### Multiple imputation and Three-mode analysis. A research programme

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### **Three-mode data**





ee-mode data

nbining results

Examples

omatography E-M solution **MI** solutions

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sing data l estimation tiple Imputation

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## **Missing data**

- ee-mode data sing data many types creation origin procedures I estimation tiple Imputation nbining results mples Chromatography *Child development* omatography E-M solution MI solutions
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slices





linked modes (Harshman)

# **Creation of missing data**





#### **Missing values**

- -Second-order signals
- Light scattering
- Detector out of range

Source data:KVL, Bro & Ander

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300

Excitation [nm]

250

# **Origin of missing data**

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#### sing data

- many types creation
- origin
- procedures
- estimation
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#### • Missing completely at random

- Data are missing because of a random generating process
- Cause of missingness is unrelated to the variable with the missing data
- Deleting cases with missing data has no influence on representativeness, but diminishes power

#### Missing at random

- Cause of missing is systematic and correlated with the variable containing the missing data.
- Cause is accessible and can be included in the analysis to correct for bias

#### Missing not at random

- Cause of missing is systematic and correlated with the variable containing the missing data. Often the variable is the cause itself and thus the cause not accessible
- Cause is not accessible and cannot be included in the analysis to correct for bias

Little & Rubin (1987). *Statistical analysis with missing data*. Wiley; Schafer(1997). *Analysis of incomplete multivariate data*. Chapman & Hall

### **Procedures**

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- Expectation-Maximisation (EM) via three-mode model:

Estimate the missing data during iterations to determine the estimates of the model parameters

• Multiple imputation via data augmentation: Create several data sets with different values for the missing data and analyse each of them with a three-mode model, then combine the results



# **E(xpectation)-M(aximization)**

- ee-mode data sing data I **estimation**
- Tucker3
- Limitations
- tiple Imputation
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### **Tucker3 Model:** $x_{ijk} = \sum_{p} \sum_{q} \sum_{r} a_{ip} b_{jq} c_{kr} g_{pqr} + e_{ijk}$

- 1. *Tuckals*: Express <u>G</u> in A, B, C and <u>X</u>; *Gepcam*: Skip this step
- 2. (Preprocess: centre and normalise)
- 3. Find reasonable *starting values* for **A**, **B**, **C(, G)** and for missing data.
- 4. Estimate model parameters of three-mode model
- 5. Estimate *missing values* using model parameters
- 6. (Recentre and renormalise)
- 7. Iterate till convergence

Eigenvalue-eigenvector based (Kroonenberg & De Leeuw) Regression based (Weesie & Van Houwelingen - Gepcam). Missing data estimates are continously updated.

# **E(xpectation)-M(aximization)**

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- nples Chromatography *Child development* matography E-M solution MI solutions earch programme & ussion

### Limitations

- Single imputation
- Missing data estimates are tailored to the model.
- Model fits the (augmented) data too well
  - Underestimation of sampling variability
  - No estimate of uncertainty due to missing data
  - Missing data estimates have no sampling errors



## **Multiple imputation: Basics**

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Multiple imputation is a Monte Carlo technique in which missing data are replaced by m > 1 simulated versions, where *m* is typically small, say 3 - 10.





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# **Multiple imputation: Basics**

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- Validity imputations depends on the method of generation of the imputations
- Often normality of the original scores assumed

#### Rubin:

- Specify a parametric model for the complete data
- Apply a prior distribution to unknown model parameters
- Simulate *m* independent draws from conditional distribution of missing values given the observerd ones by Bayes' theorem



# **Multiple imputation: Generation**

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#### • iterative two-step process:

- alternatingly sample missing values from their conditional predictiv distribution
- then sample unknown parameters from a simulated complete-data posterior distribution.
- given initial values of the parameters this defines a Markov chair which converges to a stationary distribution of the missing values and the parameters, given the observed data
- iteration produces a draw of the parameters from its observed day posterior distribution and a draw of the missing values from the distribution of the missing values given the observed ones



\*Description taken from Schafer, J. L. (1999) in Stat. Meth. Med. Res., 8, 3-15

Multiple imputation and three-mode analysis

# **Multiple imputation: Three-way**

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Long matrix



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#### Wide matrix

- less data per variable,
- means and variance per *jk* taken into consideration (means - ok, variances - not?; see preprocessing)
- problematic if missing columns jk

#### Long matrix

- more data per variable
- mixtures of distributions (means confounded, variances - ok?)
- missing column = missing slice =>delete it
   Special procedures necessary?

