

A closer look at fixed effects regression in structural equation modeling using **lavaan**

Henrik Kenneth Andersen

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Abstract

This article provides an in-depth look at fixed effects regression in the structural equation modeling (SEM) framework, specifically the application of fixed effects in the **lavaan** package for R. It is meant as a applied guide for researchers, covering the underlying model specification, syntax, and summary output. Online supplementary materials further discuss various common extentsions to the basic fixed-effect model, demonstrating how to relax model assumptions, deal with measurement error in both the dependent and independent variables, and include time-invariant predictors in a type of hybrid fixed-/ random effects model.

Keywords: Fixed effects, structural equation modeling, lavaan, R, panel analysis

1 Introduction

Several years ago, Curran and Bauer (2011) reflected positively on the growing use of panel studies in empirical social research. Some of the strengths of panel data are well-known, e.g., the ability to establish temporal precedence, increased statistical power and the reduction of potential alternative models. However, perhaps the greatest strength of panel data is that they allow for a more rigorous testing of substantive theories. Panel data, i.e., repeated measures of the same observed units (people, schools, firms, countries, etc.), allow researchers to decompose the error term into a part that stays constant within units and the part that changes over time. The part that does not change over time can be seen as the combined effect of all time-invariant influences (e.g., sex, date of birth, nationality) on the dependent variable. Fixed effects (FE) regression involves controlling for these time-invariant influences via a number of various methods. It thus drastically reduces the number of potential confounders of the relationship between variables of interest.

Structural equation modeling (SEM) is a popular regression framework. One of its main strengths is its flexibility. Not only can complex causal structures with multiple dependent variables be tested simultaneously, but in longitudinal (and, more generally, hierarchical) studies both time-varying and invariant predictors can be included, and effects can easily be allowed to vary over time. Thus researchers can allow for and study effects that increase or fade over time, or that appear only in specific periods. Beyond that, with the use of latent variables, SEM provides a way to deal with measurement error and get closer to the true underlying constructs of interest.

There are a number of articles describing basic concept of panel model regression, and FE regression in SEM (e.g., Allison 2011; Bollen and Brand 2010; Teachman et al. 2001). This article is intended as a *practical guide* for researchers looking for in-depth help with specifying FE models in SEM. It focuses on the `lavaan` (Rosseel 2012) package for R (R Core Team 2017). While `Mplus` (Muthén and

Muthén, [n.d.](#)) is arguably the most robust SEM software currently available (in terms of features like alignment, latent variable interactions, for example) , the `lavaan` package has many benefits. First, it and R are open source and completely free. For researchers dipping their toes into SEM, there is no financial barrier to try, and no risk if they decide it is not for them. Second, the implementation of `lavaan` in the larger R environment is an enormous advantage. Instead of poring over reams of plain text, copying out coefficients by hand, every part of the `lavaan` output is available as an object. This means that all aspects of the model, from fit indices, to coefficients and standard errors, to the model matrices, can be accessed and easily integrated into tables and plots. Furthermore, R can be used for a great deal of applications. It can be used to manage and manipulate as well as simulate data, perform symbolic algebra, run more traditional analyses (e.g., multiple regression, logistic regression, principal component analysis), etc. Once one is comfortable using R, there is no longer any need to switch between different software for data preparation and analysis.

The following article outlines the basic idea of FE regression, the particularities of FE in SEM, and shows its implementation in `lavaan`. Using [simulated data](#), it demonstrates and annotates the code for the most basic FE model and provides an overview of the summary output. A number of potential extensions to the basic model, including relaxing various assumptions, dealing with measurement error in both the independent and dependent variables, as well as the inclusion of time-invariant predictors in the form of a hybrid fixed-/ random effects model, are shown in detail in the form of [online supplementary materials](#).

2 Panel models

To begin, let us start by reviewing a general panel model (Bollen and Brand [2010](#)), also referred to as the ‘unobserved effects model’ (Wooldridge [2012](#)) (we will return to this model in the [online section](#) of the article when we discuss loosening assumptions)

$$y_{it} = \mathbf{x}_{it}\boldsymbol{\beta} + \mathbf{z}_i\boldsymbol{\gamma} + \alpha_i + \varepsilon_{it} \quad (1)$$

where y_{it} is the dependent variable for unit i , $i = 1, \dots, N$ at time t , $t = 1, \dots, T$, \mathbf{x}_{it} is a $1 \times K$ vector of time-varying covariates (which could include a constant) linked to the dependent variable by the $K \times 1$ vector of coefficients $\boldsymbol{\beta}$. \mathbf{z}_i is a $1 \times M$ vector of time-invariant covariates linked to the dependent variable by the $M \times 1$ vector of coefficients in $\boldsymbol{\gamma}$, α_i represents the combined effect of all unobserved unit-specific variables affecting the dependent variable and ε_{it} is the idiosyncratic error.

We can make stating some of the model assumptions easier by rewriting it in matrix notation

$$\mathbf{y}_i = \mathbf{X}_i\boldsymbol{\beta} + \mathbf{Z}_i\boldsymbol{\gamma} + \boldsymbol{\iota}_T\alpha_i + \boldsymbol{\varepsilon}_i$$

where \mathbf{y}_i and $\boldsymbol{\varepsilon}_i$ are $T \times 1$ vectors, \mathbf{X}_i and \mathbf{Z}_i are $T \times K$ and $T \times M$ matrices, respectively, $\boldsymbol{\iota}_T$ is a $T \times 1$ vector of ones and α_i is a scalar. $\boldsymbol{\beta}$ and $\boldsymbol{\gamma}$ are unchanged from Equation (1).

