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(1)

**Nonlinear Path Analysis
with
Optimal Scaling**

**Jan de Leeuw
Department of Data Theory FSW
University of Leiden**

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Path analysis: general idea [History]

$$\left\{ \begin{array}{l} Y = \beta_1 x_1 + \dots + \beta_m x_m + \epsilon \\ \epsilon \perp (x_1, \dots, x_m) \end{array} \right.$$

abundance environmental gradients.

Or:

$$Y_i = \beta_{i1} x_1 + \dots + \beta_{im} x_m + \epsilon_i$$

abundance of species i

$$\epsilon_i \perp (x_1, \dots, x_m).$$

[perhaps some β_{ij} are known to be zero]

Or:

$$Y = \beta_1 x_1 + \dots + \beta_m x_m + \epsilon,$$

$$x_m = \alpha_{m1} x_1 + \dots + \alpha_{m,m-1} x_{m-1} + \delta_m,$$

$$x_{m-1} = \alpha_{m-1,1} x_1 + \dots + \alpha_{m-1,m-2} x_{m-2} + \delta_{m-1}, \dots$$

....

$$\epsilon \perp (x_1, \dots, x_m)$$

$$\delta_m \perp (x_1, \dots, x_{m-1})$$

¶

perhaps some α_{ij} are known
to be zero [or equal to
each other].

Example.
time series

$$\left. \begin{array}{l} x_m = \alpha x_{m-1} + \delta_m \\ \vdots \\ x_{m-1} = \alpha x_{m-2} + \delta_{m-1} \\ \vdots \end{array} \right\}$$

First methodological point

Why introduce this structure (zero regression coeffs) (structure within the predictors)?

Answer.

The models are special, restricted case of models in which every variable in the system depends on all other variables. These very general models are not restrictive [$M = X$].

Reframed question

Why use restrictive models?

Answer. - Trade-off of bias and precision
- Curse of dimensionality.

Second methodological point

Are these structural models causal models?

Answer. ??

(a) you can formulate them in causal language

[X influences, explains, determines Y]

(b) you can interpret them in causal language

[the effect of X on Y is .74]

(c) but this has nothing to do with

{ necessary connections
 productivity
 manipulation } HUME

(d) Thus the word causal is used in a colloquial, imprecise sense, or in a statistical, precise sense. These two uses must be kept strictly separate. [DANGER]

Third methodological question

What is non linear?

It is not

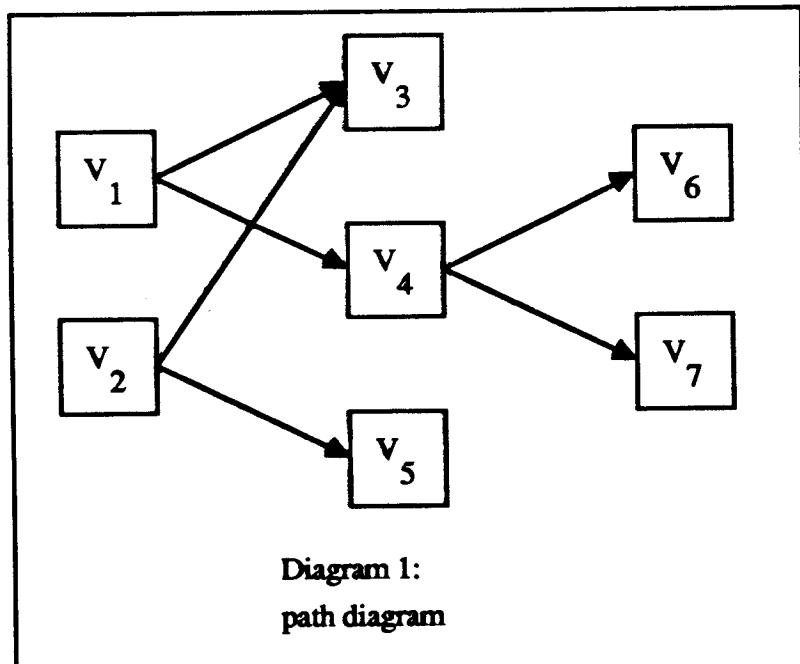
$$y = \phi(x_1, \dots, x_m) + \epsilon$$

but it is

$$\begin{cases} y = \phi_1(x_1) + \dots + \phi_m(x_m) + \epsilon \\ \phi_j \in \Phi_j \end{cases}$$

as in the previous lecture.

Perhaps quasilinear, semilinear is better.



	level	causes	direct causes	predecessors
Var 1	0	****	****	****
Var 2	0	****	****	****
Var 3	1	{1,2}	{1,2}	{1,2}
Var 4	1	{1}	{1}	{1,2}
Var 5	1	{2}	{2}	{1,2}
Var 6	2	{1,4}	{4}	{1,2,3,4,5}
Var 7	2	{1,4}	{4}	{1,2,3,4,5}

Table 1:
causal relations in Diagram 1

Example

(a) where do you get your arrows from?