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# **Multiple imputation: Stochastics**

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### • With sampling framework

cases x variables x conditions

### • Without sampling framework (single observation (or mean) per cell)

- varieties x attributes x locations
- wavelengths x wavelengths x concentrations
- solutes x eluents x adsorbents
- Distributional assumptions for multiple imputation valid?
- Estimate missing values some way and add normal error distributions per cell of three-way array with external standard errors for parameters to create multiple data sets? (add measurement error)

# Multiple data sets, multiple solutions

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- nbining results
- Subjects,variables Procrustes
- mples
- Chromatography *Child development* omatography E-M solution
- MI solutions
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- 10 imputed data sets
- 10 Tucker3 (Parafac) solutions
  - 10 Solutes component spaces
  - 10 Adsorbents component spaces
  - 10 Eluents component spaces
  - 10 Fit measures

### How to combine it all?

Standard MI - per parameter standard errors



### Here: Invariant subspaces with rotatable axes



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- Generalised Prokrousthj analysis on all imputed spaces
  - including the E-M solution
  - only imputed data, fit E-M solution into the centroid space for comparison
- First E-M solution and use that solution as target from imputed data: Target rotations



# Matching spaces via Generalised Prokrousthj analysis

Α



A. Translation B. Reflection C. Isotropic Scaling D. Rotation





First find iteratively a centroic then determine the optimal transformation to the centroic



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Options Procrustes

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E-M solution MI solutions

Child development

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#### ee-mode data sing data I estimation tiple Imputation nbining results **mples** Chromatography *Child Development* omatography E-M solution MI solutions earch programme &

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### Chromatography

- Data from De Ligny et al.
- Liquid chromatography

### Child development

- Data from the child care study of the NICHD
- Development in family background variables



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# Chromatography

### De Ligny, Spanjer, et al.



# Liquid chromatography

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1st mode: Solutes - monosubstituted phenols, anilines, pyridines
2nd mode: Stationary phase = adsorbents
3rd mode: Mobile phase = eluents
Measurement: Retention rate = log (net retention volume)/weight of absorbent

Source picture: http://falcon.sbuniv.edu/~ggray/CHE3345/chp24.html

# Data De Ligny et al.

### Data

- Dependent variable: Retention rate in High Performance Liquid Chromatography (HPLC)
- 39 solutes (bisubstituted benzenes) x 3 adsorbents x 2 eluents
- 21 missing data (= 9%); cause? retention too long?
- 5 rows for the 1st eluent have 1 valid and 2 missing observations. No missing for 2nd eluent.
- No preprocessing (=> 1st components primarily means)

### Purpose of the orginal analysis (De Ligny et al.)

Get estimates for the missing data, but not today **Structure** is also interesting; present focus.

### Question

How does the presence of the missing data influence the relationships between solutes?

Source data: De Ligny, C.L., et al. (1984). Journal of Chromatography, 301,311-323



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### **Data description**

3CI

3Br

**Pyridines** 

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<i>р</i> вг	<i>p</i> Br		
mCH3	mCH3		-
<i>р</i> СН3	<i>p</i> CH3	4CH3	-
mOCH3	mOCH3		-
pOCH3	<i>р</i> ОСН3		
mNO2	mNO2		
<i>p</i> NO2	<i>p</i> NO2		_
<i>m</i> CN	<i>m</i> CN	3CN	-
<i>p</i> CN	<i>p</i> CN	4CN	-
mCOOCH			_
<i>р</i> СООСН			
mCOCH3	mCOCH3		_
pCOCH3	<i>р</i> СОСН3		