We typically make the following assumptions (see, e.g., Wooldridge 2002; Schmidheiny 2019) about this model:

- Linearity: the model is linear in its parameters and $\mathbb{E}[\varepsilon_{it}] = 0$ and $\mathbb{E}[\alpha_i] = 0$.
- Independence: the observations are independent across individuals (assured by random sampling in the cross-section), but not necessarily across time.
- Strict exogeneity: $\mathbb{E}[\varepsilon_{it}|\mathbf{X}_i, \mathbf{z}_i, \alpha_i] = 0$, i.e., the idiosyncratic error at each point in time is mean independent of \mathbf{X}_i at *all* points in time, $t = 1, \dots, T$, as well as \mathbf{z}_i and α_i .
- Error variance: homoscedastic and non-serially correlated errors, i.e., $\text{Var}(\varepsilon_i|\mathbf{X}_i, \mathbf{z}_i, \alpha_i) = \sigma_\varepsilon^2 \mathbf{I}$.

2.1 Random effects

If we are willing to make some assumptions about the distribution of the individual effects and, more importantly, their lack of a relationship with the observed time-varying covariates, we are left with the random effects model (Wooldridge 2002, 2012; Schmidheiny 2019; Skrondal and Rabe-Hesketh 2004). Namely, we assume the individual effects are normally distributed, with a constant (homoscedastic) variance conditional on the model covariates, i.e., $\alpha_i | \mathbf{X}_i, \mathbf{z}_i \sim N(0, \sigma_\alpha^2)$.¹ Further, we assume that the expectation of the individual effects conditional on the model covariates is zero, i.e., $\mathbb{E}[\alpha_i | \mathbf{X}_i, \mathbf{z}_i] = 0$, which implies they are also uncorrelated.

The random effects model works by defining a composite error term: $\nu_{it} = \alpha_i + \varepsilon_{it}$ and rewriting the model in Equation (1) as

$$y_{it} = \mathbf{x}_{it}\boldsymbol{\beta} + \mathbf{z}_i\boldsymbol{\gamma} + \nu_{it}, \text{ or} \\ \mathbf{y}_i = \mathbf{X}_i\boldsymbol{\beta} + \mathbf{Z}_i\boldsymbol{\gamma} + \boldsymbol{\nu}_i.$$

By adding the individual effects to the composite error in each time period, the composite error becomes serially correlated over time, to give the following conditional covariance matrix for the errors,

$$\text{Var}(\boldsymbol{\nu} | \mathbf{X}, \mathbf{z}) = \boldsymbol{\Omega}_{\boldsymbol{\nu}} = \begin{pmatrix} \boldsymbol{\Omega}_{\boldsymbol{\nu}_1} & \dots & 0 & \dots & 0 \\ \vdots & \ddots & \vdots & \ddots & \vdots \\ 0 & \dots & \boldsymbol{\Omega}_{\boldsymbol{\nu}_i} & \dots & 0 \\ \vdots & \ddots & \vdots & \ddots & \vdots \\ 0 & \dots & 0 & \dots & \boldsymbol{\Omega}_{\boldsymbol{\nu}_N} \end{pmatrix}$$

with typical elements

¹As long as \mathbf{X}_i contains a constant, the assumption that α_i has a mean of zero is unproblematic (Wooldridge 2012).

$$\text{Var}(\boldsymbol{\nu}_i | \mathbf{X}_i, \mathbf{z}_i) = \boldsymbol{\Omega}_{\nu_i} = \begin{pmatrix} \sigma_\nu^2 & \sigma_\alpha^2 & \dots & \sigma_\alpha^2 \\ \sigma_\alpha^2 & \sigma_\nu^2 & \dots & \sigma_\alpha^2 \\ \vdots & \vdots & \ddots & \vdots \\ \sigma_\alpha^2 & \sigma_\alpha^2 & \dots & \sigma_\nu^2 \end{pmatrix} \quad (2)$$

where $\sigma_\nu^2 = \sigma_\alpha^2 + \sigma_\varepsilon^2$. This means that in the conditional covariance matrix of the errors, given the time-varying and -invariant covariates, units over time will be correlated due to the individual effects. We should keep the covariance structure of the errors in mind as it will help make sense of the use of latent variables to decompose the dependent variable into between- and within-variance components, discussed below in Section 3.

2.2 Fixed effects

What if the individual effects are *not* independent of the observed time-varying covariates, i.e., $\mathbb{E}[\alpha_i | \mathbf{X}_i] \neq 0$? Then grouping them in with the error will cause $\boldsymbol{\beta}$ to be biased (Wooldridge 2002, 2012). In order to drop assumptions involving the individual effects, a number of methods are available (e.g., differencing, least squares dummy variable regression), but the most common approach is to *demean* the equation (Brüderl and Ludwig 2015). Demeaning involves subtracting the per-unit, over-time average from each of the model terms, i.e.,

$$\begin{aligned} (y_{it} - \bar{y}_i) &= (\mathbf{x}_{it} - \bar{\mathbf{x}}_i)\boldsymbol{\beta} + (\mathbf{z}_i - \bar{\mathbf{z}}_i)\boldsymbol{\gamma} + (\alpha_i - \bar{\alpha}_i) + (\varepsilon_{it} - \bar{\varepsilon}_i) \\ \ddot{y}_{it} &= \ddot{\mathbf{x}}_{it}\boldsymbol{\beta} + \ddot{\varepsilon}_{it}, \text{ or} \\ \ddot{\mathbf{y}}_i &= \ddot{\mathbf{X}}_i\boldsymbol{\beta} + \ddot{\boldsymbol{\varepsilon}}_i \end{aligned} \quad (3)$$

where, for example, $\bar{y}_i = T^{-1} \sum_{t=1}^T y_{it}$ (and the other over-time means are calculated analogously), and the variables with the dots above them represent the demeaned versions. Because the average of something that does not change is that thing itself, the individual effects, along with any time-invariant predictors, get wiped out by the demeaning. This means that no assumptions about the

relatedness of the model covariates and the unit-specific portion of the error are needed. The unbiasedness of the estimate is related solely to the strict exogeneity assumption imposed on the idiosyncratic errors, i.e., $\mathbb{E}[\tilde{\varepsilon}_{it}|\tilde{\mathbf{x}}_{it}] = \mathbb{E}[\tilde{\varepsilon}_{it}] = 0$ which also implies $\mathbb{E}[\tilde{\mathbf{x}}_{is}^\top \tilde{\varepsilon}_{it}] = \mathbf{0}$, $\forall s, t = 1, \dots, T$ (Bröderl and Ludwig 2015; Wooldridge 2002). The downside to this approach is that no time-invariant predictors can be included in the model. In the [online section](#), it will be discussed how to get around this restriction using SEM.