- prior knowledge (theory)
- time and space

(b) colloquial interpretations.

- V_3 does not influence V_6
- V_1 influences V_6 through V_4

(c) ~~signs~~ - definitions [qualitative]

- graph theory

Table 1

exogenous - endogenous

transitive - recursive

Quantitative translation

$$\left\{ \begin{array}{l} x_3 = \beta_{31} x_1 + \beta_{32} x_2 + \epsilon_3 \\ x_4 = \beta_{41} x_1 + \epsilon_4 \\ x_5 = \beta_{52} x_2 + \epsilon_5 \\ x_6 = \beta_{64} x_4 + \epsilon_6 \\ x_7 = \beta_{71} x_1 + \epsilon_7 \end{array} \right.$$

weak orthogonality

$$\left\{ \begin{array}{l} \epsilon_3 \perp (x_1, x_2) \\ \epsilon_4 \perp x_1 \\ \epsilon_5 \perp x_2 \\ \epsilon_6 \perp x_4 \\ \epsilon_7 \perp x_4 \end{array} \right.$$

saturated model.

strong orthogonality

(a) disturbances uncorrelated with exogenous

$$\{\epsilon_3, \epsilon_4, \epsilon_5, \epsilon_6, \epsilon_7\} \perp \{x_1, x_2\}$$

(b) disturbances different levels uncorrelated

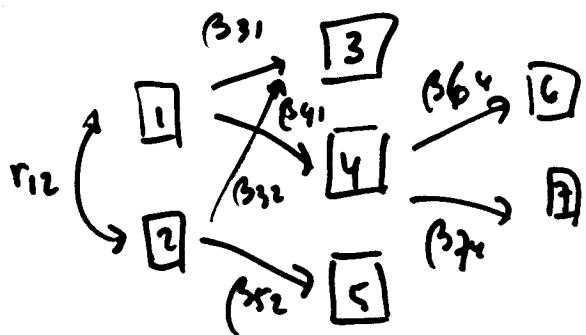
$$\{\epsilon_3, \epsilon_4, \epsilon_5\} \perp \{\epsilon_6, \epsilon_7\}.$$

Main results under strong orthogonality

(1) Causal interpretation

(Given the direct causes a variable is independent of its other predecessors)

(b) Calculus of path coefficients



$$r_{15} = r_{12}\beta_{52}$$

$$r_{26} = r_{12}\beta_{14}\beta_{64}$$

$$r_{34} = \beta_{41}\beta_{31} + \beta_{41}r_{12}\beta_{23}$$

useful decompositions

$$r_{13} = \beta_{31} + \beta_{32}r_{12}$$

direct indirect

(c) ML = LS

Classes of models

① Recursive models [path calculus
cond. independence

ⓐ Saturated recursive models

- block recursive models

- causal chains

- multiple regression

$$y = \begin{matrix} Ax \\ By \\ \vdots \end{matrix} + \epsilon$$

exogenous

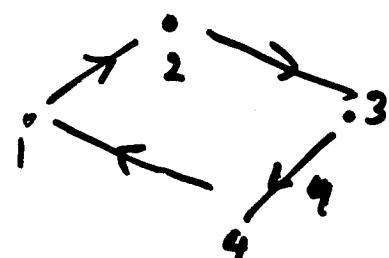
Strong orthogonality \equiv weak orthogonality

