

Unified Framework Notation

Manuel Arnold

Humboldt-Universität zu Berlin

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This document is intended to facilitate a better understanding of the code used in the backend. Terminology and notation are not meant to be used in the frontend or in the paper.

Following Usami et al., I limit the scope to single-indicator models. Furthermore, I assume stationarity meaning that all models parameters are time-invariant.

I specify the unified framework in reticular action model (RAM) notation. In RAM notation, SEMs are described by a symmetrical \mathbf{S} matrix of dimensions $V \times V$, an asymmetrical matrix \mathbf{A} of dimensions $V \times V$, a filter matrix \mathbf{F} of dimensions $M \times V$, and a mean vector \mathbf{m} of length V , where M denotes the number of manifest variables and V the total number of variables including latent variables from the model.

Usami et al. use a element-wise notation of the unified framework. In the following, I will convert to Usami et al.'s notation to matrix notation so that is easier for programming. First, we look at the dimensions of the RAM matrices. Let \mathbf{y}_t be a J -variate vector of manifest variables that are measured at the time points $t = 1 \dots, T$. Thus, the total number of manifest variables is $JT = V$. The unified framework contains the following latent variables: \mathbf{f}_t ($J \times 1$) are latent true scores for the measurements $t = 1, \dots, T$. \mathbf{f}_t^* ($J \times 1$) are temporal deviation terms from group means that are measured at $t = 1, \dots, T$. $\mathbf{R} = [\mathbf{I}^\top, \mathbf{S}^\top]^\top$ ($2J \times 1$) are random effects, where \mathbf{I} ($J \times 1$) is a random intercept and \mathbf{S} ($J \times 1$) is a random slope. Finally, $\mathbf{C} = [\mathbf{A}^\top, \mathbf{B}^\top]^\top$ ($2J \times 1$) are accumulating factors, where \mathbf{A} ($J \times 1$) has a constant effect over time and \mathbf{B} ($J \times 1$) has a changing effect over time. Hence, the total number of latent variables is $2JT + 4J = 2J(T + 2) = V$. I sort all variables in the following order:

$\mathbf{y}_1, \dots, \mathbf{y}_T, \mathbf{f}_1, \dots, \mathbf{f}_T, \mathbf{f}_1^*, \dots, \mathbf{f}_T^*, \mathbf{I}, \mathbf{S}, \mathbf{A}, \mathbf{B}$.

Table 1 summarizes the building blocks of the unified framework. I ignored the

mean structure in my first draft. $\boldsymbol{\tau}_y$ and $\boldsymbol{\tau}_A$ are uninteresting in stationary models and could be replaced by centering. $\boldsymbol{\tau}_B$ contains potentially interesting parameters. I suppose that \mathbf{I} and \mathbf{S} also have means but Usami et al. do not mention them. I am open to including a mean structure later. Because we only consider single-indicator models, Λ_f and Λ_{f^*} are identity matrices of size $J \times J$.

Symbol	Usami ¹	Description	Used in model
$\boldsymbol{\tau}_y$	μ_{xt}	observed means	CLPM, FCLPM, RICLPM, STARTS
Λ_f	-	Factor loadings of \mathbf{f}_t	all
Λ_{f^*}	-	Factor loadings of \mathbf{f}_t^*	all
Γ	β_x, γ	ARCLP	all but LCS
Ω	same	residual var/cov	all
Ψ	same	unique var/cov	FCLPM, STARTS, LCS
Φ_I	-	random intercepts var/cov	RICLPM, STARTS, LCMSR
Φ_S	-	random slopes var/cov	LCMSR
$\Phi_{I,S}$	-	covariance between RE	LCMSR
$\boldsymbol{\tau}_A$	-	constant accumulating factors mean	ALT, LCS
Φ_A	-	constant accumulating factors var/cov	ALT, LCS
$\boldsymbol{\tau}_B$	-	changing accumulating factors mean	ALT
Φ_B	-	changing accumulating factors var/cov	ALT
$\Phi_{A,B}$	-	covariance between CF	ALT
Γ^*	β_x^*, γ_x^*	LCS ARCLP	LCS
		Not used. See below.	

Table 1

¹: Notation used in Usami et al.

The matrices Φ_I , Φ_S , Φ_A , and Φ_B are symmetrical variance-covariance matrices and are structured similarly. Consider a bivariate process with two random intercepts

named I_1 and I_2 . Then, it follows that

$$\Phi_{\mathbf{I}} = \begin{bmatrix} \text{Cov}(I_1, I_1) & \text{Cov}(I_2, I_1) \\ \text{Cov}(I_2, I_1) & \text{Cov}(I_2, I_2) \end{bmatrix}$$

Conversely, $\Phi_{\mathbf{I}, \mathbf{S}}$ and $\Phi_{\mathbf{I}, \mathbf{S}}$ are not symmetrical. Both matrices have the following pattern (example given for a bivariate process with random intercepts and slopes):

$$\Phi_{\mathbf{I}, \mathbf{S}} = \begin{bmatrix} \text{Cov}(I_1, S_1) & \text{Cov}(I_1, S_2) \\ \text{Cov}(I_2, S_1) & \text{Cov}(I_2, S_2) \end{bmatrix}$$

We use the endogenous version of the ALT model where the factor loadings of the constant accumulating factors \mathbf{A} and the factor loadings of the changing accumulating factors \mathbf{B} are constrained at the first time point $t = 1$. The factor loadings of \mathbf{A} are constrained to $(\mathbf{I}_J - \mathbf{\Gamma})^{-1}$ at $t = 1$ and are \mathbf{I}_J for $t \in 2, \dots, T$. The factor loadings of \mathbf{B} are constrained to $\mathbf{\Gamma}[(\mathbf{I}_J - \mathbf{\Gamma})(\mathbf{I}_J - \mathbf{\Gamma})]^{-1}$ at $t = 1$ and to $(t - 1)\mathbf{I}_J$ for $t \in 2, \dots, T$.

The LCS model can be respecified as an ALT model by removing the changing accumulating factors \mathbf{B} from the ALT model and using $\mathbf{\Gamma}^* = \mathbf{\Gamma} + \mathbf{I}_J$. Thus, it is not necessary to use $\mathbf{\Gamma}^*$ in the backend.