Basic Latent Markov model

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Outline

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Introduction

Background:

Latent Markov (LM) models (Wiggins, 1973; Bartolucci et al., 2012) are successfully applied in the analysis of longitudinal data: they allow to take into account several aspects, such as serial dependence between observations, measurement errors, unobservable heterogeneity

LM models assume that one or more occasion-specific response variables depends only on a discrete latent variable characterized by a given number of latent states which in turn depends on the latent variables corresponding to the previous occasions according to a first-order Markov chain

LM models are characterized by several parameters: the initial probabilities to belong to a given latent state, the transition probabilities from a latent state to another one, the conditional response probabilities given the discrete latent variable

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The basic LM model may be seen as

- a generalization of a discrete-time Markov chain model to account for measurement errors in the observed variables of interest
- a generalization of a latent class (LC) model for longitudinal data, in which each subject may move between latent classes
- E.g., in the univariate case



Notation

- Repeated measurements of the same response variable on the same subjects at different occasions
- $\tilde{Y} = (Y^{(1)}, \dots, Y^{(T)})$: vector of values assumed by the categorical response variable *Y* at time *t* (*t* = 1, ..., *T*), having *c* categories
- $U^{(t)}$: latent state at time *t* with state space $\{1, \ldots, k\}$
- $U = (U^{(1)}, \dots, U^{(T)})$: vector describing the latent process

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Main assumptions

- local independence: response variables in Y are conditionally independent given the latent process U, i.e.,
 each occasion-specific observed variable Y^(t) is independent of all its previous values Y^(t-1),...,Y⁽¹⁾, given U^(t)
- latent process *U* follows a first-order Markov chain with *k* latent states, i.e.,

each latent variable $U^{(t)}$ is independent of $U^{(t-2)}, \ldots, U^{(1)}$, given $U^{(t-1)}$

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Parameters

• k(c-1) conditional response probabilities

$$\phi_{y|u}^{(t)} = p(Y^{(t)} = y|U^{(t)} = u)$$
 $t = 1, \dots, T; \ u = 1, \dots, k; \ y = 0, \dots, c-1$

• (k-1) initial probabilities

$$\pi_{\boldsymbol{u}} = p(\boldsymbol{U}^{(1)} = \boldsymbol{u}) \quad \boldsymbol{u} = 1, \dots, k$$

• (T-1)k(k-1) transition probabilities = (T-1)k(k-1) + (T-1) + (T-1)k(k-1) + (T-1)k(k-1)k(k-1) + (T-1)k(k

$$\pi_{u|v}^{(t)} = p(U^{(t)} = u|U^{(t-1)} = v) \quad t = 2, \dots, T; \ u, v = 1, \dots, k$$

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par = $k(c-1) + (k-1) + (T-1)k(k-1)$

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Probability distributions

•
$$p(\boldsymbol{U} = \boldsymbol{u}) = \pi_u \prod_{t=2}^T \pi_{u|v}^{(t)} = \pi_u \cdot \pi_{u_2|u}^{(2)} \dots \pi_{u_T|u_{T-1}}^{(T)}$$

•
$$p(\tilde{Y} = y | U = u) = \prod_{t=1}^{T} \phi_{y|u}^{(t)} = \phi_{y|u}^{(1)} \cdot \phi_{y|u}^{(2)} \dots \phi_{y|u}^{(T)}$$

• manifest distribution of \tilde{Y}

$$p(\tilde{Y} = \mathbf{y}) = \sum_{u} p(\tilde{Y} = \mathbf{y}, U = u) = \sum_{u} p(U = u) \cdot p(\tilde{Y} = \mathbf{y} | U = u)$$
$$= \sum_{u} \pi_{u} \phi_{y|u}^{(1)} \cdot \sum_{u_{2}} \pi_{u_{2}|u}^{(2)} \phi_{y|u}^{(2)} \dots \sum_{u_{T}} \pi_{u_{T}|u_{T-1}}^{(T)} \phi_{y|u}^{(T)}$$
$$= \sum_{u} \sum_{u_{2}} \dots \sum_{u_{T}} \pi_{u} \prod_{t=2}^{T} \pi_{u|v}^{(t)} \prod_{t=1}^{T} \phi_{y|u}^{(t)}$$