ⓑ nonsaturated recursive models

[fig 1]. Strong \neq weak

② Nonrecursive models

weak \Rightarrow saturated



[no path calculus
no cond. independence

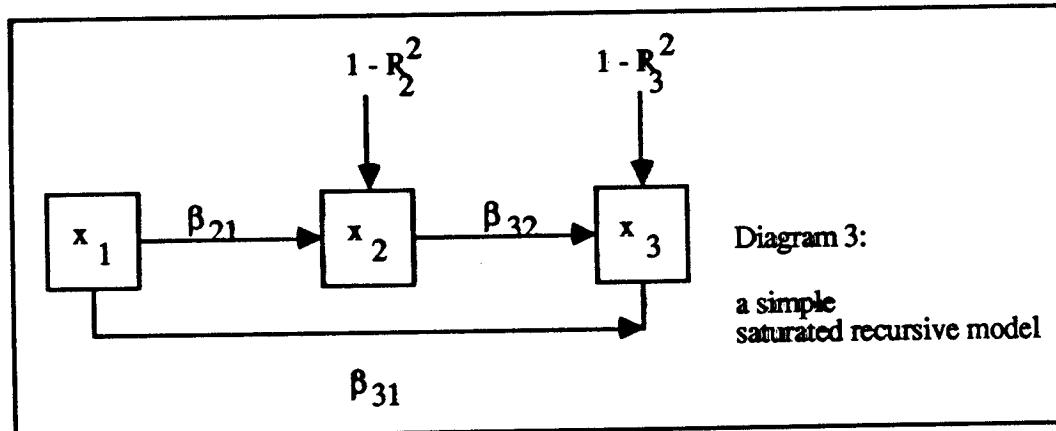
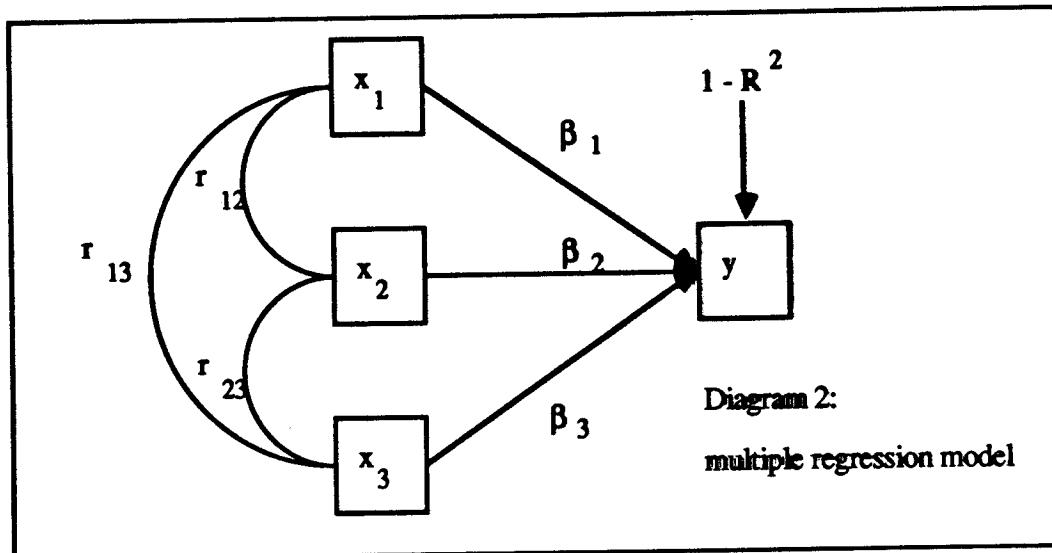
no level assignment

$$r(x_1, y) = \beta_1 + \beta_2 r_{12} + \beta_3 r_{13}$$

$$r(x_2, y) = \beta_1 r_{12} + \beta_2 + \beta_3 r_{23}$$

- - - - -

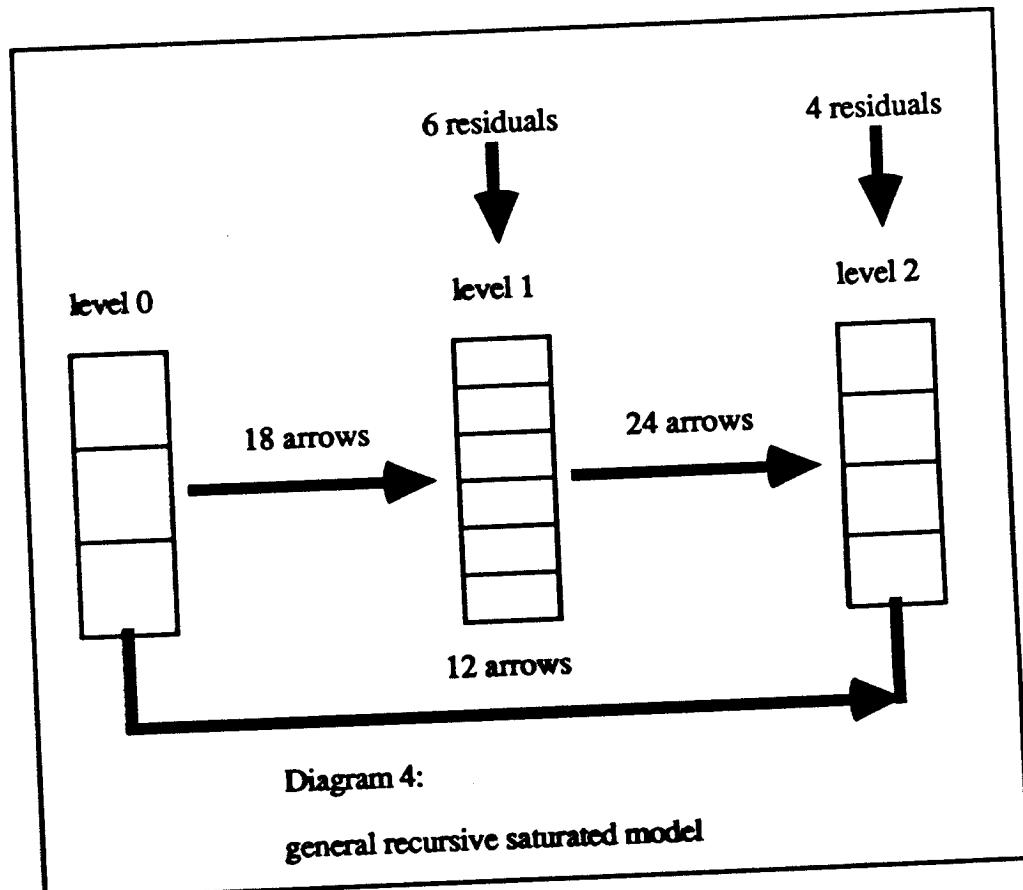
$$\begin{cases} r = R(\beta) \\ \beta = R^{-1}r \end{cases}$$



