists like to put restrictions on models, whereas psychologists refused to do so until “Jöreskog (1969) came up with the catchy terms ‘exploratory’ and ‘confirmatory factor analysis’” with zero-restrictions on loading matrices, along with computer programs for maximum likelihood estimation. Jöreskog certainly popularized the terms, but the distinction was originally due to Tucker (1955).

out, Frisch and Waugh (1933: 389-390) were first to define “*structural* relation” as a “theoretical relation postulated *a priori*” in a single equation multivariate linear model in which the partial regression coefficient represented a “structural coefficient”: “An empirically determined relation is ‘true’ if it approximates fairly well a certain well-defined *theoretical* relationship, assumed to represent the nature of the phenomenon studied.”

Frisch (1934), however, was critical of the use of probability models for economic data (such as variations in the business cycle), which were rarely the result of a sampling process, and of OLS regression because measurement errors existed not only on dependent variables, but independent variables as well. This led him to confluence analysis, which treated observed variables as fallible indicators of latent variables, and then examined the interrelationships among all latent and observed variables to distinguish “true relations” from “confluent relations.” Frisch (1934) developed the method of bunch maps—a graphical presentation of regression coefficients—as a tool to discover underlying structure, often obtaining approximate bounds for relationships (for details, see Hendry and Morgan 1989).

According to Qin (1989), Frisch—who coined the term “econometrics” and helped found the Econometric Society and its journal *Econometrica*—had developed many of the abstract principles of identification of simultaneous equation models—although in a manner confounded with issues of estimation and testing—particularly in his critique of Tinbergen (1939). Tinbergen himself had discussed a formal way of identifying a two-equation model from reduced-form parameters (Tinbergen 1930 cited in Magnus and Morgan 1987), although in his monumental models of the Dutch and U.S. economies, he “cleverly constructed his model in the causal chain fashion,” using OLS to estimate its parameters, including effects of lagged

dependent variables (Anderson 1991).⁵ In his classic works on demand, Schultz (1938) had developed the cobweb model in which lagged price identified the supply-demand model. Remarkably, Schultz was unaware of Sewell Wright's more general instrumental variable solution to the identification problem despite the two being colleagues, friends, and hiking partners (Goldberger 1972a:985-86).

Within this context, Haavelmo (1943, 1944) made two key contributions to structural equation models in economics. First, he built on the work of Wald, Koopmans (1937) and others in specifying a probability model for economic models, presenting clearly and concisely the Neyman-Pearson (e.g., Neyman and Pearson 1933) approach to hypothesis testing, and using the probability approach for estimation, testing, and forecasting (see Morgan 1990). He also distinguished between two models of the source of stochastic components: errors-in-variables models emphasized by Frisch, and random shocks models introduced by Slutsky (1937).⁶ This framework is often referred to as the "probabilistic revolution" in econometrics (see Morgan 1990) and has had a lasting impact on the field, particularly in cementing the Neyman-Pearson approach to inference over others, such as Bayesian approaches (e.g., Jeffreys 1935). Second, Haavelmo made major advances in simultaneous equation models, showing that OLS estimates are biased in a two-equation supply-demand model, and distinguishing between structural form equations and what Mann and Wald (1943) would term, the reduced-form equation. He applied maximum likelihood estimation to the system of equations, showing its equivalence to OLS when applied to the reduced form, and specifying necessary and sufficient conditions for identification in terms of partial derivatives of the likelihood function (Haavelmo 1943, 1944).

⁵ This recursive causal chain model was later formalized and extended by Wold (1964), who criticized simultaneous equation models for ignoring the presumption that causality proceeds through time and is best modeled by recursive causal chains.

⁶ Koopmans (1937) appears to be the first to argue that residual variance in structural models was due not solely to measurement error—which implies deterministic relationships in the absence of such errors—but also due to omitting numerous minor variables from the model (see Epstein 1987:55).

Haavelmo (1944) also refined the term, “autonomy”: parameters representing *relatively* autonomous relations are more likely to be stable, intelligible, and useful for policy analysis (Aldrich 1989). Parameters, then, are structural when they represent autonomous relations, which are invariant to policy interventions.

The advances made by Haavelmo and Mann and Wald led to work on the simultaneous equation model at the Cowles Commission, which moved to Chicago in 1942, led by Marschak, and including Koopmans, Haavelmo, Wald, Lurwicz, Klein, and Anderson (Rubin and Leipnik were graduate assistants in Mathematics, and Simon joined later). Work at the Cowles Commission solved the major problems of identification, estimation, and testing of simultaneous equation models. In particular, Koopmans et al. (1950) gave a general treatment of the model’s structural and reduced forms:

$$By + \Gamma x = u \quad (7)$$

where y is a vector of p endogenous variables, x is a vector of q predetermined or exogenous variables, u is a vector of p disturbances (assumed normally-distributed) and Γ and B are coefficient matrices in which B is non-singular.⁷ The reduced form is

$$y = \Pi x + v \quad (8)$$

where $\Gamma = -B\Pi$, $u = Bv$, $\Sigma = B\Omega B'$, and Ω is the covariance matrix of v . Anderson (1991) provides an intuitive way of stating the identification problem. Suppose that in (7) some elements of B and Γ are constrained to be zero. If we rearrange the matrices so that the first row of (B, Γ) is written as $(\beta, 0, \gamma, 0)$, then the first row of $\Gamma = -B\Pi$ becomes $(\beta, 0)\Pi = -(\gamma, 0)$.

Then partition Π :

$$\Pi = \begin{pmatrix} \Pi_{11} & \Pi_{12} \\ \Pi_{21} & \Pi_{22} \end{pmatrix} \quad (9)$$

⁷ For a discussion of the concept of exogeneity, and of super-exogeneity, see Engle, et al. 1993.

and we obtain $\beta \Pi_{11} = -\gamma$, and

$$\beta \Pi_{12} = 0. \quad (10)$$

The vector β is identified (except for a multiplicative constant) by (10) if and only if the rank of Π_{12} is at least one less than the number of elements in β (Anderson 1991:7). If an equation does not satisfy this condition, it is under-identified and cannot be estimated. If an equation's restrictions on β are exactly one fewer than the rank of Π_{12} , then the equation is just-identified; if the restrictions are more than one fewer than the rank of Π_{12} , the equation is over-identified.

Koopmans et al. (1950) also specified a maximum-likelihood estimator for the general simultaneous equations model, which made Haavelmo's model accessible for empirical research.⁸

Perhaps the most important empirical applications of simultaneous equation models were Klein's (1950) Keynesian models, culminating with the 15-equation Klein-Goldberger model estimated by limited-information methods (Klein and Goldberger 1955). Others at Cowles had worried about the finite sample properties of estimation and introduced limited information methods as a solution (e.g., Anderson and Rubin 1949). Later, Theil (1953) developed a two-stage least squares estimator (2SLS), which is consistent but asymptotically inefficient. He applied OLS to the reduced form, obtained predicted values for endogenous predictors, and applied OLS to the structural form having replaced endogenous predictors with their predicted counterparts. Zellner (1962) developed a joint generalized least squares (GLS) approach to seemingly-unrelated regressions, which incorporates information on covariances among errors of equations that are otherwise unrelated. He showed that GLS estimates and standard errors are minimum variance for linear models, and gain efficiency over OLS when the x s differ across

⁸ For excellent discussion of the history of simultaneous equation models in economics, see Epstein (1987) and especially Merry (1990).

equations and covariances among errors of equations are non-zero. Zellner and Theil (1962) developed a three-stage least squares (3SLS) estimator, which applies joint GLS to the 2SLS estimates—using information from the disturbance covariances—and showed that 3SLS is consistent, asymptotically efficient, and asymptotically equivalent to full-information maximum-likelihood.⁹

The emphasis of Haavelmo and the Cowles Commission on models of errors in equations led most econometricians to abandon the errors-in-variables model emphasized by Frisch (1934). It was not until 1970 that Zellner revived interest in errors-in-variables by presenting GLS (a modification of his joint GLS estimator) and Bayesian approaches to estimating a model with a fallible endogenous predictor. Goldberger (1972b) showed that GLS is equivalent to ML only when errors are normally distributed with known variances. He also showed that when error variances are unknown, an iterated GLS will converge to ML.¹⁰

INTERDISCIPLINARY INTEGRATION

Nineteen-seventy was a watershed year for structural equation modeling: Jöreskog (1970) published his general method of analyzing covariance structures, Hauser and Goldberger (1971) presented, at the sociology meetings, their work on unobservables in path analysis, and Zellner (1970) published his GLS results on unobservable independent variables. The year 1970 was also marked by the Conference on Structural Equation Models, an interdisciplinary forum—featuring economists, sociologists, psychologists, statisticians, and political scientists—originating from an SSRC recommendation and culminating with a published volume, *Structural Equation Models in the Social Sciences* (Goldberger and Duncan 1973).