Note that computing $p(\tilde{Y} = y)$ involves all the possible k^T configurations of vector u

Computing of the manifest distribution: example

- We assume three occasions (T = 3) and three latent states (k = 3)
- We have $3^3 = 27$ possible configurations of vector *u*
- The manifest distribution of \tilde{Y} is given by:

$$p(\tilde{\mathbf{Y}} = \mathbf{y}) = \pi_1 \pi_{1|1}^{(2)} \pi_{1|1}^{(3)} \phi_{y|1}^{(1)} \phi_{y|1}^{(2)} \phi_{y|1}^{(3)} + \\ + \pi_1 \pi_{1|1}^{(2)} \pi_{2|1}^{(3)} \phi_{y|1}^{(1)} \phi_{y|1}^{(2)} \phi_{y|2}^{(3)} + \\ + \dots + \\ + \pi_3 \pi_{3|3}^{(2)} \pi_{3|3}^{(3)} \phi_{y|3}^{(1)} \phi_{y|3}^{(2)} \phi_{y|3}^{(3)}$$

Notation

- Y^(t) = (Y^(t)₁,...,Y^(t)_r): vector of categorical response variables Y_j (j = 1,...,r) observed at time t (t = 1,...,T), having c_j categories
- Y = (Y⁽¹⁾,...,Y^(T)): vector of observed responses made of the union of vectors Y^(t); usually, it is referred to repeated measurements of the same variables Y_j (j = 1,...,r) on the same individuals at different time points
- $U^{(t)}$: latent state at time *t* with state space $\{1, \ldots, k\}$
- $\boldsymbol{U} = (U^{(1)}, \dots, U^{(T)})$: vector describing the latent process

Main assumptions

• Local independence: vectors $Y^{(t)}$ (t = 1, ..., T) are conditionally independent given the latent process U and the response variables in each $Y^{(t)}$ are conditionally independent given $U^{(t)}$, i.e.,

each occasion-specific observed variable $Y_j^{(t)}$ is independent of $Y_j^{(t-1)}, \ldots, Y_j^{(1)}$ and of each $Y_h^{(t)}$, for all $h \neq j = 1, \ldots, r$, given $U^{(t)}$

• latent process *U* follows a first-order Markov chain with *k* latent states, i.e.,

each latent variable $U^{(t)}$ is independent of $U^{(t-2)}, \ldots, U^{(1)}$, given $U^{(t-1)}$

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Parameters

• $k \sum_{j=1}^{r} (c_j - 1)$ conditional response probabilities $\phi_{jy|u}^{(t)} = p(Y_j^{(t)} = y|U^{(t)} = u) \quad j = 1, \dots, r; \ t = 1, \dots, T; \ u = 1, \dots, k; \ y = 0, \dots, c_j - 1$ $\phi_{\mathbf{y}|u}^{(t)} = \prod_{j=1}^{r} \phi_{jy|u}^{(t)} = p(Y_1^{(t)} = y_1, \dots, Y_r^{(t)} = y_r|U^{(t)} = u)$

• (k-1) initial probabilities $\pi_u = p(U^{(1)} = u) \quad u = 1, \dots, k$

•
$$(T-1)k(k-1)$$
 transition probabilities
 $\pi_{u|v}^{(t)} = p(U^{(t)} = u|U^{(t-1)} = v)$ $t = 2, ..., T; u, v = 1, ..., k$

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par = $k \sum_{j=1}^{r} (c_j - 1) + (k - 1) + (T - 1)k(k - 1)$

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Probability distributions

•
$$p(\boldsymbol{U} = \boldsymbol{u}) = \pi_u \prod_{t=2}^T \pi_{u|v}^{(t)} = \pi_u \cdot \pi_{u_2|u}^{(2)} \dots \pi_{u_T|u_{T-1}}^{(T)}$$