$$\beta_{31} = c \quad \Leftrightarrow \quad r_{13}|_{x_2} = 0.$$

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Computation

Recursive models

LS [projection] [on direct causes]

Nonrecursive models

[Weak orth : LS]

Strong or other : ??

Example from L&L

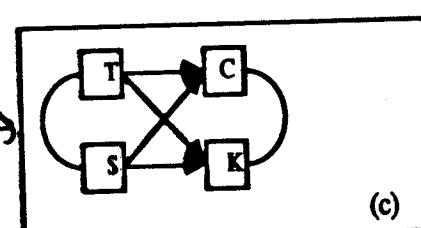
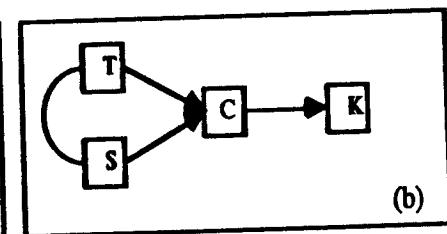
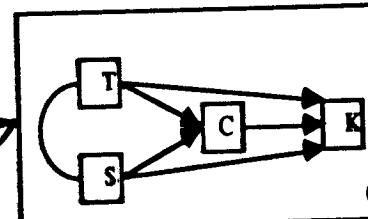
$\left\{ \begin{array}{l} N: \\ C: \\ S: \\ T: \end{array} \right.$	primary production Chlorophyll Salinity Temperature
---	--

Alternative models : fit

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	K	C	T
C	+.842		
T	.043	.236	
S	-.146	-.369	-.925

Correlations
Baie des Chaleurs



Three recursive models

Table 1:
Legendre and Legendre
Primary Production Data

	(a)	(b)	(c)
T \Rightarrow C	-0.730	-0.730	-0.730
S \Rightarrow C	-1.044	-1.044	-1.044
T \Rightarrow K	+0.031	*****	-0.638
S \Rightarrow K	+0.220	*****	-0.736
C \Rightarrow K	+0.916	+0.842	*****
VERR C	0.787	0.787	0.787
VERR K	0.260	0.291	0.920
CERR C,K		A	0.721

Seems fine

Latent variables

- (a) to extend the notion of a path model (and make it cover other existing techniques)
- (b) to model the idea of measurement error or of indicators

Examples in figure.

General example:

LISREL (Jöreskog)

EQS (Bentler)

COSAN (McDonald)

$$\begin{aligned} x &= A\gamma + B\chi + \xi \text{ or errors in equations} \\ x &= C\chi + \epsilon \quad \text{errors in variables.} \\ y &= D\eta + \delta \end{aligned}$$