⁹ The decline of applications of simultaneous equation models in economics appears traceable to three events: (1) self-criticism by members of Cowles; Lucas' (1976) critique, in which economic agents anticipate policy interventions, and then act contrary to linear models—implying that structural parameters are not policy-invariant; and empirical research suggesting the model was not superior in forecasting the future (e.g., Nelson 1972), leading to alternative time-series models, such as vector autoregressions (Sims 1980) (see Epstein 1987).

¹⁰ For a lucid discussion of various ways that econometricians have approached measurement error, see Judge et al. (1980), chapter 13.

In this section, I will focus on two key papers published in this period: Hauser and Goldberger (1971) and Jöreskog (1973). Hauser and Goldberger's (1971) far-ranging examination of unobservable variables is an exemplar of cross-disciplinary integration, drawing on path analysis and moment estimators from Wright and sociologists, factor-analytic models from psychometrics, and efficient estimation and Neyman-Pearson hypothesis testing from statistics and econometrics. The authors analyze two models: the two-factor multiple indicator "walking dog" model (considered in factor analysis and by Costner and Blalock) and the multiple-indicator, multiple-indicator cause (MIMIC) model.¹¹ Figure 2 presents a simple walking dog model with four observables and two latent factors. We can express the model in matrix form:

$$\begin{pmatrix} y_1 \\ y_2 \\ y_3 \\ y_4 \end{pmatrix} = \begin{pmatrix} \lambda_{11} & 0 \\ 1 & 0 \\ 0 & \lambda_{32} \\ 0 & 1 \end{pmatrix} \begin{pmatrix} \eta_1 \\ \eta_2 \end{pmatrix} + \begin{pmatrix} \varepsilon_1 \\ \varepsilon_2 \\ \varepsilon_3 \\ \varepsilon_4 \end{pmatrix} \quad (11)$$

$$y = \Lambda \eta + \varepsilon$$

$$\begin{pmatrix} \eta_1 \\ \eta_2 \end{pmatrix} = \begin{pmatrix} 0 & 0 \\ \beta_{21} & 0 \end{pmatrix} \begin{pmatrix} \eta_1 \\ \eta_2 \end{pmatrix} + \begin{pmatrix} \zeta_1 \\ \zeta_2 \end{pmatrix} \quad (12)$$

$$\eta = B \eta + \zeta$$

It then follows that the covariance structure model is::

$$\Sigma_{yy} = \Lambda(I - B)^{-1}\Psi(I - B)^{-1'} + \theta_{\varepsilon} \quad (13)$$

where $\Sigma_{yy} = E(y y')$ is the (*population*) covariance matrix of observable indicators, Λ is a matrix of loadings, B is a matrix of regression coefficients among latent variables, η , $\Psi = E(\zeta \zeta')$ is the covariance matrix of structural disturbances, and $\theta_{\varepsilon} = E(\varepsilon \varepsilon')$ is a covariance matrix of measurement errors (diagonal in this example). This model is overidentified with one degree of

¹¹ The term, "walking dog model," originated with Beverly Duncan, who noted that the path diagram (see Figure 1) resembled a walking dog (Hauser, personal communication).

freedom (10 moments and 9 parameters). The overidentifying restriction implies that there is more than one way of computing parameters in terms of moments and there is a testable overidentifying restriction in the data. This can be seen by computing moments in terms of parameters (normal equations) and then solving for parameters in terms of moments. For example,

$$\lambda_{32} = \sigma_{23}/\sigma_{24} = \sigma_{13}/\sigma_{14} \quad (14)$$

By cross-multiplying the second two terms and rearranging, we obtain the identical tetrad-difference restriction as found by Spearman (1927) and given above in equation (4), but in unstandardized form: $\sigma_{23}\sigma_{14} = \sigma_{24}\sigma_{13}$. Because (14) can be satisfied by many different models, a traditional structural equation approach tests a specific nested parameterization of the restriction, rather than testing the tetrad-difference constraint on moments directly.

In estimating overidentified models, the question becomes which moment estimator(s) should be used (Hauser and Goldberger 1971). We can see this by replacing the population moments in (14) with their sample counterparts, and noting we have two moment estimators for λ_{32} . In overidentified fully-recursive models, the OLS estimator is unbiased and efficient; therefore, using the moment estimator corresponding to OLS (giving it a weight of one) and ignoring other instrumental variable moment estimators (giving them a weight of zero), is optimal. In the general case, however, one would not want to use only one estimator or a simple unweighted average, but instead weight the moments inversely to their sampling variability. Hauser and Goldberger (1971) show that this is precisely what maximum likelihood does when minimizing the fit function in equation in equation (6) above. We can illustrate this point by noting that minimizing equation (6) is asymptotically equivalent to minimizing a quadratic form (Anderson 1973; Browne 1974):

$$F_{GLS} = [s - \sigma(\theta)]' W^{-1} [s - \sigma(\theta)] \quad (15)$$

where s is a vector of non-redundant elements from the sample covariance matrix S , $\sigma(\theta)$ is the corresponding vector of elements of the parametric structure of the covariance matrix Σ —which makes $s - \sigma(\theta)$ a discrepancy vector to be minimized—and W is a weight matrix consisting of the *covariance matrix of the sample moments*. Under normality, the latter consists of products of second-order moments about the mean. Thus, the parameters in $\sigma(\theta)$ are expressed as a function of sample moments s , each of which is weighted inverse to its sampling variability by W . The estimator in equation (15), termed GLS by Browne (1974), has been applied to econometric models by Hansen (1982), who terms it the generalized method of moments.

The second model analyzed by Hauser and Goldberger (1971), the MIMIC model, is presented in a simple four-variable three-equation form in Figure 3. This model has nine parameters, ten observable moments, and therefore, one overidentifying restriction. In matrix form, the model is:

$$\begin{pmatrix} y_1 \\ y_2 \end{pmatrix} = \begin{pmatrix} \beta_{11} \\ \beta_{21} \end{pmatrix} (\eta_1) + \begin{pmatrix} \varepsilon_1 \\ \varepsilon_2 \end{pmatrix} \quad (16)$$

$$y = B \eta + \varepsilon$$

$$(\eta_1) = (\gamma_{11} \quad \gamma_{12}) \begin{pmatrix} x_1 \\ x_2 \end{pmatrix} + (\zeta_1) \quad (17)$$

$$\eta = \Gamma x + \zeta$$

and the covariance structure is:

$$\begin{aligned} \Sigma_{yy} &= B(\Gamma \Phi \Gamma' + \Psi) B' + \theta_\varepsilon \\ \Sigma_{xy} &= \Phi \Gamma' B' \end{aligned} \quad (18)$$

where $\Sigma_{xy} = E(x y')$ contains the covariances between x and y , $\Sigma_{yy} = E(y y')$ the covariances among the y s, $\Phi = \Sigma_{xx} = E(x x')$ (covariances among x s are unconstrained), $\theta_\varepsilon = E(\varepsilon \varepsilon')$ the

covariance matrix of measurement errors in y (assumed diagonal here), and $\Psi = E(\zeta \zeta')$ the covariance matrix of the structural disturbance(s). Let us fix $\beta_{21} = 1.0$ to normalize the latent variable η and give it a metric; one could also normalize by fixing Ψ to a constant.