3 Fixed effects in structural equation modeling

Moving from the conventional methods outlined above to SEM, we must state the FE model in a different way. We turn to latent variables to account for time-invariant unobserved heterogeneity. In fact, besides accounting for measurement error and the representation of abstract hypothetical concepts, unobserved heterogeneity has historically been one of the main uses of latent variables in SEM (Skrondal and Rabe-Hesketh 2004).

3.1 Modeling time-invariant unobserved heterogeneity as a latent variable

We first need to convert the data from stacked, long-format vectors of length NT into T individual vectors of length N . To see why this is necessary, consider what effect this has on the vector of responses y_{it} . Let us, for a minute ignore any covariates and focus just on the dependent variable (a so-called ‘intercept-only’ model) so that we have $y_{it} = \alpha_i + \varepsilon_{it}$. When we convert the data to wide-format, we get T individual equations,

$$\begin{aligned} y_{1i} &= \alpha_i + \varepsilon_{1i} \\ y_{2i} &= \alpha_i + \varepsilon_{2i} \\ &\vdots \\ y_{Ti} &= \alpha_i + \varepsilon_{Ti}. \end{aligned} \tag{4}$$

Because the idiosyncratic errors are assumed to be uncorrelated across units and across time, the covariance between any two of the new wide vectors $\text{Cov}(y_{ti}, y_{si}) = \text{Var}(\alpha_i)$, $t \neq s$. Otherwise, when $t = s$, the covariance $\text{Cov}(y_{ti}, y_{ti}) = \text{Var}(\alpha_i) + \text{Var}(\varepsilon_{ti})$. This is the structure we saw above in a typical element of $\mathbf{\Omega}_\nu$.

And in fact this is exactly how a latent variable is used to account for time-invariant unobserved heterogeneity. The dependent variable at each timepoint is regressed onto the latent variable, see Figure 1. Here, the regression weights or ‘factor loadings’ are fixed to one to represent our assumption that the effect of the time-invariant unobserved heterogeneity is constant over time. It also means that the estimated variance of the latent variable is equal to the *average covariance between the wide-format columns of the dependent variable over time*. If $y_{it} = \alpha_i + \varepsilon_{it}$ is the true data generating process, then the relationship between two units over time is just α , regardless of the time distance. Referring back to the random effects structure of $\mathbf{\Omega}_{\nu_i}$ in Equality (2) for a generic unit i , we see the covariance on all of the off-diagonals is σ_α^2 . And, as we know, the average of something that does not change is that thing itself. I.e., if $T(T-1)/2 = h$ is the number of elements on either the upper- or lower triangle of $\mathbf{\Omega}_{\nu_i}$, then we have $h^{-1} \sum_{i=1}^h \sigma_\alpha^2 = \frac{h\sigma_\alpha^2}{h} = \sigma_\alpha^2$.

To show this, consider the following matrix equation of the variances and the nonredundant covariances in a three-wave intercept-only model that follows directly from Equation (4) (assuming $\text{Cov}(\varepsilon_{ti}, \varepsilon_{si}) = 0$, $t \neq s$), and which we can solve easily with least squares:

$$\mathbf{b} = \mathbf{A}\mathbf{x}$$

$$\begin{pmatrix} \text{Var}(y_1) \\ \text{Cov}(y_2, y_1) \\ \text{Cov}(y_3, y_1) \\ \text{Var}(y_2) \\ \text{Cov}(y_3, y_2) \\ \text{Var}(y_3) \end{pmatrix} = \begin{pmatrix} \text{Var}(y_1) \\ \text{Cov}(y_2, y_1) \\ \text{Cov}(y_3, y_1) \\ \text{Var}(y_2) \\ \text{Cov}(y_3, y_2) \\ \text{Var}(y_3) \end{pmatrix} \begin{pmatrix} \psi & \phi_1 & \phi_2 & \phi_3 \\ 1 & 1 & 0 & 0 \\ 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 \\ 1 & 0 & 1 & 0 \\ 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 1 \end{pmatrix} \begin{pmatrix} \psi \\ \phi_1 \\ \phi_2 \\ \phi_3 \end{pmatrix}$$

where $\psi = \text{Var}(\alpha)$, $\phi_t = \text{Var}(\varepsilon_t)$. We can solve this equation to show

$$\begin{aligned} \mathbf{b} &= \mathbf{A}\mathbf{x} \\ (\mathbf{A}^\top \mathbf{A})^{-1} \mathbf{A}^\top \mathbf{b} &= (\mathbf{A}^\top \mathbf{A})^{-1} \mathbf{A}^\top \mathbf{A}\mathbf{x} \\ (\mathbf{A}^\top \mathbf{A})^{-1} \mathbf{A}^\top \mathbf{b} &= \mathbf{x} \\ \begin{pmatrix} .33 \text{Cov}(y_2, y_1) + .33 \text{Cov}(y_3, y_1) + .33 \text{Cov}(y_3, y_2) \\ \text{Var}(y_1) - \psi \\ \text{Var}(y_2) - \psi \\ \text{Var}(y_3) - \psi \end{pmatrix} &= \begin{pmatrix} \psi \\ \phi_1 \\ \phi_2 \\ \phi_3 \end{pmatrix}. \end{aligned}$$

This shows that if our assumption of constant covariances between the dependent variable across time holds, the estimated variance of α is just what it should be: the average covariance between units of y over time. Once we add in observed covariates, the estimated covariance of α then become the *conditional* covariance of y over time, given those covariates.