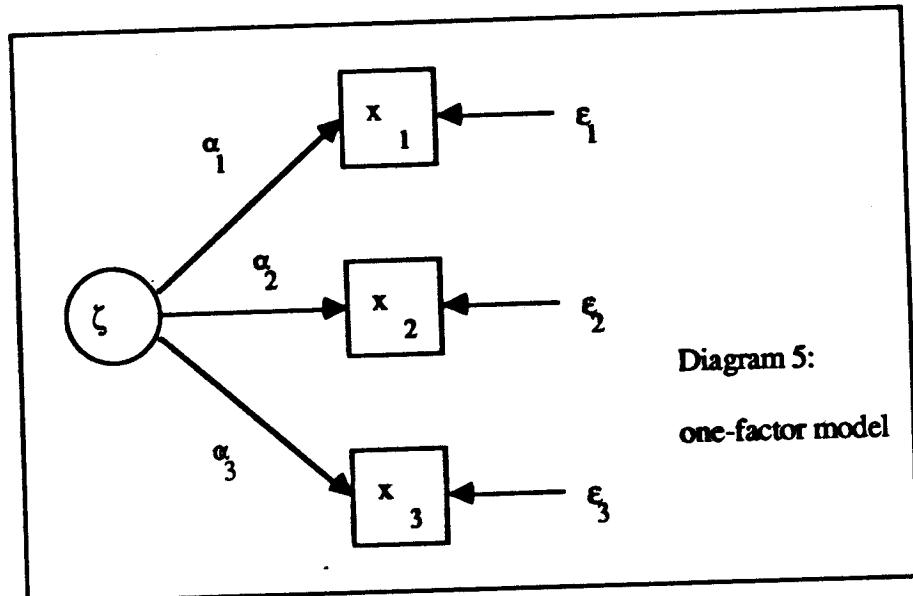


Diagram 5:
one-factor model

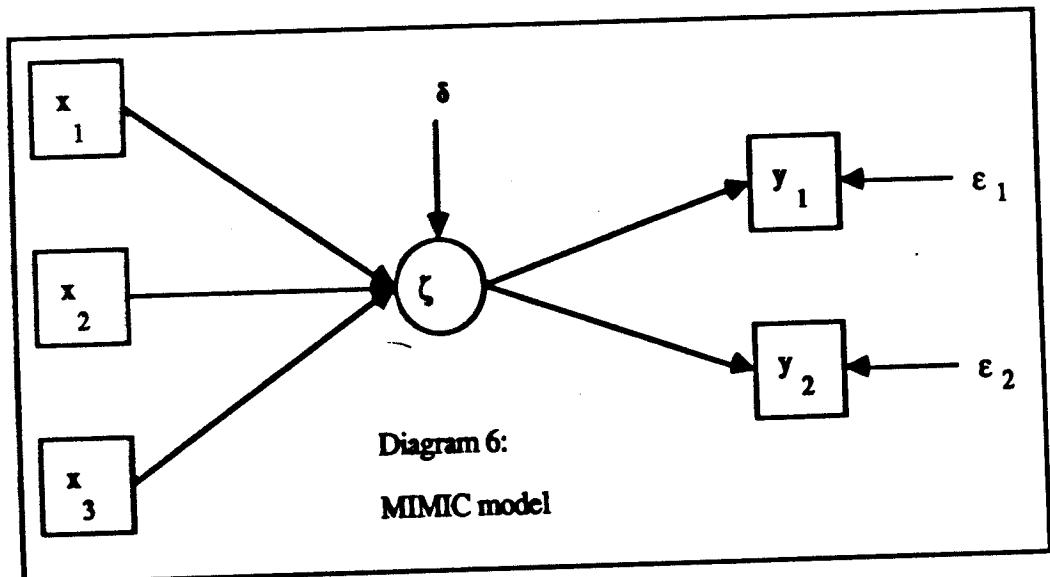


Diagram 6:
MIMIC model

Generalities to more factors

linear dynamic systems (1)

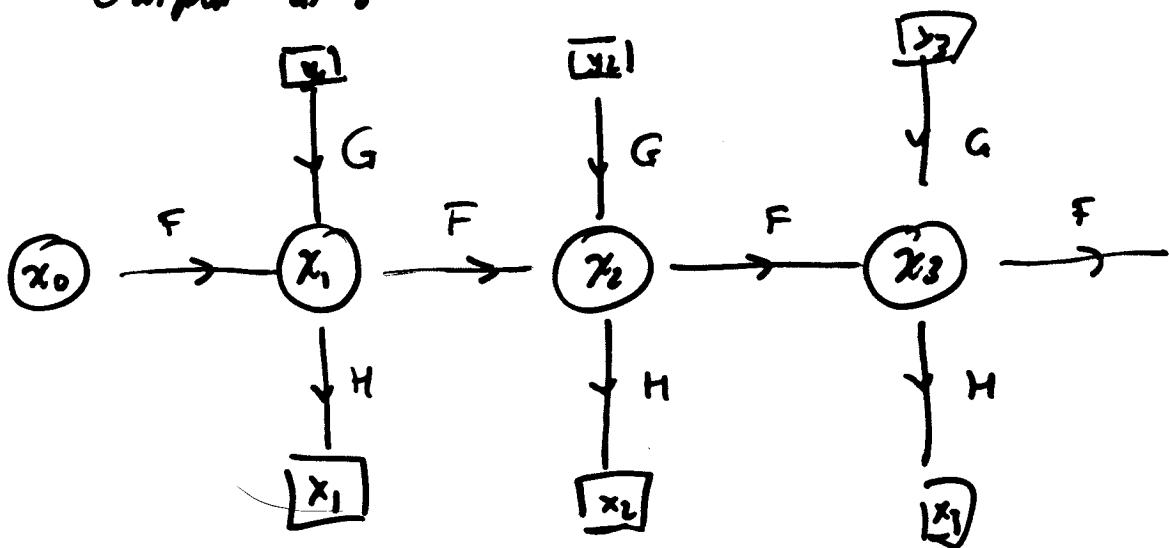
$$x_t = F x_{t-1} + G y_t + \epsilon_t$$