Using path analysis, we can compute moments in terms of parameters, solve for the parameters, and obtain two ways of expressing parameters in terms of moments (Hauser and Goldberger 1971). For example,

$$\beta_{11} = \sigma_{y_1 x_1} / \sigma_{y_2 x_1} = \sigma_{y_1 x_2} / \sigma_{y_2 x_2} \quad (19)$$

Replacing the population moments with their sample counterparts gives us two sample moment estimators of β_{11} . Also, if we cross-multiply the right two terms in (19) we get a single tetrad-difference overidentifying restriction, $\sigma_{y_1 x_1} \sigma_{y_2 x_2} = \sigma_{y_2 x_1} \sigma_{y_1 x_2}$. Note that this is the same restriction on observable moments we found for the walking dog model above (if we denote all variables as y s), which illustrates an important difficulty for structural equation models: overidentifying restrictions can be satisfied by substantively different models. In general, maximum likelihood will provide consistent and asymptotically efficient estimates by weighting sample moments inverse to their sampling variability, resulting in optimal weights for multiple moment estimators. Again, minimizing the likelihood function will provide a likelihood ratio χ^2 test of overidentifying restrictions.

Hauser and Goldberger (1971) also use econometric methods to study the identification and estimation. By substituting (17) into η in (16), we obtain the reduced form:

$$\begin{pmatrix} y_1 \\ y_2 \end{pmatrix} = \begin{pmatrix} \pi_{11} & \pi_{12} \\ \pi_{21} & \pi_{22} \end{pmatrix} \begin{pmatrix} x_1 \\ x_2 \end{pmatrix} + \begin{pmatrix} \pi_{\epsilon 1} \\ \pi_{\epsilon 2} \end{pmatrix} \quad (20)$$

$$y = \Pi x + \Pi_\epsilon$$

where $\pi_{11} = \beta_{11} \gamma_{11}$, $\pi_{12} = \beta_{11} \gamma_{12}$, $\pi_{21} = 1.0 \gamma_{11}$, $\pi_{22} = 1.0 \gamma_{12}$, $\pi_{\varepsilon 1} = \beta_{11} \zeta_1 + \varepsilon_1$, and

$\pi_{\varepsilon 2} = 1.0 \zeta_1 + \varepsilon_2$. The reduced form can always be efficiently estimated using OLS. The

estimation issue arises because there are two ways of expressing structural parameters in terms of reduced form parameters:

$$\beta_{11} = \pi_{11} / \pi_{21} = \pi_{12} / \pi_{22} \quad (21)$$

This also implies a proportionality constraint on reduced form parameters, providing a test of the MIMIC specification. Maximum likelihood weights the reduced form parameter estimates $\hat{\pi}_{pq}$ inverse to their sampling variability to obtain asymptotically efficient estimates (Hauser and Goldberger 1971). In this example, there is one degree of freedom and the single constraint can be expressed in terms of observed moments or reduced-form parameters. Generally, in more complex models, both kinds of restrictions exist, and maximum likelihood will use both forms of restrictions in estimating parameters.¹² Jöreskog and Goldberger (1975) later expanded on maximum likelihood estimation of the MIMIC model and Goldberger (1973) discussed estimation in overidentified models with latent variables and simultaneity.

In a series of papers, Jöreskog (1970, 1973, 1978) outlined a general approach to covariance analysis and a computer program he called LISREL, which, following econometricians as far back as Frisch and Waugh (1933), stood for “LInear Structural RELations.” At about the same time, Keesling (1972) in his Ph.D. dissertation, and Wiley (1973) in the Goldberger-Duncan volume presented nearly identical models. However, it was Jöreskog’s version and software package that came to dominate the field. The LISREL model incorporates factor analysis, simultaneous equation models, and path analysis (as discussed above) into a general covariance structure model (e.g., Jöreskog and Sörbom 2001)

¹² Hauser and Goldberger (1971) also show that in the special case in which all tetrad-difference constraints are satisfied—such as when measurement errors are mutually correlated—modified GLS (GLS with unknown error variances) is equivalent to maximum likelihood.

$$\Sigma = \begin{pmatrix} \Sigma_{yy} & \Sigma_{yx} \\ \Sigma_{xy} & \Sigma_{xx} \end{pmatrix} = \begin{pmatrix} \Lambda_y (\mathbf{I} - \mathbf{B})^{-1} (\Gamma \Phi \Gamma' + \Psi) (\mathbf{I} - \mathbf{B})^{-1'} \Lambda_y' + \theta_\varepsilon & \Lambda_y (\mathbf{I} - \mathbf{B})^{-1} \Gamma \Phi \Lambda_x' \\ \Lambda_x \Phi \Gamma' (\mathbf{I} - \mathbf{B})^{-1'} & \Lambda_x \Phi \Lambda_x' + \theta_\delta \end{pmatrix} \quad (22)$$

Jöreskog showed that maximum likelihood estimates are obtained by minimizing the following fit function and solving for parameters:

$$F_{ML} = \log |\Sigma| - \log |S| + tr(S \Sigma^{-1}) - p + q \quad (23)$$

Where S is the sample estimate of the population covariance matrix, Σ , and p and q are constants reflecting the number of observed ys and xs respectively. If we let θ be a vector of r parameters, then the $t \times t$ covariance matrix of parameter estimates, V , is a function of the inverse of Fisher's information matrix:

$$V = \left(\frac{2}{n} \right) \left[E \left(\frac{\partial^2 F}{\partial \theta \partial \theta'} \right) \right]^{-1} \quad (24)$$

in which the square roots of the diagonal elements are asymptotic standard errors. Finally, if F_{H_0} is the minimum of (23) under the null hypothesis, and F_{H_A} is the minimum under the less-restrictive alternative, then -2 times the log likelihood ratio is

$$v = N(F_{H_0} - F_{H_A}) \quad (25)$$

which is asymptotically distributed χ^2 with $(p + q) - r$ degrees of freedom. Equation (25) can be applied to tests of nested models and the overall goodness-of-fit of the model. Jöreskog (1971b) also generalized this result to estimate the model in multiple populations, and showed how the model can be applied to simultaneous equations, MIMIC models, confirmatory factor models, panel data, simplex models, growth models, and factorial designs (for reviews, see Bielby and Hauser 1977; Bentler (1980, 1986).

In 1975, Duncan authored an excellent introductory text for path analysis and structural equation models, in which he echoed Frisch and Haavalmo's concept of autonomy—"The

structural form is that parameterization—among the various possible ones—in which the coefficients are (relatively) unmixed, invariant, and autonomous” (Duncan 1975:151). He also distinguished forms of social change, from trivial changes in sampling or exogenous variables (that leave structural coefficients intact) to deeper changes in structural coefficients (which provide fodder for explanation by multi-level models), and changes in the model’s structure itself (167), and provided sage advice for applying structural models (150):

Do not undertake the study of structural equation models (or, for that matter, any other topic in sociological methods) in the hope of acquiring a technique that can be applied mechanically to a set of numerical data with the expectation that the result will automatically be “research.”

Furthermore, he noted that if research using structural models “are contributions to science (and not merely exercises in quantitative technique), it is because the models rest on creative, substantial, and sound sociological theory” (p. 151).

The next two decades saw an explosion of the use of structural equation models in many areas of the social sciences, including stratification (e.g., Bielby, Hauser, and Featherman 1977), social psychology (e.g., Kohn and Schooler 1982), psychology (e.g., Bentler and Speckart 1981), marketing (Bagozzi 1980), mental health (e.g., Wheaton 1978, 1985), sociology of science (e.g., Hargens, Reskin, and Allison 1976), criminology (e.g., Matsueda 1982; Matsueda and Heimer 1987), adolescence (e.g., Simmons and Blyth 1987), and population genetics (e.g., Li 1975). Some extensions of the model were developed during this period. Alwin and Hauser (1975) wrote a systematic treatment of decomposing effects into total, direct, and indirect effects using path analysis. Sobel (1982, 1986) applied the delta method to obtain asymptotic standard errors for total and indirect effects. Kenny and Judd (1984) showed how to estimate a LISREL model with product terms among latent exogenous variables, and Jöreskog and Yang (1996) showed

that Browne's ADP estimator used on the matrix of augmented moments provides consistent estimates of parameters in the Kenny-Judd model as well as consistent standard errors and fit statistics.