•
$$p(\mathbf{Y} = \mathbf{y} | \mathbf{U} = \mathbf{u}) = \prod_{t=1}^{T} \phi_{\mathbf{y}|u}^{(t)} = \phi_{\mathbf{y}|u}^{(1)} \cdot \phi_{\mathbf{y}|u}^{(2)} \dots \phi_{\mathbf{y}|u}^{(T)}$$

• Manifest distribution of Y

$$p(\mathbf{Y} = \mathbf{y}) = \sum_{\mathbf{u}} p(\mathbf{Y} = \mathbf{y}, \mathbf{U} = \mathbf{u}) = \sum_{\mathbf{u}} p(\mathbf{U} = \mathbf{u}) \cdot p(\mathbf{Y} = \mathbf{y} | \mathbf{U} = \mathbf{u})$$
$$= \sum_{u} \pi_{u} \phi_{\mathbf{y}|u}^{(1)} \cdot \sum_{u_{2}} \pi_{u_{2}|u}^{(2)} \phi_{\mathbf{y}|u}^{(2)} \dots \sum_{u_{T}} \pi_{u_{T}|u_{T-1}}^{(T)} \phi_{\mathbf{y}|u}^{(T)}$$
$$= \sum_{u} \sum_{u_{2}} \dots \sum_{u_{T}} \pi_{u} \prod_{t=2}^{T} \pi_{u|v}^{(t)} \prod_{t=1}^{T} \phi_{\mathbf{y}|u}^{(t)}$$

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Maximum likelihood (ML) estimation

• Log-likelihood of the model

$$\ell(\boldsymbol{\theta}) = \sum_{\mathbf{y}} n_{(\mathbf{y})} \log[p(\mathbf{y} = \mathbf{y})]$$

- θ : vector of all model parameters $(\pi_u, \pi_{u|v}^{(t)}, \phi_{jy|u}^{(t)})$
- *n*(*y*): frequency of the response configuration *y* in the sample
- ℓ(θ) may be maximized with respect to θ by an Expectation-Maximization (EM) algorithm (Dempster et al., 1977)

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EM algorithm

Complete data log-likelihood of the model

$$\ell^{*}(\boldsymbol{\theta}) = \sum_{j=1}^{r} \sum_{t=1}^{T} \sum_{u=1}^{k} \sum_{y=0}^{c-1} a_{juy}^{(t)} \log \phi_{jy|u}^{(t)} + \sum_{u=1}^{k} b_{u}^{(1)} \log \pi_{u} + \sum_{t=2}^{T} \sum_{y=1}^{k} \sum_{u=1}^{k} b_{vu}^{(t)} \log \pi_{u|v}^{(t)}$$

- a^(t)_{juy}: frequency of subjects responding by y for the j-th response variable and belonging to latent state u, at time t
- $b_u^{(1)}$: frequency of subjects in latent state *u* at time 1
- $b_{vu}^{(t)}$: frequency of subjects which move from latent state v to u at time t

EM algorithm

- The algorithm *alternates two steps* until convergence in $\ell(\theta)$:
 - **E**: compute the expected values of frequencies $a_{juy}^{(t)}$, $b_u^{(1)}$, and $b_{vu}^{(t)}$, given the observed data and the current value of θ , so as to obtain the expected value of $\ell^*(\theta)$
 - **M**: update θ by maximizing the expected value of $\ell^*(\theta)$ obtained above; explicit solutions for θ estimations are available
- The E-step is performed by means of certain recursions

Forward and backward recursions

To efficiently compute the probability p(Y = y) and, then, the posterior probabilities $f_{u|y}^{(t)}$ and $f_{u|y,y}^{(t)}$ we can use forward and backward recursions for obtaining the following intermediate quantities