3.2 Model notation

Having explained how a latent variable is used to estimate the individual effects, we need to state the FE model using SEM-compatible matrix notation. There are a number of different model notations (see, for example Bollen (1989) for an overview), but the one that will serve us best is one that was proposed by Graff (1979):

$$\begin{aligned}\mathbf{y}^+ &= \mathbf{\Lambda}_y^+ \boldsymbol{\eta}^+, \\ \boldsymbol{\eta}^+ &= \mathbf{B} \boldsymbol{\eta}^+ + \boldsymbol{\zeta}^+, \end{aligned}$$

where $\boldsymbol{\eta}^+ = (\mathbf{y}, \mathbf{x}, \boldsymbol{\eta}, \boldsymbol{\xi})^\top$, $\boldsymbol{\zeta}^+ = (\boldsymbol{\varepsilon}, \boldsymbol{\delta}, \boldsymbol{\zeta}, \boldsymbol{\xi})^\top$, $\mathbf{y}^+ = (\mathbf{y}, \mathbf{x})^\top$. \mathbf{y} is a vector of observed dependent variables and \mathbf{x} is a vector of observed independent variables. $\boldsymbol{\eta}$ is a vector of the latent dependent variables and $\boldsymbol{\xi}$ is a vector of latent independent variables. $\boldsymbol{\varepsilon}$ and $\boldsymbol{\delta}$ are vectors of the errors of the observed dependent and independent variables, respectively, and $\boldsymbol{\zeta}$ is a vector of the errors, or disturbances, of the latent variables. Notice the $^+$ symbol is just meant to differentiate the vectors with them from those without them. That means, $\boldsymbol{\eta}^+$ is a vector that holds the observed and latent variables, both dependent (in SEM they are referred to as ‘endogenous’) and independent (i.e., ‘exogenous’)², $\boldsymbol{\zeta}^+$ holds the errors for the observed variables and the disturbances of the latent variables. \mathbf{y}^+ holds just the observed variables, both dependent and independent, and $\mathbf{\Lambda}_y^+$ is a matrix of ones and zeros that selects the observed variables from $\boldsymbol{\eta}^+$. Lastly \mathbf{B} is a matrix that holds the regression coefficients.

If we say that p and q stand for the number of observed dependent and independent variables, respectively, and m and n stand for the number of latent dependent and independent variables, respectively, then $\boldsymbol{\eta}^+$ and $\boldsymbol{\zeta}^+$ are $p+q+m+n$, \mathbf{y}^+ is $p+q$, $\mathbf{\Lambda}_y^+$ is $(p+q) \times (p+q+m+n)$ and \mathbf{B} is $(p+q+m+n) \times (p+q+m+n)$ (Bollen 1989).

This notation may be confusing at first, but it has advantages. First, it allows us the flexibility we need for the models. For example, it allows observed \mathbf{x} to directly influence observed \mathbf{y} (more common notation assumes that substantive effects occur only between latent variables, observed ones are only used as indicators, see for example Bollen (1989), Kline (2016)). It also allows $\boldsymbol{\xi}$, i.e., any latent exogenous variables, to influence \mathbf{y} directly. These two scenarios cover the

²For our purposes, the terms endogenous and dependent, on the one hand, and exogenous and independent, on the other, can be used interchangeably.

traditional FE model with observed variables, and one in which latent variables are used to account for measurement error in the independent variables. It is also consistent with the notation used for these models in `lavaan`. In fact, `lavaan` switches automatically between matrix notations depending on the specified model. That means the matrix representation of the model one sees if they type in `lavInspect(model, what = "est")` after specifying their model in `lavaan` will match up with the notation used here.³ It does have a potential disadvantage however. Besides being less intuitive than the typical $\boldsymbol{\eta} = \mathbf{B}\boldsymbol{\eta} + \mathbf{\Gamma}\boldsymbol{\xi} + \boldsymbol{\zeta}$ notation, it means that by including the observed covariates in the stacked long vector \mathbf{y}^+ , they are treated as another response variable with variances and covariances to be estimated by the model (instead of just using the sample statistics). This means that the assumption of multivariate normality (otherwise just imposed on the dependent variables) also applies to the independent ones (Skrondal and Rabe-Hesketh 2004, 75). This can be problematic for noncontinuous independent variables like sex, nationality dummies, marriage status (married/unmarried), etc. See Skrondal and Rabe-Hesketh (2004) for more on this topic.

Let us, however, make things more concrete and take a look at a simple, three-wave version of the typical FE-SEM using this notation (shown graphically in Figure 1). For that, we have:

$$\mathbf{y}^+ = \mathbf{\Lambda}_y^+ \boldsymbol{\eta}^+ \quad (5)$$

$$\begin{pmatrix} y_1 \\ y_2 \\ y_3 \\ x_1 \\ x_2 \\ x_3 \end{pmatrix} = \begin{matrix} y_1 & y_2 & y_3 & x_1 & x_2 & x_3 & \alpha \\ y_1 & \begin{pmatrix} 1 & 0 & 0 & 0 & 0 & 0 & 0 \end{pmatrix} \\ y_2 & \begin{pmatrix} 0 & 1 & 0 & 0 & 0 & 0 & 0 \end{pmatrix} \\ y_3 & \begin{pmatrix} 0 & 0 & 1 & 0 & 0 & 0 & 0 \end{pmatrix} \\ x_1 & \begin{pmatrix} 0 & 0 & 0 & 1 & 0 & 0 & 0 \end{pmatrix} \\ x_2 & \begin{pmatrix} 0 & 0 & 0 & 0 & 1 & 0 & 0 \end{pmatrix} \\ x_3 & \begin{pmatrix} 0 & 0 & 0 & 0 & 0 & 1 & 0 \end{pmatrix} \end{matrix} \begin{pmatrix} y_1 \\ y_2 \\ y_3 \\ x_1 \\ x_2 \\ x_3 \\ \alpha \end{pmatrix},$$