input or background
at t

$$x_t = H x_t + \delta_t$$

state of system at t

output at t



- General AR-model
- Kalman filter
- Also space
- Also nonstationary

And now the most original part

PATHS

Step 1: Take an arc diagram

Step 2: Translate it into regressions [linear structural equations]

Step 3: Translate these into a loss function

Step 4: Minimise over

(a) path coefficients

(b) latent variables, transformations, errors.

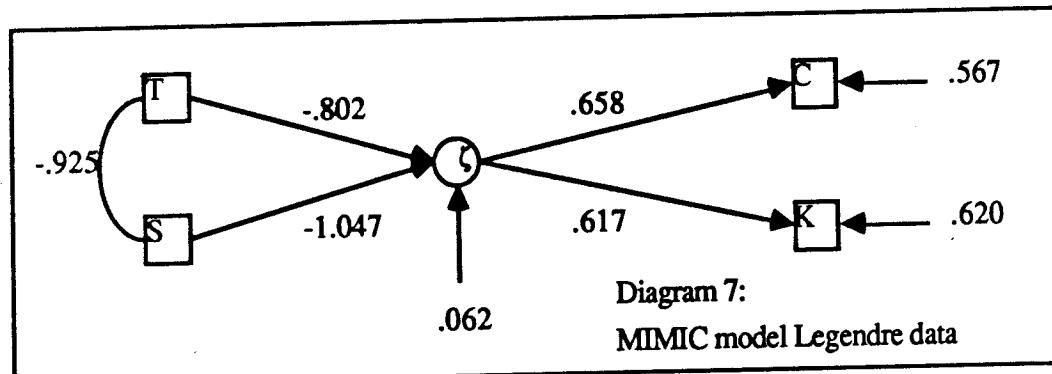
Example MIMIC from (6)

$$\text{loss} \approx \sum_j \alpha_j x_j - \beta_j - \delta_j^2 + \sum_k \|y_k - \beta_j - \epsilon_k\|^2$$

$$\text{loss} = \left\| \sum_j \alpha_j x_j - \beta_j - \delta_j \right\|^2 + \sum_k \|y_k - \beta_j - \epsilon_k\|^2$$

$$\begin{aligned} \text{weak : } & \delta \perp (x_1, x_2, x_3) \\ & \beta \perp (\epsilon_1, \epsilon_2) \end{aligned}$$

Perhaps transform. ALS algorithm.



	weights metric		weights nonmetric		explained variances	
	metric	nonmetric	metric	nonmetric	metric	nonmetric
WC	-.82	.10	-.96	-.20		
BS	-.06	.11	-.56	.39		
CM	.13	.21	-.14	-.33		
LR	-.02	.56	.28	.12		
FT	.72	-.24	.22	-.15		
CH	-.29	.10	-.71	.43		
S1	-.79	-.09	-.89	.22	.39	.21
S2	.04	-.79	.30	-.87	.36	.21
S3	-.85	-.35	-.88	-.16	.22	.16
S4	-.95	-.10	-.99	.21	.13	.04
S5	-.97	-.06	-.99	.22	.08	.04
S6	-.91	-.13	-.95	.19	.21	.10
S7	-.93	-.48	-.98	.01	.07	.04
S8	-.77	-.11	-.85	.00	.43	.27
S9	-.36	.52	.74	-.48	.53	.32
S10	.18	.88	.07	.90	.25	.16
S11	.52	.71	.48	.71	.36	.18
S12	.53	.53	.57	.54	.54	.31

Table 2: hunting spider data: metric and nonmetric MIMIC analysis

The important thing in PATHS

- scaling of variables , including latent variables
- latent variables are like other variables , but with a very low measurement level

Instead of

(linear
Latent)