• Forward recursions

$$q_{u,\mathbf{y}}^{(t)} = p(U^{(t)} = u, \mathbf{Y}^{(1)}, \dots, \mathbf{Y}^{(t)}) = \sum_{\nu=1}^{k} q_{\nu,\mathbf{y}}^{(t-1)} \pi_{u|\nu}^{(t)} \phi_{\mathbf{y}|u}^{(t)} \quad u = 1, \dots, k$$

starting with $q_{u,y}^{(1)} = \pi_u \phi_{y|u}^{(1)}$

Backward recursions

$$\bar{q}_{v,\mathbf{y}}^{(t)} = p(\mathbf{Y}^{(t+1)}, \dots, \mathbf{Y}^{(T)} | U^{(t)} = v) = \sum_{u=1}^{k} \bar{q}_{u,\mathbf{y}}^{(t+1)} \pi_{u|v}^{(t+1)} \phi_{\mathbf{y}|u}^{(t+1)} \quad v = 1, \dots, k$$

starting with $\bar{q}_{v,y}^{(T)} = 1$

Variants of basic LM model

- LM model with covariates: An LM model may be generalized in a similar way to the basic LC model introducing individual covariates
- Constrained LM model: Several interesting constraints may be introduced in an LM model to reduce the number of parameters and to make easier the interpretation of results
 - constraints on the conditional distribution
 - constraints on the transition probabilities
- In what follows we describe some of the most interesting constraints on the transition probabilities

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Constraints on $\Pi^{(t)}$

- We denote by $\Pi^{(t)} = \{\pi^{(t)}_{u|v}\}$ the matrix of transition probabilities
- Linear constraint: $\rho_{\nu}^{(t)} = \mathbf{Z}_{\nu}^{(t)} \boldsymbol{\delta}$, with $\rho_{\nu}^{(t)}$ denoting a column vector containing the off-diagonal elements of the *v*-th row of $\mathbf{\Pi}^{(t)}$
- More in general, a GLM may be imposed on the transition probabilities $\lambda_{\nu}^{(t)} = \mathbf{Z}_{\nu}^{(t)} \boldsymbol{\delta}$, with $\lambda_{\nu}^{(t)} = g(\pi_{\nu}^{(t)})$; e.g., $g(\cdot)$ may be a logit link function, so that the generic element of $\lambda_{\nu}^{(t)}$ is $\lambda_{u|\nu}^{(t)} = \log \frac{\pi_{u|\nu}^{(t)}}{\pi_{v|\nu}^{(t)}}$; $u = 1, \ldots, k, \ u \neq v$

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Examples of constraints on $\Pi^{(t)}$

- C1 Time-homogeneous Markov-chain: $\Pi^{(t)} = \Pi$ \rightarrow Transition probability from state *v* to state *u* is independent of the occasion *t*
- C2 All the off-diagonal transition probabilities are equal to each other

$$\mathbf{\Pi}^{(t)} = \begin{pmatrix} 1 - 2\pi^{(t)} & \pi^{(t)} & \pi^{(t)} \\ \pi^{(t)} & 1 - 2\pi^{(t)} & \pi^{(t)} \\ \pi^{(t)} & \pi^{(t)} & 1 - 2\pi^{(t)} \end{pmatrix}, \quad t = 2, \dots, T$$

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C3 Symmetric transition matrix: transition probability from state *v* to state *u* is the same as the reverse transition

$$\mathbf{\Pi}^{(t)} = \begin{pmatrix} 1 - (\pi_{2|1}^{(t)} + \pi_{3|1}^{(t)}) & \pi_{2|1}^{(t)} & \pi_{3|1}^{(t)} \\ \\ \pi_{2|1}^{(t)} & 1 - (\pi_{2|1}^{(t)} + \pi_{3|2}^{(t)}) & \pi_{3|2}^{(t)} \\ \\ \\ \pi_{3|1}^{(t)} & \pi_{3|2}^{(t)} & 1 - (\pi_{3|1}^{(t)} + \pi_{3|2}^{(t)}) \end{pmatrix},$$

C4 Upper-triangular transition matrix: a subject in state v may move only in state u = v + 1, ..., k

$$\mathbf{\Pi}^{(t)} = \begin{pmatrix} \pi_{1|1}^{(t)} & \pi_{2|1}^{(t)} & \pi_{3|1}^{(t)} \\ \mathbf{0} & \pi_{2|2}^{(t)} & \pi_{3|2}^{(t)} \\ \mathbf{0} & \mathbf{0} & 1 \end{pmatrix}, \quad t = 2, \dots, T$$