³In `Mplus`, the model matrices can be requested by including `OUTPUT: TECH1` in the input file.

$$\boldsymbol{\eta}^+ = \mathbf{B}\boldsymbol{\eta}^+ + \boldsymbol{\zeta}^+$$

$$\begin{pmatrix} y_1 \\ y_2 \\ y_3 \\ x_1 \\ x_2 \\ x_3 \\ \alpha \end{pmatrix} = \begin{matrix} y_1 \\ y_2 \\ y_3 \\ x_1 \\ x_2 \\ x_3 \\ \alpha \end{matrix} \begin{pmatrix} y_1 & y_2 & y_3 & x_1 & x_2 & x_3 & \alpha \\ 0 & 0 & 0 & \beta & 0 & 0 & 1 \\ 0 & 0 & 0 & 0 & \beta & 0 & 1 \\ 0 & 0 & 0 & 0 & 0 & \beta & 1 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 \end{pmatrix} \begin{pmatrix} y_1 \\ y_2 \\ y_3 \\ x_1 \\ x_2 \\ x_3 \\ \alpha \end{pmatrix} + \begin{pmatrix} \varepsilon_1 \\ \varepsilon_2 \\ \varepsilon_3 \\ x_1 = \delta_1 \\ x_2 = \delta_2 \\ x_3 = \delta_3 \\ \alpha = \xi \end{pmatrix}. \quad (6)$$

Notice in $\boldsymbol{\zeta}^+$, for the independent variables we could either write, for example x_t or δ_t . As mentioned above, this is due to the model notation treating the independent variables like dependent variables with variances/covariances to be estimated. For the sake of simplicity, we will ignore this subtlety and refer to the observed variable from now on, keeping in mind that if the multivariate normality assumption holds, the estimated statistics will likely be sufficiently close to the sample ones for it to not make much of a difference.

Admittedly, Equations (5) and (6) may not look like much yet. We can remedy this by first putting the equation for $\boldsymbol{\eta}^+$ in reduced form, i.e., by getting rid of the dependent variable on the r.h.s.:

$$\begin{aligned} \boldsymbol{\eta}^+ &= \mathbf{B}\boldsymbol{\eta}^+ + \boldsymbol{\zeta}^+ \\ \boldsymbol{\eta}^+ - \mathbf{B}\boldsymbol{\eta}^+ &= \boldsymbol{\zeta}^+ \\ (\mathbf{I} - \mathbf{B})\boldsymbol{\eta}^+ &= \boldsymbol{\zeta}^+ \\ \boldsymbol{\eta}^+ &= (\mathbf{I} - \mathbf{B})^{-1}\boldsymbol{\zeta}^+, \end{aligned}$$

where \mathbf{I} is the identity matrix. By substituting this back into the equation for the observed variables we get $\mathbf{y}^+ = \boldsymbol{\Lambda}_y^+[(\mathbf{I} - \mathbf{B})^{-1}\boldsymbol{\zeta}^+]$, which works out to:

$$\begin{pmatrix} y_1 \\ y_2 \\ y_3 \\ x_1 \\ x_2 \\ x_3 \end{pmatrix} = \begin{pmatrix} \alpha + \beta x_1 + \varepsilon_1 \\ \alpha + \beta x_2 + \varepsilon_2 \\ \alpha + \beta x_3 + \varepsilon_3 \\ x_1 \\ x_2 \\ x_3 \end{pmatrix},$$

which is of course exactly what we should expect given Equation (3)⁴.

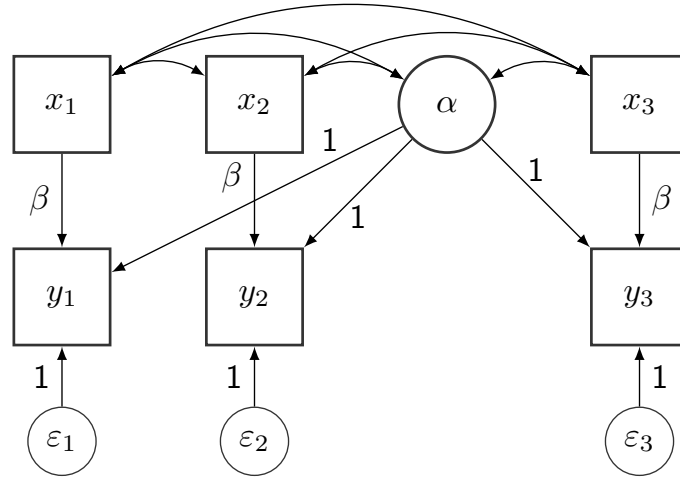


Figure 1: Typical three-wave FE-SEM model with contemporary effects

3.3 Assumptions

What essentially differentiates an FE from a random effects (RE) model is our assumption concerning the relationship between the unobserved individual effects and the model covariates (Bollen and Brand 2010). The FE model assumes that $\mathbb{E}[\alpha x_t] \neq 0$. As such, if we fail to control for the correlation of the covariate and the time-invariant part of the error, then the coefficient of interest, here β , will be biased. Our assumption regarding whether the individual effects are correlated with the model covariates occurs in $\mathbb{E}[\zeta^+ \zeta^{+\top}] = \Psi$, the covariance matrix of the errors

⁴We can use the `sympy` package in `python` to verify and show the steps for this and other examples, see the [supplementary materials](#) for the code.