{ linear
monotonic
polynomial
:
Latent }

Example Hunting spiker data

[and earlier

SPECIES
NITRO

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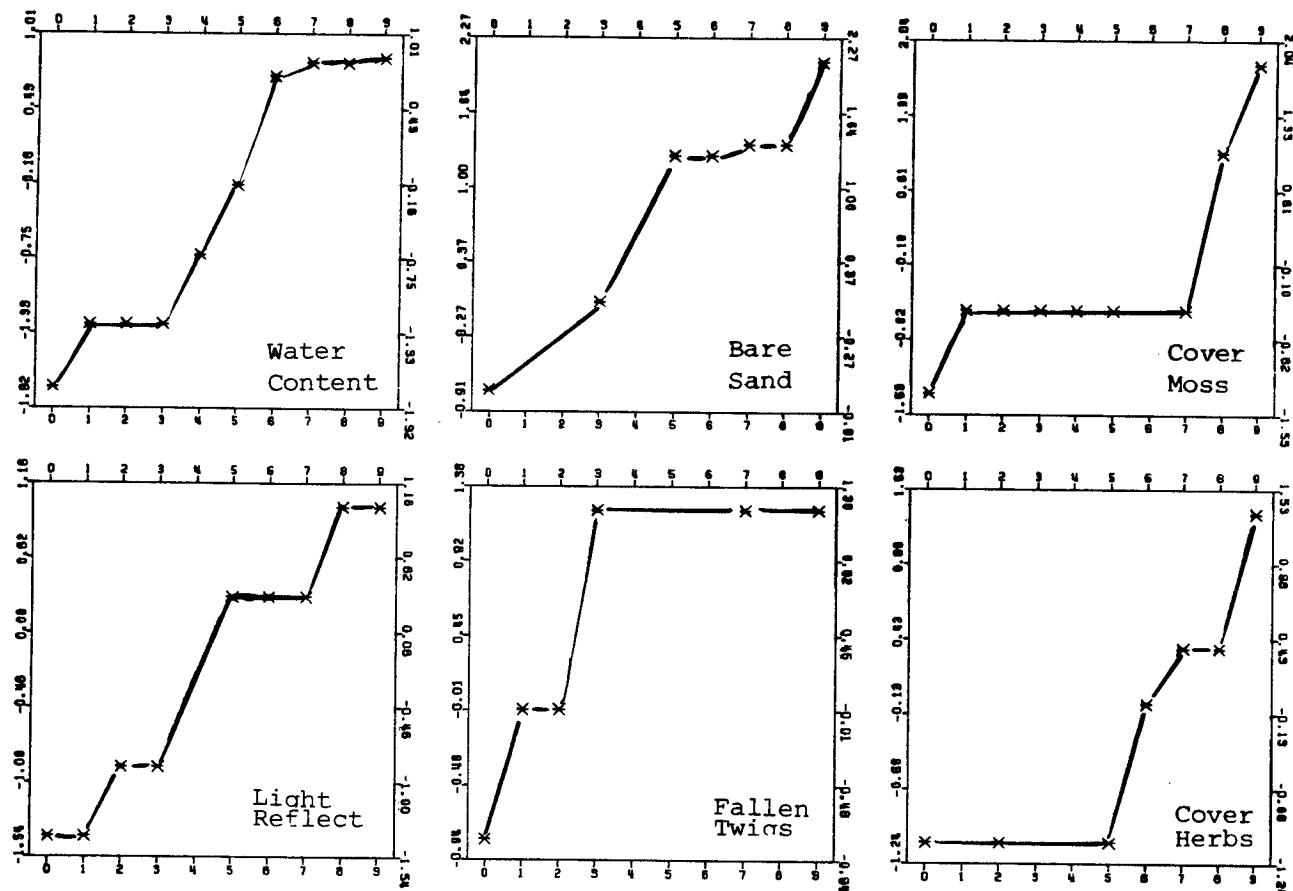
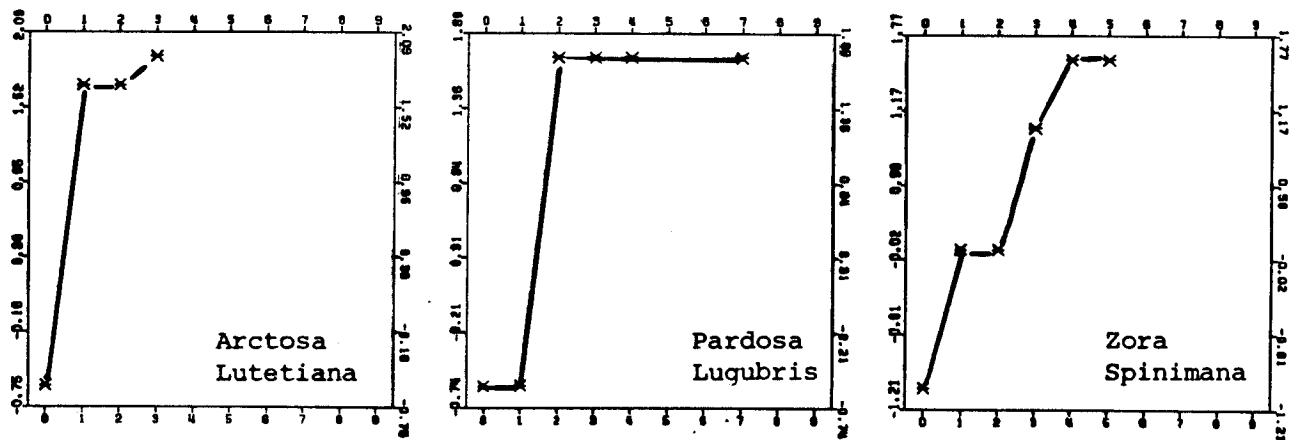