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C5 Tridiagonal transition matrix: transition from state *v* is only allowed to state u = v - 1, v + 1

$$\mathbf{\Pi}^{(t)} = \begin{pmatrix} \pi_{1|1}^{(t)} & \pi_{2|1}^{(t)} & \mathbf{0} & \mathbf{0} \\ \pi_{1|2}^{(t)} & \pi_{2|2}^{(t)} & \pi_{3|2}^{(t)} & \mathbf{0} \\ \pi_{2|3}^{(t)} & \pi_{3|3}^{(t)} & \pi_{4|3}^{(t)} \\ \mathbf{0} & \mathbf{0} & \pi_{3|4}^{(t)} & \pi_{4|4}^{(t)} \end{pmatrix}, \quad t = 2, \dots, T$$

C6 Basic LC model: transition from state v to state u equals 0, for all $v \neq u$

$$\mathbf{\Pi}^{(t)} = \begin{pmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{pmatrix}, \quad t = 2, \dots, T$$

Model selection

- Given the number k of latent states, it is convenient to sequentially introduce the constraints on Π^(t) and retain the constraint that, at each attempt, leads to a reduction of the BIC index
- Also an LR test is possible, based on the test statistic

$$LR = -2(\hat{\ell}_0 - \hat{\ell}_1)$$

- $\hat{\ell}_1$: maximum log-likelihood value of the unconstrained model
- $\hat{\ell}_0$: maximum log-likelihood value of the constrained model
- What's about the distribution of LR statistic?
 - when the usual regularity conditions hold (e.g., case C1), LR statistic has a chi-square distribution with a number of degrees of freedom equal to the number of free parameters
 - in presence of constraints on the boundary space (e.g., cases C4 and C5), LR statistic has a chi-bar-squared distribution (i.e., a mixture of chi-squared distributions)

Analysis of marijuana consumption

- We again consider the dataset about marijuana consumption, assuming that individuals may move from one state to another one during the time
- We first estimate an unconstrained basic LM model, characterised by a completely general transition matrix $\Pi^{(t)}$ (model LM1)
- Then, we estimate some more realistic constrained LM models, characterised by
 - a time-homogeneous transition matrix Π (constraint C1; model LM2)
 - a time-homogeneous and tridiagonal transition matrix Π (constraints C1 and C5; model LM3)
 - a time-homogeneous and upper-triangular transition matrix Π (constraints C1 and C4; model LM4)
- Note that LM2 is nested in LM1, LM3 is nested in LM2, and LM4 is nested in LM2; LM3 and LM4 are not nested

Unconstrained basic LM model (LM1)

Table : Estimates of conditional response probabilities, $\hat{\phi}_{y|u}$

	y = 0	y = 1	y = 2
u = 1	0.996	0.000	0.004
u = 2	0.305	0.687	0.008
<i>u</i> = 3	0.012	0.083	0.905

- Results are coherent with those obtained by the LC model (Lecture 1, Example 1): state 1 corresponds to the lowest tendency of marijuana consumption, state 2 to an intermediate tendency, and state 3 to the highest tendency
- Note that the results outline the presence of measurement errors, as some off-diagonal values differ from 0

Table : Estimates of initial, $\hat{\pi}_u$, and transition probabilities, $\hat{\pi}_{u|v}^{(t)}$

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		u = 1	u = 2	<i>u</i> = 3
t = 1		0.898	0.083	0.019
t = 2	v = 1	0.831	0.154	0.015
	v = 2	0.318	0.228	0.454
	v = 3	0.057	0.000	0.943
t = 3	v = 1	0.810	0.190	0.000
	v = 2	0.056	0.482	0.461
	v = 3	0.000	0.147	0.853
t = 4	v = 1	0.908	0.064	0.028
	v = 2	0.059	0.718	0.224
	v = 3	0.000	0.186	0.814
t = 5	v = 1	0.789	0.163	0.048
	v = 2	0.099	0.821	0.080
	v = 3	0.020	0.035	0.945

- Individuals tend to be in state 1 (low tendency to marijuana consumption) at the beginning of the study (t = 1)
- However, the completely general transition matrices Π^(t) are not easy to be interpreted
- In order to have information about the time trend of the tendency of marijuana consumption, we may
 - calculating the marginal distribution of latent states for each time occasion
 - constraining the transition matrices in order to reduce the number of parameters