$$\begin{aligned}
\mathbf{y}^+ \mathbf{y}^{+\top} &= \mathbb{E}[(\mathbf{\Lambda}_y^+ (\mathbf{I} - \mathbf{B})^{-1} \boldsymbol{\zeta}^+) (\mathbf{\Lambda}_y^+ (\mathbf{I} - \mathbf{B})^{-1} \boldsymbol{\zeta}^+)^{\top}] \\
&= \mathbb{E}[(\mathbf{\Lambda}_y^+ (\mathbf{I} - \mathbf{B})^{-1} \boldsymbol{\zeta}^+) (\boldsymbol{\zeta}^{+\top} (\mathbf{I} - \mathbf{B})^{-1\top} \mathbf{\Lambda}_y^{+\top})] \\
&= \mathbf{\Lambda}_y^+ (\mathbf{I} - \mathbf{B})^{-1} \mathbb{E}[\boldsymbol{\zeta}^+ \boldsymbol{\zeta}^{+\top}] (\mathbf{I} - \mathbf{B})^{-1\top} \mathbf{\Lambda}_y^{+\top} \\
&= \mathbf{\Lambda}_y^+ (\mathbf{I} - \mathbf{B})^{-1} \boldsymbol{\Psi} (\mathbf{I} - \mathbf{B})^{-1\top} \mathbf{\Lambda}_y^{+\top}.
\end{aligned}$$

In the case of an FE model, $\boldsymbol{\Psi}$ will reflect our belief that the individual effects are correlated with the model covariates, here again for demonstration the three-wave model:

$$\boldsymbol{\Psi} = \begin{matrix} & \begin{matrix} \varepsilon_1 & \varepsilon_2 & \varepsilon_3 & x_1 & x_2 & x_3 & \alpha \end{matrix} \\ \begin{matrix} \varepsilon_1 \\ \varepsilon_2 \\ \varepsilon_3 \\ x_1 \\ x_2 \\ x_3 \\ \alpha \end{matrix} & \begin{pmatrix} \varepsilon_1^2 & & & & & & \\ 0 & \varepsilon_2^2 & & & & & \\ 0 & 0 & \varepsilon_3^2 & & & & \\ 0 & 0 & 0 & x_1^2 & & & \\ 0 & 0 & 0 & x_2 x_1 & x_2^2 & & \\ 0 & 0 & 0 & x_3 x_1 & x_3 x_2 & x_3^2 & \\ 0 & 0 & 0 & \alpha x_1 & \alpha x_2 & \alpha x_3 & \alpha^2 \end{pmatrix} \end{matrix}.$$

Knowing this, we can work out the equation for the coefficient of interest, β . For the sake of simplicity, assume here and throughout mean-centered variables:

$$\begin{aligned}
\text{Cov}(y_t, x_t) &= \mathbb{E}[y_t x_t] \\
&= \mathbb{E}[(\alpha + \beta x_t + \varepsilon_t) x_t] \\
&= \mathbb{E}[\alpha x_t + \beta x_t^2 + \varepsilon_t x_t] \\
&= \text{Cov}(\alpha, x_t) + \beta \text{Var}(x_t) \\
\hat{\beta} &= \frac{\text{Cov}(y_t, x_t) - \text{Cov}(\alpha, x_t)}{\text{Var}(x_t)}.
\end{aligned}$$

This should make intuitive sense. From the observed covariance between the dependent and the independent variable, we are partialling out the part that is due to the covariance between the independent variable and the individual effects per unit, and then dividing by the variance of the independent variable, as usual. For the RE model, we assume $\mathbb{E}[\alpha x_t] = 0$ and the equation reduces to $\hat{\beta} = \text{Cov}(y_t, x_t) / \text{Var}(x_t)$. The rest of the model-implied covariance matrix

results from $\mathbf{y}^+ \mathbf{y}^{+\top}$.

4 Fixed effects in lavaan

The package `lavaan` needs to be installed once with `install.packages("lavaan")`.

To be able to use it, we need to load it for every new R session:

```
library( lavaan)
```

For users unfamiliar with R, SEM analyses can be carried out with almost no knowledge of the language. Typically, someone unfamiliar with R would prepare their data using some other statistical software, and then save the intended dataset as a `.csv`, `.xlsx`, `.dta`, `.sav`, etc. file. The user must then import the data, preferably as a dataframe, and the rest occurs using the `lavaan` syntax.⁵

Specifying the most basic fixed effects model, like the one shown in Bollen and Brand (2010) (the same model as Equation (3) but with just one time-varying predictor) involves four components. First, we define the latent individual effects variable using the `=~` ‘measured by’ or ‘manifested by’ (Rosseel 2012) operator at the same time constraining the factor loadings at each timepoint to one. I will call the latent variable `a` to stand for α . Constraining all of the factor loadings to one reflects our implicit assumption that the combined effect of the unit-specific unobserved factors is constant over time. This is the default behaviour of traditional POLS-based approaches to FE that use the stacked long-format data.

```
a =~ 1*y1 + 1*y2 + 1*y3 + 1*y4 + 1*y5
```

Second, we regress the dependent variable on the independent variable using the `~` regression operator. With stacked, long-format data, only one regression coefficient is estimated over all observed timepoints. To have our FE-SEM model mimic this behaviour, we need to constrain the the estimated coefficient to equal

⁵There are many online tutorials for importing data in various formats, see, for example some from [datacamp](#) or [Quick-R](#), or any of the many posts on [stackoverflow](#).

over time. We do so by adding the same label to the regression coefficient at every time point. We will use the label `b` (this label was chosen arbitrarily, we could have used any letter or string of characters) and have it act as an equality constraint for the regression coefficient of interest β :

```
y1 ~ b*x1
y2 ~ b*x2
y3 ~ b*x3
y4 ~ b*x4
y5 ~ b*x5
```