Matsueda and Bielby (1986) and Satorra and Saris (1985) independently showed how to calculate the power of the likelihood ratio test in covariance structure models—using the non-central χ^2 distribution—and independently presented a nearly identical way of approximating the non-centrality parameter. They showed that the likelihood ratio test statistic is asymptotically equivalent to a quadratic form:

$$\nu = (\hat{\theta}_r - \theta_{r0})' V_r^{-1} (\hat{\theta}_r - \theta_{r0}) \quad (26)$$

where $\hat{\theta}_r$ is the ML estimator for the r th parameter θ_{r0} is the corresponding null hypothesis, and V_r is the asymptotic covariance matrix of $\hat{\theta}_r$, or in other words, r -dimensional submatrix of (24). Under the null hypothesis, ν has a central χ^2 with r degrees of freedom. Under the alternative hypothesis, ν has a non-central χ^2 distribution with r degrees of freedom and non-centrality parameter:

$$\tau = (\theta_r - \theta_{r0})' V_r^{-1} (\theta_r - \theta_{r0}) \quad (27)$$

where θ_r is the population parameter corresponding to the alternative hypothesis and θ_{r0} is the population parameter corresponding to the null hypothesis (see Kendall and Stuart 1979:246-7). Matsueda and Bielby (1986) then drew on Hauser and Goldberger (1971) and Goldberger (1973) to show analytically that, in a walking dog model (Figure 2), adding indicators to the endogenous latent variable increases the power of the test of β_{21} , depending on the reliability of the indicators. This is analogous to adding cross-sections to a pooled time-series cross-section

econometric model estimated by GLS. They also gave simulation results for adding indicators to the exogenous latent variable (see Matsueda and Bielby 1986).

Applied researchers obsessed over the global goodness of fit likelihood ratio χ^2 test because in large samples, models with many overidentifying restrictions tend to be rejected even when each restriction only departs trivially from the null hypothesis. This gave rise to a cottage industry of fit indices designed to offset the effect of sample size on test statistics. From this literature, it seems that a consensus is emerging that the most useful fit indices are Steiger's (1990) root mean squared error of approximation (RMSEA) (see Browne and Cudeck 1993) and Raftery's (1993, 1995) application of Schwartz's (1978) Bayesian information criterion (BIC). (For details, see West et al., this volume).

ADDRESSING VIOLATIONS OF DISTRIBUTIONAL ASSUMPTIONS

At this time, a major criticism of structural equation models argued that the assumptions of continuous observed variables, multivariate normal distributions, and large sample sizes—needed to capitalize on the asymptotic properties of maximum likelihood estimation and testing—were rarely met in practice. Some early Monte Carlo studies, such as Boomsma (1983), which created non-normal errors by categorizing continuous variables, found that estimators were robust when samples were greater than 200, but that skewness due to categorization produced spurious measurement error correlations, and biased standardized coefficients (see Bollen 1989 for a review).¹³

Asymptotic Distribution-Free Estimator

As noted above, Browne (1974) introduced the quadratic form estimator he termed generalized least squares, which yielded optimal estimates for normally-distributed observable variables when W is the covariance matrix of the sample moments, (see equation 15). Subsequently,

¹³ Wold's (1982) partial least squares "soft modeling" approach to causal chain models provides estimates that are consistent but not necessarily asymptotically efficient under arbitrary distributions.

Browne (1984) made a landmark contribution by developing what he termed an “asymptotic distribution-free” (ADF) estimator, by incorporating information about higher-order moments into the weight matrix of equation (15), which can be written in scalar form as

$$F_{GLS} = \sum_{g=1}^k \sum_{h=1}^k \sum_{i=1}^k \sum_{j=1}^k w^{gh,ij} [s_{gh} - \sigma_{gh}(\theta)] [s_{ij} - \sigma_{ij}(\theta)] \quad (28)$$

where s_{gh} is the sample covariance between variables g and h , $\sigma_{gh}(\theta)$ is the corresponding element of $\Sigma(\theta)$ implied by the model, and $w^{gh,ij}$ is a typical element of W^{-1} , which is $u \times u$, where $u = k(k + 1)$, and k is the number of observables. Browne (1984) showed that if W is a matrix with typical element

$$w_{gh,ij} = m_{ghij} - s_{gh}s_{ij} \quad (29)$$

where m_{ghij} is the fourth-order moment about the mean, then minimizing equation (15) yields the ADF estimator, which is minimum variance consistent within the class of estimators in the form of (15) under the mild assumption that eighth-order moments are finite (Browne 1984:710).¹⁴

Browne (1984) presented the asymptotic covariance matrix for $\hat{\theta}_{ADF}$ and an asymptotic χ^2 test statistic, as well as an estimator for elliptical distributions, which have zero skewness but kurtosis that departs from multivariate normality.

Browne’s (1984) ADF and elliptical estimators first appeared in Bentler’s (1995) EQS program, followed by Jöreskog’s LISREL program. Recent work has examined the finite sample properties of ADF, finding that it works well in very large samples. Other techniques available are using corrections to covariance matrix of ML estimators (e.g., Browne 1984) or a bootstrap method (Bollen and Stine 1993). Browne’s ADF estimator was also crucial for a second important advance: developing models for ordinal, limited, and discrete outcomes.

¹⁴ Note that $m_{ghij} = 1/n \sum (x_g - \bar{x}_g)(x_h - \bar{x}_h)(x_i - \bar{x}_i)(x_j - \bar{x}_j)$ is a sample estimator of $\sigma_{ghij} = 1/n \sum [x_g - E(x_g)][x_h - E(x_h)][x_i - E(x_i)][x_j - E(x_j)]$

Models for Ordinal, Limited, and Discrete Outcomes

Structural equation models are often applied to survey data, in which items are measured on dichotomous or ordinal scales, violating the assumption of continuous and normally-distributed observed variables. Muthén (1984) has made seminal contributions for analyzing dichotomous, ordinal, and limited dependent variables within a covariance structure framework. The trick is to estimate scale-appropriate correlation coefficients (e.g., polychoric and polyserial) and then use a variation of Browne's (1984) ADF estimator. The polychoric correlation, which goes back to Pearson (1901b), computes a correlation under the assumption that the ordered categories can be represented by contiguous intervals on a continuous scale (correlations between ordinal and continuous variables are termed polyserial correlations).¹⁵ Thus, the ordinal variable is related to the underlying normally-distributed continuous latent variable through a threshold model. Early work on factor models for dichotomous variables include Bock and Lieberman (1970) who used tetrachoric correlations and an ML estimator for a single factor model, and Christofferson (1975) who generalized this to multiple factors using a GLS estimator (see also Muthén 1978). Muthén (1979) developed a multiple-indicator structural probit model and Winship and Mare (1983, 1984) applied multivariate probit models estimated by ML to multiple-indicator structural equation models and path analysis.