Diagram 8a:
hunting spider example
transformations of environmental variables



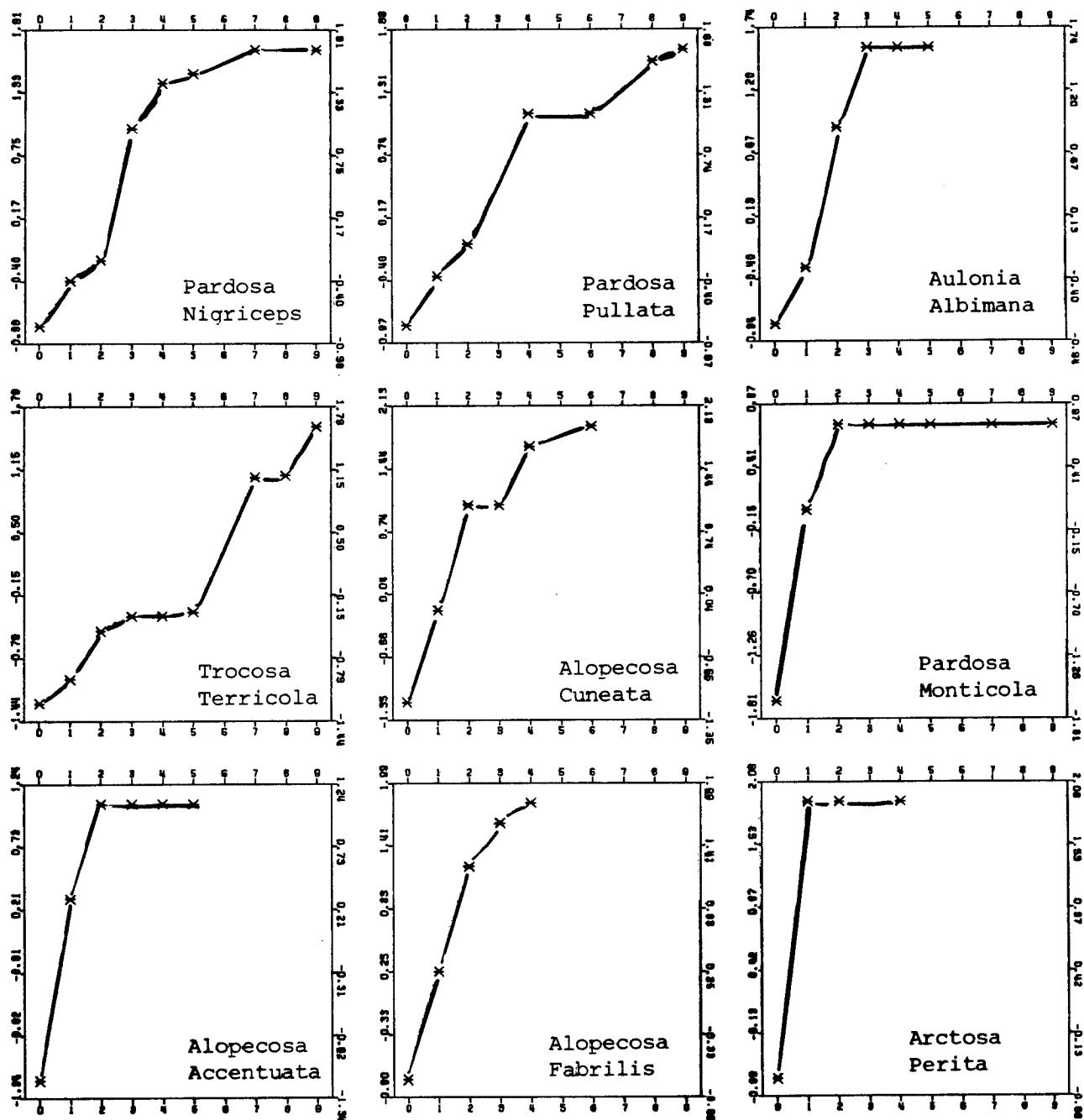


Diagram 8b:
hunting spider example
transformations of abundance variables