The key to a FE model, as opposed to an RE model are our assumptions about the relatedness of our covariate and the individual effects, i.e., $\mathbb{E}[x_t\alpha]$. For an FE model, we want to partial out any potential covariance between the independent variable and the individual effects. This accounts for any linear relationship between x_t and the unit-specific characteristics influencing the dependent variable. Further, allowing unrestricted covariances between the independent variable itself over time will not affect how the coefficient β is estimated, but will have an effect on the standard errors. To mimic the behaviour of a conventional FE model, we allow the independent variable to be correlated with the individual effects and itself over time. Covariances (including covariances between a variable and itself, i.e., variances) are specified using the `~~` operator:

```
a ~~ x1 + x2 + x3 + x4 + x5
x1 ~~ x2 + x3 + x4 + x5
x2 ~~ x3 + x4 + x5
x3 ~~ x4 + x5
x4 ~~ x5
```

The last component of our code involves the variances of the residuals. This component is optional, but we can constrain the residual variances to be equal over time to again mimic the behaviour of a conventional FE model using POLS on stacked data. Here, again, we use labels to make equality constraints. Because y_t is endogenous, the `~~` operator specifies the variances of *residuals*, i.e. ϵ_t .


```

y1 ~~ e*y1
y2 ~~ e*y2
y3 ~~ e*y3
y4 ~~ e*y4
y5 ~~ e*y5

```

5 A simulated example

To demonstrate the application of FE models in SEM, a dataset can be simulated that embodies the FE assumptions. Again, the code for data simulation can be found in the [online supplementary materials](#).

To show that the latent individual effects variables represent the *combined* effect of all time-invariant characteristics, the dependent variable will be influenced by two separate unit-specific variables, which we can call α_1 and α_2 . We will construct the simulated data such that the independent variable is correlated with both of the time-invariant variables. This means that approaches that fail to account for this confounding influence, such as pooled ordinary least squares (POLS) or RE, will be biased.

The wide-format equations for the data generating process can be described as:

$$\begin{aligned}
\mathbf{x}_t &= \boldsymbol{\alpha}_1 \beta_{x_t, \alpha_1} + \boldsymbol{\alpha}_2 \beta_{x_t, \alpha_2} + \boldsymbol{\delta}_t, \\
\mathbf{y}_t &= \mathbf{x}_t \beta_{y_t, x_t} + \boldsymbol{\alpha}_1 \beta_{y_t, \alpha_1} + \boldsymbol{\alpha}_2 \beta_{y_t, \alpha_2} + \boldsymbol{\varepsilon}_t
\end{aligned}$$

where, for the sake of simplicity, $\boldsymbol{\alpha}_1$, $\boldsymbol{\alpha}_2$, $\boldsymbol{\delta}_t$ and $\boldsymbol{\varepsilon}_t$ are $\sim N(0, 1)$.

For the following example, a sample size of 1,000, observed over five waves, was chosen. The unique variance of \mathbf{x} , as well as both the individual-effect variables is also $\sim N(0, 1)$. The coefficient of interest, $\beta_{y, x}$ is set to be equal to 0.3. A correlation between \mathbf{x} and the individual effects is induced through $\beta_{x, \alpha_1} = 0.85$ and $\beta_{x, \alpha_2} = 0.50$. With the variances above set to one, the covariances will be roughly $\text{Cov}(x_t, \alpha_1) = 0.85$ and $\text{Cov}(x_t, \alpha_2) = 0.5$. The dependent variable

is also influenced by the individual effects variables with $\beta_{y_t, \alpha_1} = 0.75$ and $\beta_{y_t, \alpha_2} = 0.45$. These values were chosen arbitrarily.

Now, we run the FE-SEM in `lavaan`.

```
fe_sem <- '  
# Define individual effects variable  
a =~ 1*y1 + 1*y2 + 1*y3 + 1*y4 + 1*y5  
# Regressions, constrain coefficient to be equal over time  
y1 ~ b*x1  
y2 ~ b*x2  
y3 ~ b*x3  
y4 ~ b*x4  
y5 ~ b*x5  
# Allow unrestricted correlation between eta and covariates  
a ~~ x1 + x2 + x3 + x4 + x5  
x1 ~~ x2 + x3 + x4 + x5  
x2 ~~ x3 + x4 + x5  
x3 ~~ x4 + x5  
x4 ~~ x5  
# Constrain residual variances to be equal over time  
y1 ~~ e*y1  
y2 ~~ e*y2  
y3 ~~ e*y3  
y4 ~~ e*y4  
y5 ~~ e*y5  
'  
fe_sem.fit <- sem(model = fe_sem,  
                  data = dfw,  
                  estimator = "ML")
```

We can get a summary of the model with `summary()`. The first portion of the summary output gives an overview of some basic information and fit statistics. The maximum likelihood estimator is the default, so it did not have to be explicitly selected in the fitting function call. Other estimators are available, including generalized and unweighted least squares (GLS and ULS, respectively), robust standard errors maximum likelihood (MLM) and several others (see [the lavaan online tutorial for more](#)).

This part of the summary output also tells us that the analysis is based on 1,000 observations (missings would be shown here as well if there were any), and that

the χ^2 statistic is 30.138 based on 32 degrees of freedom (55 observed covariances minus 1 error variance, 1 coefficient, 1 latent variable variance, 5 exogenous variable variances and 15 covariances for $55 - 23 = 32$ df). The p-value on the χ^2 statistic is not significant with $p = 0.561$ which tells us the differences between the model-implied and observed covariance matrices are likely due to chance, and that the model fits the data well (given how the data was generated, it would be surprising if this were not the case). Other fit measures including typical comparative fit indices can be requested by either adding `fit.measures = TRUE` as a secondary argument to the `summary()` call, or by asking for a complete list of all available fit statistics using `lavInspect(model, "fit")` where `model` stands for the name of the fitted model, in this case `fe_sem.fit`.

```
summary(fe_sem.fit)

## lavaan 0.6-6 ended normally after 37 iterations
##
##      Estimator                      ML
##      Optimization method          NLMINB
##      Number of free parameters      31
##      Number of equality constraints    8
##
##      Number of observations          1000
##
## Model Test User Model:
##
##      Test statistic                  30.138
##      Degrees of freedom              32
##      P-value (Chi-square)            0.561
##
## Parameter Estimates:
##
##      Standard errors                  Standard
##      Information                      Expected
##      Information saturated (h1) model  Structured
...
```

Next the summary output shows the measurement models for the latent variables, if any. In this case the latent variable `a` for α is measured by each of the five observed dependent variables with factor loadings fixed to 1.0.