Muthén (1984) provided a general framework for analyzing ordinal variables. I will focus on the polychoric and ADF approach with a simple example of a pair of three-category ordinal variables. Each ordered variable is related to an underlying continuous variable by two thresholds:

¹⁵ Pearson's tetrachoric correlation, involving pairs of dichotomous variables, led to his famous quarrel with Yule (1912), who argued that many binary outcomes, such as death, cannot be viewed as reflections of a continuous scale, and proposed his Q-coefficient instead (see Pearson and Heron 1913).

$$\begin{aligned}
y_1 &= 1 && \text{if } y^* \leq \alpha_1 \\
y_2 &= 2 && \text{if } \alpha_1 \leq y^* < \alpha_2 \\
y_3 &= 3 && \text{if } \alpha_2 \leq y^*
\end{aligned} \tag{30}$$

where the subscript for y indexes the ordinal category for y , y^* is a latent continuous variable, and α_1 and α_2 are thresholds. If we specify a distribution for y^* —we will assume it is normal—we can then estimate the thresholds by the general formula:

$$\alpha_i = \Phi^{-1} \sum_{k=1}^i n_k / N \quad i = 1, 2, 3 \quad k = 1, 2 \tag{31}$$

where i indexes the category of y , k indexes the number of thresholds, $\Phi^{-1}(\cdot)$ is the inverse of the standard normal distribution function, n_k is the sample size of the k th category, and N is the total sample size and $N = n_1 + n_2 + \dots + n_k$. If we apply the above to a second three-category ordered variable x , but with thresholds β_1 and β_2 , and define π_{ij} as the population parameter denoting that an observation falls into cell (i, j) , we can then define the log-likelihood function of the sample (Olsson 1979):

$$\log L(n_{ij} | \pi_{ij}) = c \sum_{i=1}^3 \sum_{j=1}^3 n_{ij} \log \pi_{ij} \tag{32}$$

where $\pi_{ij} = \Phi_2(\alpha_i, \beta_j) - \Phi_2(\alpha_{i-1}, \beta_j) - \Phi_2(\alpha_i, \beta_{j-1}) + \Phi_2(\alpha_{i-1}, \beta_{j-1})$, and $\Phi_2(\cdot, \cdot)$ is the bivariate normal distribution function with population correlation ρ . Maximizing (32) will yield the ML estimator of the polychoric correlation, $\hat{\rho}_{ML}$. Alternatively, one can use a two-step procedure, estimating the thresholds α_i and β_j from the marginals of the contingency table (e.g., 31), and then solving for ρ . See Olsson (1979) for a parallel treatment of the polyserial correlation between continuous and ordered variables, and Poon and Lee (1987) for multivariate ML estimators of both polychoric and polyserial correlations.

Once the polychoric and polyserial correlations $\hat{\rho}$ and their asymptotic covariances have been estimated, Browne's (1984) ADF fitting function can be used to obtain optimal estimates:

$$F_{ADF} = [\hat{\rho} - \sigma(\theta)]' S_{\rho\rho}^{-1} [\hat{\rho} - \sigma(\theta)] \quad (33)$$

where $\hat{\rho}$ is a vector of scale-appropriate correlation estimates, $\sigma(\theta)$ is the corresponding vector of the parametric structure generating the correlations, and $S_{\rho\rho}^{-1}$ is the inverse of the asymptotic covariance matrix of the correlation estimates. Standard errors and χ^2 test statistics are obtained as above (Muthén 1984). Muthén (1989) has also developed a tobit factor analysis for censored observed variables. A general method for handling dichotomous and limited dependent variables in SEM was initially programmed in Muthén's LISCOMP program, and then in his recent more comprehensive Mplus program (Muthén and Muthén 2004), and later in Jöreskog and Sörbom's (1996, 2001) PRELIS and LISREL programs, and Bentler's (1995) EQS. Much of this material is covered in Bollen's (1989) excellent intermediate-level SEM text.

RECENT ADVANCES: DISCOVERY, CAUSALITY, GENERALIZED MULTILEVEL LINEAR MODELS, AND BAYESIAN INFERENCE

Major contemporary advances in structural equation modeling make it an exciting and growing field. These include the development of latent growth and latent class growth models for longitudinal data, the application of Bayesian methods, the integration of generalized linear models and multi-level models within an SEM framework, the adoption of algorithms from artificial intelligence to discover causal structure, and a formal treatment of causality within an SEM framework.

Latent Growth and Latent Class Growth Models

Although the use of factor analysis for modeling panel data on growth was introduced by Tucker (1958) and Rao (1958), it was not until 1990 that Meredith and Tisak (1990) published the

treatment within an SEM framework that is still relevant today (see Bollen and Curran 2006). Meredith and Tisak showed that individual growth curves, often modeled within a multi-level or mixed model framework (e.g. Raudenbush and Bryk 2002), can be modeled within a standard SEM framework by treating the shape of growth curves as latent variables with multiple indicators consisting of the variable at multiple time points. This latent growth curve approach models both covariances and means of observed variables. Figure 4 presents a path diagram of a four-wave quadratic latent growth curve model. Here, the intercept, α , gives the value of y implied by the model at the first time point, β_1 is the linear growth component (giving the growth rate at the first time point), and β_2 is the quadratic growth component (giving the change in the growth rate over time). One can then impose a parametric structure on the growth parameters, α , β_1 and β_2 , which would correspond to the second-level equation in a multi-level model.

In criminology, Nagin and Land (1993) develop a finite mixture model for latent classes of individual trajectories. This group-based trajectory model estimates individual trajectories using polynomials and then classifies the trajectories into discrete groups. The latent classes can be viewed as points of support in approximating a continuous distribution of unobserved heterogeneity or as reflections of theoretically important groups (see Nagin 2005). In criminology, this model has been used to distinguish criminal careers, such as chronic offenders, early starters, and adolescence-limited offenders (see Nagin and Tremblay 2005). Muthén (2004) shows how to estimate this model within an SEM framework with Mplus. Moreover, Muthén's approach, termed growth mixture modeling, allows for within-class variation among individual trajectories—a mean curve with variation around it— whereas Nagin's approach does not. The latter is nested within the former and can be subjected to statistical test. These models have become important features of research in child development, adolescent substance abuse, and criminal careers (e.g., Nagin and Tremblay 2005; Connell and Frye 2006).

Bayesian Approaches

As noted above, work at the Cowles Commission helped cement the Neyman-Pearson frequentist approach to estimation and hypothesis testing in econometrics, which spread to psychology through Anderson and Jöreskog, and to sociology through Goldberger and Duncan. In recent years, alternatives—particularly Bayesian approaches—have been proposed for SEM (for an early and persistent advocate in economics, see Zellner 1971). From a Bayesian perspective, estimation is less about deducing the values of population parameters and more about updating, sharpening, and refining our beliefs about the empirical world.

Bayesian estimation begins with a probability model of the data D in terms of a vector of parameters θ (e.g., Raftery 1995). The analyst's prior beliefs about the uncertainty of θ is denoted by the prior probability density, $p(\theta)$. The probability model for the data, then, is the likelihood function, $p(D|\theta)$, which is the probability of the data given that θ are the true parameters. We then observe the data D and update our beliefs about θ using Bayes' theorem:

$$p(\theta | D) = [p(D | \theta)p(\theta)] / p(D) \quad (34)$$

The data are treated as a fixed set of information to be used in updating our beliefs about the parameters. Therefore, $p(D)$ does not involve θ , and equation (34) reduces to

$$p(\theta | D) \propto p(D | \theta)p(\theta) \quad (35)$$

where \propto means “proportional to.” The marginal density of the data has been dropped; to make this a proper density, a proportionality constant can be added. Thus, the posterior density is proportional to the likelihood times the prior density. Inferences about θ are made from summary statistics about the posterior density, $p(\theta | D)$, such as the posterior mode or Bayesian confidence intervals (“credible intervals”), which have an intuitive interpretation: “the probability that the true value of the parameter lies in the interval is—for example—95%.”

Bayesian hypothesis testing entails comparing hypotheses to determine which has the highest probability of being correct. Suppose we have two hypotheses, H_0 and H_1 , with *prior* probabilities, $p(H_0)$ and $p(H_1)$ before the data are examined, and define the prior odds ratio as

$$Odds_{prior} = p(H_0)/p(H_1) \quad (36)$$

After examining the data, the prior probability will be updated, resulting in posterior probabilities for each hypothesis, $p(H_0 | D)$ and $p(H_1 | D)$, and a posterior odds ratio:

$$p(H_0 | D)/p(H_1 | D) = Odds_{posterior} = B_{01} \times Odds_{prior} \quad (37)$$

where B_{01} is the Bayes factor:

$$B_{01} = p(D | H_0)/p(D | H_1) \quad (38)$$

and $p(D | H_0)$ and $p(D | H_1)$ are the marginal probabilities of the data. Equation (37), the posterior odds ratio, gives the probability that the data support H_0 over H_1 . Note that the posterior odds are equal to the Bayes factor when the prior odds are equal to 1.