Table : Estimated marginal probabilities of latent states, $p(U_i^{(t)} = u), u = 1, ..., 5$

	u = 1	u = 2	<i>u</i> = 3
t = 1	0.8980	0.0835	0.0185
t = 2	0.7736	0.1576	0.0688
<i>t</i> = 3	0.6355	0.2331	0.1314
t = 4	0.5905	0.2325	0.1770
<i>t</i> = 5	0.4924	0.2933	0.2143

- We observe that the tendency to stay in state 1 (low marijuana consumption) decreases with the age
- The tendency to consume marijuana (states 2 and 3) increases with the age

Constrained basic LM model (LM1)

Table : Estimates of conditional response probabilities, $\hat{\phi}_{y|u}$

	y = 0	y = 1	y = 2
u = 1	0.996	0.000	0.004
u = 2	0.305	0.687	0.008
<i>u</i> = 3	0.012	0.083	0.905

- Results are coherent with those obtained by the LC model (Lecture 1, Example 1): state 1 corresponds to the lowest tendency of marijuana consumption, state 2 to an intermediate tendency, and state 3 to the highest tendency
- Note that the results outline the presence of measurement errors, as some off-diagonal values differ from 0

Time-homogeneous LM model (LM2)

Table : Estimates of initial, $\hat{\pi}_u$, and transition probabilities, $\hat{\pi}_{u|v}^{(t)}$

		u = 1	u = 2	<i>u</i> = 3
t = 1		0.912	0.071	0.017
$t=2,\ldots,5$	v = 1	0.842	0.141	0.017
	v = 2	0.080	0.670	0.250
	v = 3	0.000	0.132	0.868

 High persistency in each latent state, but also a given tendency to move to adjacent states

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Time-homogeneous LM model with tridiagonal transition matrix (LM3)

Table : Estimates of initial, $\hat{\pi}_{u}$, and transition probabilities, $\hat{\pi}_{u|v}^{(t)}$

		u = 1	u = 2	<i>u</i> = 3
t = 1		0.896	0.089	0.015
$t=2,\ldots,5$	v = 1	0.835	0.165	0.000
	v = 2	0.070	0.686	0.244
	v = 3	0.000	0.082	0.918

 High persistency in each latent state, but also a given tendency to move to higher adjacent states

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Model selection

Table : Model selection for k = 3: maximum log-likelihood value, number of parameters, and BIC index

Model	$\hat{\ell}$	# par	BIC
LC	-658.238	32	1491.454
LM1	-646.895	32	1468.768
LM2	-658.593	14	1393.738
LM3	-660.600	12	1375.890
LM4	-661.930	11	1373.070

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R package LMest

- This package includes a set of functions to fit LM models in the basic version and in the extended version with individual covariates
- The main function for the model estimation is <code>est_lm_basic</code>

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Data structure

- > data(data_drug)
- > data_drug = as.matrix(data_drug)
- > head(data_drug)

```
> S=data_drug[,1:5]-1 # matrix of item responses
> yv=data drug[,6] # vector of weights
```

> k=3 # number of latent states

Model estimation

- > # Basic unconstrained LM model
- > LM1 = est_lm_basic(S,yv,k,mod=0)
- > # Time-homogeneous LM model
- > LM2 = est_lm_basic(S,yv,k,mod=1)

Output

- > LM1\$piv # latent states initial probabilities
- > LM1\$Pi # transition probabilities
- > LM1\$Psi # conditional response probabilities

```
> # Marginal probabilities
> marg_prob_2 = colSums(LM1$Pi[,,2]*LM1$piv)
> marg_prob_3 = colSums(LM1$Pi[,,3]*marg_prob_2)
> marg_prob_4 = colSums(LM1$Pi[,,4]*marg_prob_3)
> marg_prob_5 = colSums(LM1$Pi[,,5]*marg_prob_4)
> round(rbind(LM1$piv, marg_prob_2, marg_prob_3,
marg_prob_4, marg_prob_5), 4)
```

Main references

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