**Solutes** 

Anilines

mF

pF

**m**Cl

pCl

*m*Br

nBr

**Phenols** 

mF

pF

**mCl** 

pCl

**m**Br

nBr



Adsorbents = stationary phase

Octadecyl-silica N-cyanoethyl-N-methylamino-silicia Aminobutyl-silica

Eluents = mobile phase

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35 v/v% methylene chloride in n-hexar pure methylene chloride



### **E-M solution**

### Parameter estimation via

- Tuckals algorithm -- eigendecomposition-based
- Gepcam algorithm -- regression-based
- Gepcam slightly more stable with very high fit

### Solution:

- no preprocessing all means included
  - 3 solutes components
  - 2 adsorbants components
  - 2 eluent components (K = R)
  - Proportion fitted sum of squares =
    - .9978 -- based on valid data
    - .9984 -- SS(Total) includes estimates missing data



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### **Solutes 1st component**



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## **Solutes - 2nd and 3rd components**

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### Joint plot for solutes and adsorbents

(First (consensus) component eluents - 97.4%)





Anilines & Pyridines tend to have relatively longer retention rates for both eluents with N-Cyanoethyl and Octadecyl silicates, while the Phenols tend to have relatively longer retention rates with Aminobutyl silicate, especially those with substituents containing nitrogen.

### **Proportional fit - MI & E-M** (Components)

l estimation tiple Imputation nbining results mples Chromatography			Minimum	Maximum	Mean	Std. Deviation	E-M	
Child development	Total		.993	.997	.995	.0013	.998	
E-M solution	Solutes							
MI solutions		1	.914	.928	.922	.0041	.928	
Fit		2	.052	.055	.054	.0013	.057	
Configurations To do		3	.016	.025	.020	.0034	.014	
d development	Adsorbents	5						
earch programme &		1	.937	.946	.942	.0023	.942	
cussion		2	.051	.056	.053	.0013	.057	
	Eluents							
		1	.964	.972	.968	.0029	.974	
		2	.025	.030	.027	.0020	.024	



e-mode data

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E-M is generally higher because no error for missing data

### **Optimally matched configurations** (Variability due to imputed values)



Points coloured by solutes; arrows start in mean configuration; bold = 2 missing & 1 valid value



#### Measure for stability of solutes (cases):

Sum (Average) of the squared distances to centroid (or E-M solution)

24/05/06 11:21:49

# **Unfinished business**

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# Compare estimates missing data and their standard errors for:

- E-M solution 3x2x3-solution (De Ligny et al.)
- E-M solution 3x2x2-solution
- Multiple imputation estimates
- Estimated data values from the analyses of the 10 imputated data sets
- Evaluate the location of E-M solution with respect to the solutions of imputed data sets



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# Child Development

### NICHD

#### The National Institute of Child and Human Development Study of Early Child Care and Youth Development



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### Study of Early Child Care and Youth Development

The SECC is a large longitudinal study started in 1989 to answer all kinds of questions with respect to the effects of child care.

Origin of the samples





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Data

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Missing erns

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E-M results Fit results

Variable space

**NICHD-SECC** 

Variable means

http://secc.rti.org

### **Data description**

#### Present subset of the roughly 1300 families:

- 150 Afro-American families
  - 11 Variables (see next slide)
  - 4 Points in time: 6, 15, 24, 36 months after birth

#### **Purpose of the analysis**

Determining the structure of the family situation and its changes in the first three years after birth of the baby.

### Questions

How does the presence of the missing data influence the relationships between variables? Does the structure of the variables change over time?

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**NICHD-SECC** 

Variable means

## **Missing data**

ee-mode data sing data l estimation tiple Imputation nbining results **Id development** NICHD -SECC Data description Variable means **Missing patterns** E-M results Fit results Variable space earch programme & cussion

										Mis	sing	Patte	erns									
Number of Cases	HealthBaby 36	HealthMother 36	SatisfiedWork 36	HoursWork/Week 36	SocialSupport 36	FinanacialResources 36	Maternal Deprression 36	Parenting 36	LogTotalIncome 36	HealthMother 24	SatisfiedWork 24	HealthBaby 24	HoursWork/week 24	LogTotalIncome 24	MaternalDepression 24	FinancialResources 24	SocialSupprt 24	Parenting 24	LogIncome/NeedRatio 36	LogIncome/NeedRatio 24	LogIncome/NeedRatio 15	LogIncome/NeedRatio 6
5																				X		
3																				Χ		Χ
5																						Χ
5																					X	Χ
15																			Χ	Χ	Χ	Χ
10															X	X	X	X				
7	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	Χ	Χ		

Patterns with less than 2% cases (2 or fewer) are not displayed.