Bayesian estimation and testing is currently diffusing into the SEM literature. For example, Raftery (1993, 1995) showed how to approximate the Bayes factor with the Bayesian Information Criterion (BIC), which is computed from the likelihood-ratio test statistic. Suppose we wish to compare two models in which M_{k-1} is nested within M_k and has one more parameter than M_k . Then, if ν_{k-1} is the likelihood ratio test statistic for model M_{k-1} and ν_k is that of model M_k and $\nu = \nu_{k-1} - \nu_k$ is the test statistic for testing the one parameter (see equation 25), then

$$BIC_{k-1} - BIC_k \approx \nu - \log n \quad (39)$$

where n is the sample size. If this quantity is positive, then the less-restrictive model M_k is preferred (Raftery 1995). As a fit statistic, BIC has performed exceedingly well in a variety of contexts and is particularly useful for SEM in large samples and with many overidentifying

restrictions because trivially small departures from a reasonable model will be rejected using the likelihood ratio test.

Bayesian estimation using Markov chain Monte Carlo (MCMC) algorithms are proving useful for incorporating prior information into confirmatory factor analysis (e.g., Lee 1981), estimating complex models, such as nonlinear latent variable models (e.g., Arminger and Muthén 1998), estimating multilevel factor models (Goldstein and Browne 2002), and arriving at a semiparametric estimator (Yang and Dunson 2010). For details, see Kaplan and Depaoli, this volume.

Generalized Linear Latent and Mixed Models

When data take on a hierarchical structure—such as individuals nested within families, which are in turn, nested within neighborhoods—special methods are needed to obtain consistent estimates of standard errors and test statistics due to dependent observations within clusters. Multi-level regression models allow estimation of models in which random intercepts capture heterogeneity between clusters in the dependent variable and random coefficients capture heterogeneity in relationships among independent and dependent variables. A multi-level structural equation model would incorporate multiple-indicator measurement models into the latent variable models. Early attempts to incorporate measurement error into multilevel regression models have assumed that measurement error variances (e.g., Goldstein 1995) or factor loadings (e.g., Raudenbush and Sampson 1999) are known and have the advantage that unbalanced designs, in which the number of level-1 cases varies by level-2, is easily handled if missing at random (see Rabe-Hesketh et al. 2004).

Multi-level structural equation models have typically specified separate models for within-cluster and between-cluster covariances matrices. For example, Muthén (1994) has shown how to estimate a two-level SEM using available SEM software. The trick is to specify separate

within- and between-cluster models, and then use the multiple group option to estimate the parameters simultaneously. Muthén (1994) argues that an estimator using this method is equivalent to maximum likelihood in balanced designs, and consistent (with reasonable standard errors and test statistics) in unbalanced designs (see also Goldstein and McDonald 1988; Muthén 1997). (For a review of alternate estimators, see Yuan and Bentler 2007). This approach is easily implemented using existing SEM software, but is limited to specific models.

A more general approach is outlined in Rabe-Hesketh et al. (2004), and expanded upon in Skrondal and Rabe-Hesketh's (2004) excellent advanced text. This generalized linear latent and mixed modeling (GLLAMM) framework consists of three components: (1) a response model; (2) a structural equation model for latent variables; and (3) distributional assumptions for latent variables. The response model is simply a generalized linear model conditional on the latent variables, and consisting of a linear predictor, a link function, and a distribution from the exponential family (Rabe-Hesketh et al. 2004). The model can handle response variables that are continuous, ordinal, dichotomous, discrete time durations, counts and durations in continuous time, polytomous responses and rankings, and mixes of responses. The structural equation for latent variables takes on the usual form, $\eta = \beta\eta + \Gamma\xi + \zeta$, with the exception that latent variables are allowed to vary by different levels. Rabe-Hesketh et al. (2004) assume the latent variables at level l are distributed multivariate normal with zero mean and covariance matrix Σ_l , although other distributions can be specified. The authors have also written a program, *GLLAMM*, which maximizes the marginal likelihood using an adaptive quadrature procedure, and is available in the software package, Stata (Rabe-Hesketh et al. 2001). For more details, see Skrondal and Rabe-Hesketh (2004) and Rabe-Hesketh et al. in this volume. Most of these models can also be estimated using Mplus (Muthén and Muthén 2004).

Tetrad: The Discovery of Causal Structure

A philosophically distinct approach to SEM developed with the publication of Glymour et al.'s (1987) *Discovering Causal Structure: Artificial Intelligence, Philosophy of Science, and Statistical Modeling*. Instead of focusing on estimation and testing of structural models specified on *a priori* grounds, Glymour et al. (1987) draw on computer algorithms from artificial intelligence to “discover” causal structure with their program, TETRAD. Thus, they are returning to the earlier ideas of Spearman, Frisch, Simon, Blalock, and Costner, who tried, in various ways, to induce causal structure from patterns of association among variables.¹⁶ As we noted above, Spearman's focus on tetrad difference restrictions on observed correlations became superfluous in light of Thurstone's rotated solution to simple structure for factor models; Frisch's confluence analysis and bunch mappings became obsolete with advances in identification and estimation in simultaneous equations; and Simon and Blalock's method of ransacking three- and four-variable models became outdated with the development of estimation and testing using ML and GLS in integrated SEMs. These “outdated” approaches have been resurrected by Glymour et al. (1987).

Beginning with the observation that an infinite number of models are consistent with any covariance matrix of observed variables, Glymour et al. return to Simon and Blalock's method of identifying the vanishing partial correlations that must hold for a given model and to the writings of Wright, Spearman, and others, who identified the tetrad difference equations that must hold for a given model. They provide a fascinating philosophy of science edifice to justify the idea of discovering causal structure. Moreover, they use the terminology of directed graphs—rather than path analysis—in which variables are vertices, causal effects are directed edges that can be into a vertex (the number denoted by indegree) and out of a vertex (the number denoted by outdegree), a recursive model is acyclic, a non-recursive model (in which a path contains a

¹⁶ A feature of Wold's (1982) soft modeling approach is the further development of a model through a “dialogue between the investigator and the computer.”

subpath beginning and ending in the same vertex) is a cyclic model, and a trek is a path or a set of paths that induce a correlation. Using these notions, they provide basic theorems and definitions about causal relations and a computer program, TETRAD, for discovering causal structure. The program allows users to incorporate a priori information about the data—such as a tentative or partial model—identifies the vanishing tetrad differences and vanishing partial correlations of the model, and then provides a test of the constraints. It then modifies the model by identifying the treks needed to satisfy a “false” tetrad equation without altering “true” tetrad equations, and calculates vanishing tetrads and partial correlations implied by the new model. Bollen and Ting (1993) develop a confirmatory tetrad analysis approach for testing SEMs, including some under-identified models.

The TETRAD approach has been shown to be more efficient at modifying models than existing procedures available in SEM packages, such as using first-order partial derivatives, residuals, or univariate Lagrange multipliers (termed “modification indexes” in LISREL). Moreover, by introducing the language of directed graphs into structural equation models, it helps bridge the SEM literature with new developments in graphical theory and causal analysis (see Spirtes, et al. 1993).

Causality and Structural Equation Models

An exciting recent development, which is slowly filtering into the SEM community, has been an explicit return to causal models. As a consequence of Freedman’s (e.g., 1987) persistent criticisms of SEM applications for making causal claims when assumptions of linear models are not met, most members of the SEM community have retreated from making causal claims and using causal language.¹⁷ However, a recent literature, emerging from disparate sources, such as artificial intelligence, philosophy of science, and epidemiology, has developed graphical models

¹⁷ For a lucid response to Freedman’s (1987) critique of Blau-Duncan, see Glymour et al. (1987).

for identifying causality. Such models, which generalize and extend Wright's (1921) original path analysis, appeared in the social science literature as early as 1982, but did not catch on (see Kiiveri and Speed 1982). Since then, major advances have been made in artificial intelligence (e.g., Pearl 1988), statistics (e.g., Wermuth and Lauritsen 1983; Spirtes et al. 1993), epidemiology (Robins and Greenland 1992; Greenland et al. 1999), and culminating with Pearl's (2000) monograph, *Causality*.