...

```
## Latent Variables:
##           Estimate Std.Err z-value P(>|z|)
##    a =~
##      y1           1.000
##      y2           1.000
##      y3           1.000
##      y4           1.000
##      y5           1.000
...

```

The regressions are shown next. Here, because we have constrained the regression coefficients to be equal over time (the equality constraint label (b) is listed to the left of the estimates), the estimate of $\beta = 0.294$ (0.016) is repeated five times. The corresponding z- and p-values show that the coefficient is, unsurprisingly, significant.

```
...
## Regressions:
##           Estimate Std.Err z-value P(>|z|)
##    y1 ~
##      x1      (b)    0.294    0.016   18.809    0.000
##    y2 ~
##      x2      (b)    0.294    0.016   18.809    0.000
##    y3 ~
##      x3      (b)    0.294    0.016   18.809    0.000
##    y4 ~
##      x4      (b)    0.294    0.016   18.809    0.000
##    y5 ~
##      x5      (b)    0.294    0.016   18.809    0.000
...

```

Next, the covariance estimates are listed. First, the covariances between the latent individual effects variable and the independent variable over time are shown, and then the covariances between the independent variable with itself over time.

One should always take care to double-check that there are no unintended covariances listed here. Like `Mplus`, the `lavaan` package estimates some covariances per default, without the user explicitly having to add them to the model syntax. For example, covariances between latent variables are estimated per default. If one does not wish for them to covary, it must be explicitly stated, e.g., with `f1`

~~ 0*f2, assuming the latent variables are called f1 and f2, or by overriding the default behaviour for the entire model by adding `orthogonal = TRUE` (which sets the correlation between all latent variables to zero) to the fitting call.⁶

```
...
## Covariances:
##           Estimate Std.Err z-value P(>|z|)
## a ~~
## x1           0.844   0.055  15.355   0.000
## x2           0.867   0.056  15.441   0.000
## x3           0.845   0.055  15.400   0.000
## x4           0.822   0.053  15.455   0.000
## x5           0.820   0.053  15.572   0.000
## x1 ~~
## x2           0.908   0.070  12.900   0.000
## x3           0.935   0.069  13.466   0.000
## x4           0.921   0.067  13.661   0.000
## x5           0.914   0.067  13.716   0.000
## x2 ~~
## x3           0.889   0.070  12.675   0.000
## x4           0.922   0.069  13.423   0.000
## x5           0.889   0.068  13.165   0.000
## x3 ~~
## x4           0.865   0.067  12.976   0.000
## x5           0.901   0.066  13.554   0.000
## x4 ~~
## x5           0.850   0.064  13.285   0.000
...
```

Finally, the variance estimates are listed. Here, we see that in order to mimic the behaviour of a traditional FE model, the error variances over time were specified to be equal using the equality constraint (e). Notice the . beside y1, y2, etc.: this indicates that the listed variance refers to an endogenous variable, and that it is thus an error variance. In this case, these refer to the variances of ε_t . After that, the variances of the exogenous variables, both observed and unobserved are listed.

```
...
## Variances:
##           Estimate Std.Err z-value P(>|z|)
## .y1      (e)    1.022   0.023  44.721   0.000
## .y2      (e)    1.022   0.023  44.721   0.000
```

⁶This is at least the current behaviour of both the `cfa` and `sem` wrappers. In fact, both wrappers seem to be identical in terms of the default settings, see Rosseel et al. (2020).

##	.y3	(e)	1.022	0.023	44.721	0.000
##	.y4	(e)	1.022	0.023	44.721	0.000
##	.y5	(e)	1.022	0.023	44.721	0.000
##	x1		1.986	0.089	22.361	0.000
##	x2		2.079	0.093	22.361	0.000
##	x3		1.987	0.089	22.361	0.000
##	x4		1.860	0.083	22.361	0.000
##	x5		1.814	0.081	22.361	0.000
##	a		0.799	0.052	15.310	0.000

6 Conclusion

Fixed effects regression in SEM has been outlined in well-known articles by (Allison 2011; Bollen and Brand 2010; Teachman et al. 2001). This article provides a focused look at the implementation of the basic model, as well as common extensions using the `lavaan` package in R.

The benefits of FE-SEM as opposed to traditional OLS-based FE-models are largely the same ones that apply to the SEM framework in general: for one, SEM allows for a great deal of flexibility. For example, it is easy to loosen model constraints as necessary. Measurement error in both the dependent and independent variables can be dealt with using latent variables to achieve unbiased and more efficient results. Researchers interested in time-invariant predictors can integrate them into a hybrid FE/RE model with ease. Further extensions, like measurement invariance testing (Schout, Lugtig, and Hox 2012; Millsap 2011; Steenkamp and Baumgartner 1998) as well as lagged dependent variables (Bollen and Brand 2010; Allison, Williams, and Moral-Benito 2017) for example, can also be implemented in a straightforward fashion.

The most basic FE-SEM is furthermore the basis for a variety of currently popular extended models, such as Latent Curve Models in general (Curran and Bollen 2001; Bollen and Curran 2004), as well as special implementations like the Dynamic Panel Model (Allison, Williams, and Moral-Benito 2017), the Random-Intercept Cross-Lagged Panel Model (Hamaker, Kuiper, and Grasman 2015) and the Latent Curve Model with Structured Residuals (Curran et al.

2014). For this reason it is all the more important for researchers to have a good grasp on the method of applying FE-SEM. This article is meant to serve as a consolidated resource for researchers looking for concrete advice on specifying FE and more general panel models in SEM.

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