This approach begins with the classical SEM assumption that causality cannot be determined from observational data alone, but requires additional causal assumptions drawn from theoretical or substantive knowledge, which are translated into a structural model represented by a path model. A graphical model approach represents the causal assumptions by a graphical model and then logically infers causality from a set of theorems applied to the graph. Thus, Pearl (2000) develops a "nonparametric structural causal model" which holds regardless of distributional and other statistical assumptions about a particular dataset. Causal statements can be made that are conditional on the causal assumptions encoded into the graphical model.

Pearl's (2000) approach moves beyond SEM by (1) using new mathematical notation to reflect causality, such as replacing the algebraic equals sign with a sign that reflects a causal path; (2) deriving a theorem, the back-door criterion to determine which covariates should be controlled to arrive at a causal relationship in an SEM; (3) deriving a theorem, termed "d-separation" (directed separation), which gives the necessary and sufficient conditions for independence between two sets of variables conditioned on a third set within an acyclic directed graph; (4) provide some simple mathematical notation for making counterfactual statements, which can be analyzed within the directed graph (for an introduction, see Morgan and Winship 2007); and (5) providing an algorithm for identifying equivalent models.. Taken together, these

theorems translate the language of causality into the language of statistics and probability distributions.

I will close with an example illustrating the importance of Pearl's approach. Consider Model A in Figure 5, a bivariate regression model of Y on X , with two latent variables— ξ affects X and C , and η affects Y and C . Standard SEM texts assume that including an irrelevant variable in a linear regression model leaves estimates unbiased, but results in a loss of precision in the estimate (e.g., Greene 2003:150-1). However, when Model A is the correct model, regressing Y on X and Z —which appears irrelevant—results in biased and inconsistent estimates of β . We can see this intuitively from Model A; the covariance between Z and X and Z and Y are spurious due to the latent variables ξ and η . Spirites et al. (1995:190-1) show that if we simplify Model A to Model B, and compute parameters in terms of covariances and partial covariances, we obtain an estimator from regressing Y on X and Z , which is biased and inconsistent:

$$E(X, Y | Z) / E(X^2 | Z) = \beta \sigma_{xz} \sigma_{yz} / (\sigma_x^2 \sigma_z^2 - \sigma_{xz}^2) \quad (40)$$

Where the term left of the equality sign is the (naïve) two-variable regression coefficient, and σ_{xz}^2 is the squared covariance of X and Z . A graphical approach quickly reveals not only that an unbiased and consistent estimate of β is obtained by the bivariate regression of Y on Z , but also that a consistent estimator can be obtained by the naïve two-variable regression by also controlling for ξ , η , or both (Greenland and Brumback 2002).

For an excellent presentation of the graphical approach to SEM, see Pearl (2000), and for a lucid introduction to the principles and issues, see Pearl (this volume).

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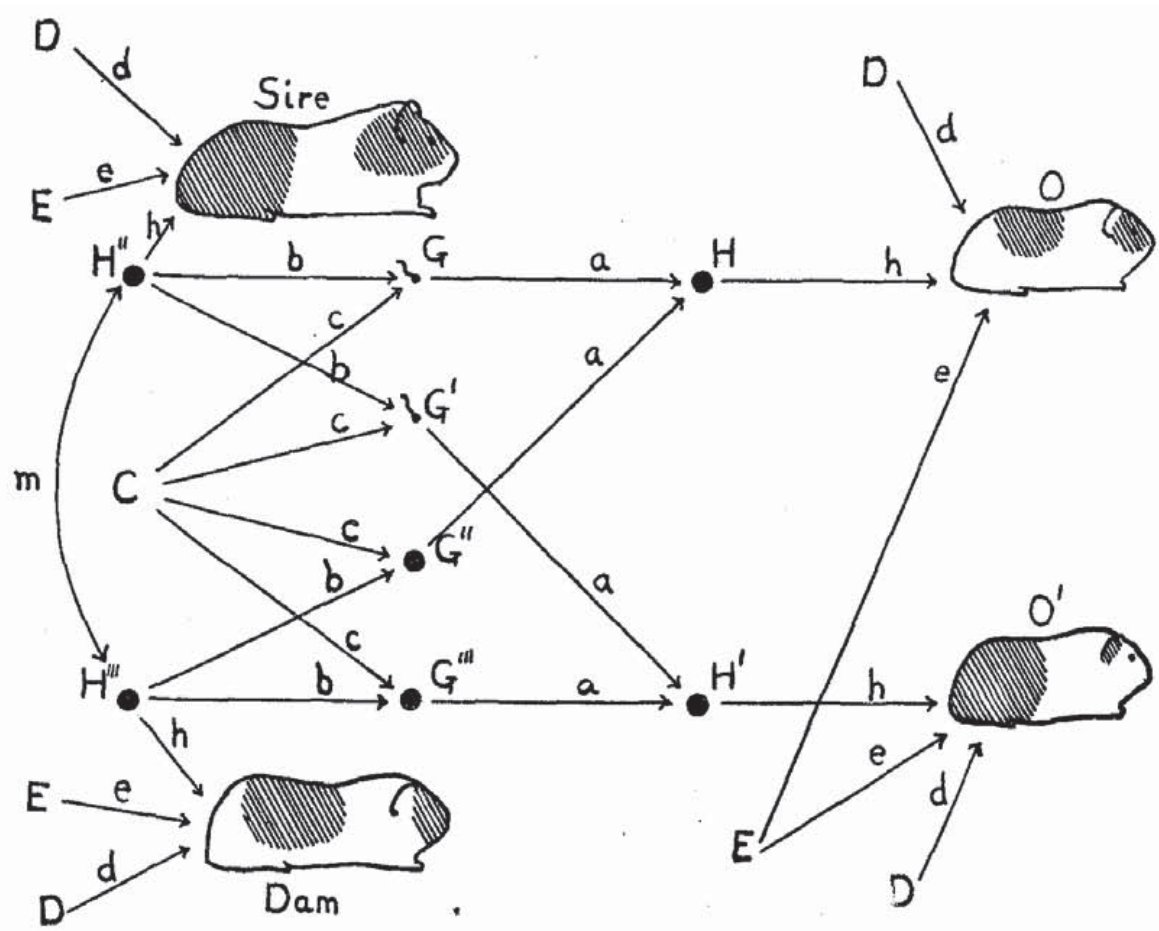


FIGURE 2.—A diagram illustrating the relations between two mated individuals and their progeny. H, H', H'' and H''' are the genetic constitutions of the four individuals. G, G', G'' and G''' are four germ-cells. E and D represent tangible external conditions and chance irregularities as factors in development. C represents chance at segregation as a factor in determining the composition of the germ-cells. Path coefficients are represented by small letters.

Figure 1. An early path diagram on the importance of heredity and environment in spotted guinea pigs from Sewall Wright (1921b).

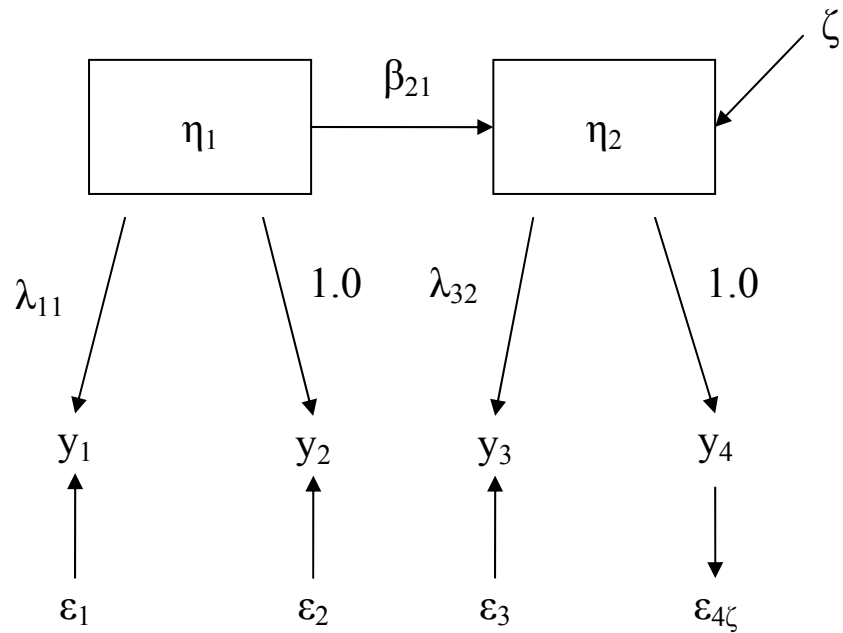


Figure 2. Path diagram of a walking dog model in four observables and two latent variables.

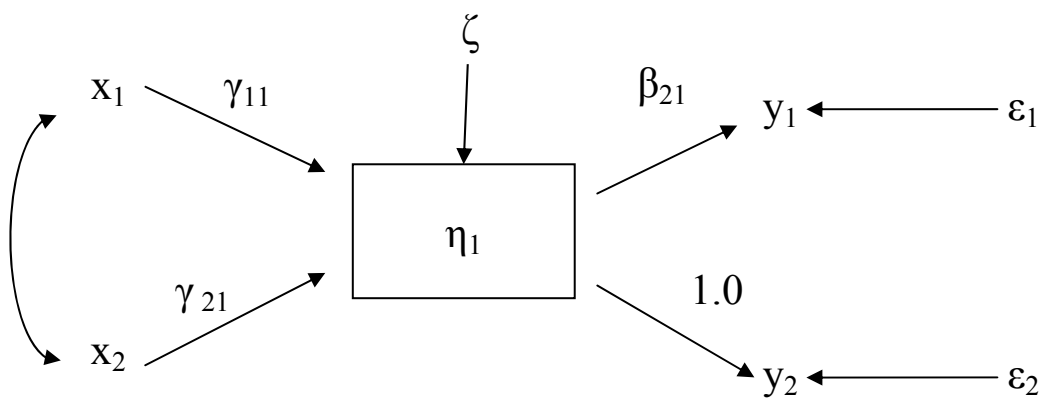


Figure 3. Path diagram of a multiple-indicator multiple-indicator cause (MIMIC) model.

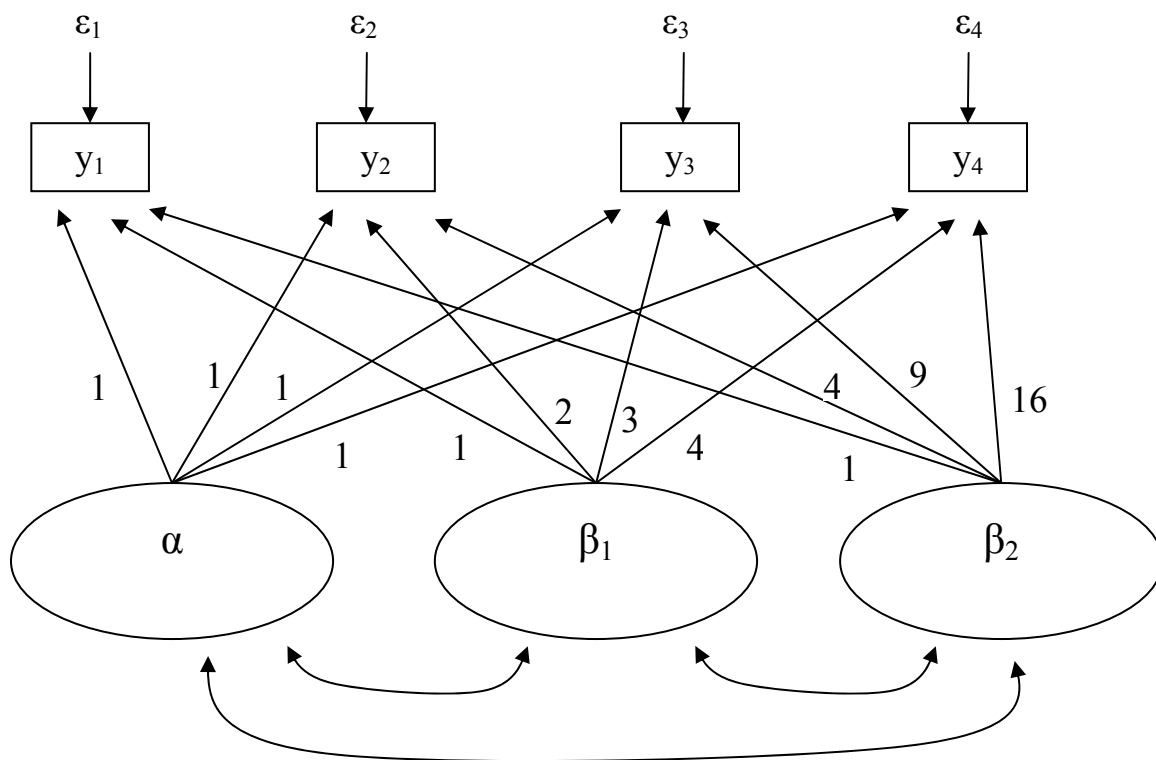


Figure 4. Path diagram of a four-wave quadratic latent curve model.

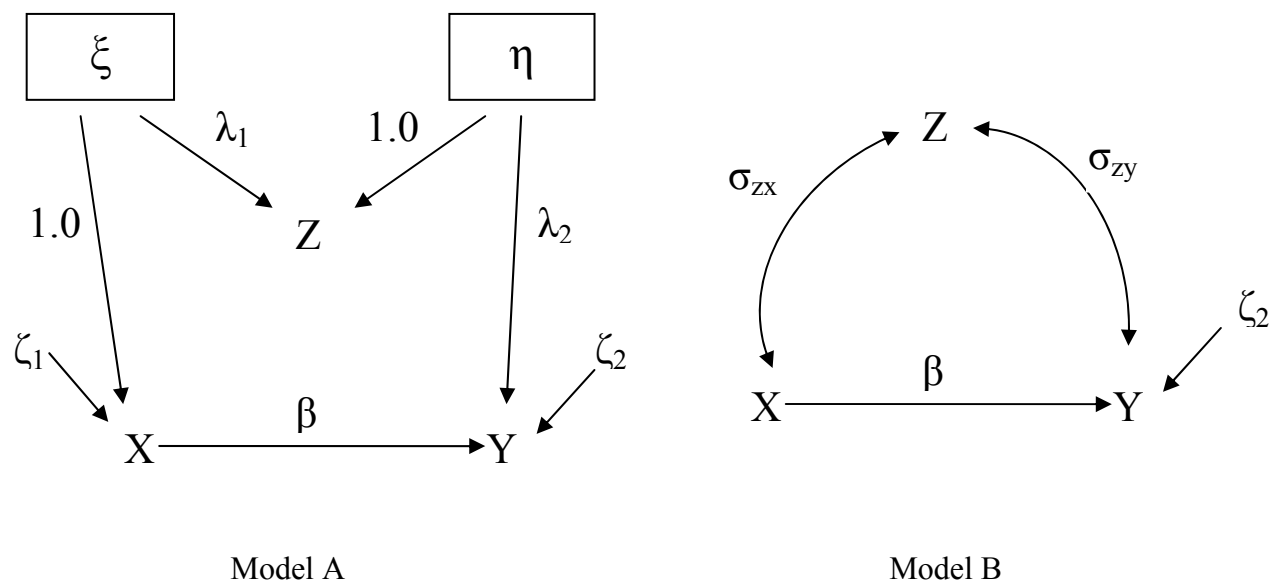


Figure 5. Path diagram of a model in which controlling for an exogenous variable creates bias.