# Variables & their means over time

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e-mode data

	1 (240) · (25) (25)	23 22222	m	onths		
Abbreviation	Description	06	15	24	36	
HrWrkM-xx	Hours/week mother works-all jobs	17.1	21.3	21.0	20.9	1
Satisf-xx	Mom satisfied with own work schedule	3.6	3.8	3.6	3.5	
Depres-xx	Maternal depression	11.9	11.2	13.1	11.9	
Suppor-xx	Social Support	5.0	4.8	4.6	4.7	
PStres-xx	Parenting stress <sup>*</sup>	51.0	34.3	35.7	34.7	
$\operatorname{HealtM-xx}$	Health of mother	3.2	3.1	3.0	2.9	
HealtB-xx	Health of baby	3.3	3.1	3.2	3.2	
HrCare-xx	Hours/week in care	23.6	26.1	24.4	26.8	
Financ-xx	Financial resources	9.3	9.3	9.2	9.4	
Income-xx	Log total income	9.7	9.7	9.8	9.9	
Need -xx	Log income to need ratio	.3	.2	.4	.4	

xx = 06, 15, 24 or 36; indicating observed in the xx month after birth.



\*different instrument at 6 months

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### **E-M results**

#### **Fundamental results**

- Number of components for a Tucker3 model:
  - 3 (subjects) x 3 (variables) x 1 (time)
- The coefficients are virtually equal for the four time points: Structure variables hardly changed over time.
- We might as well average over time points:. Tucker3 analysis is then equivalent to an SVD (PCA) on the subject-x-variable matrix averaged over time.
- Multiple imputation over wide matrix, thereafter standard preprocessing ~

 $\widetilde{x}_{iik} = (x_{iik} - \overline{x}_{ik}) / s_i$ 



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### **Fit results**

### 5 Imputed data sets

Solution	SS(Fit)	•	ortion		
		1	<b>2</b>	3	
Base solution	.415 (.382)	.229	.123	.063	
Equal weights	.414 (.381)	.228	.122	.063	evoluting missing values from SS(Total)
Time component					excluding missing values from SS(Total)
Imputation 1	.388	.220	.109	.059	<ul> <li>including missing values from SS(Total)</li> </ul>
Imputation 2	.383	.216	.110	.058	
Imputation 3	.380	.211	.113	.056	
Imputation 4	.382	.212	.111	.059	
Imputation 5	.390	.218	.113	.058	

First components of E-M explain relatively more; probably due to the tailoring of missing data to the model (to be seriously investigated)



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### **Variables spaces**

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Income/need ratio had 183 missing compared to Stress, Support, Depression with around 50.

# **Research programme**

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### • Large scale questions

- Multiple imputation via wide or long matrix?
- Multiple imputation and means, standard deviations, and recommended preprocessing, i.e. three-way multiple imputation?
- Multiple imputation and lack of stochastics in three-way data? Use external information, e.g. standard deviations from earlier studies, in multiple imputations?
  - Rotation to a target (=E-M solution) rather than to centroid?



Comments

# Some (random?) comments

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"I must add that even doing multiple imputation relatively crudely, using simple methods, is very likely to be inferentially far superior to any other equally easy method to implement (e.g., complete-cases, available cases, single imputation, Last Value Carried Forward) because the multiple copies of the data set allow the uncertainty about the values of the missing data to be incorporated into the final inferences;"

Rubin on www.statsol.ie/solas/rubin1.htm

The results suggest a reliable and efficacious basis for *imputation method for repeated measures data* is to substitute a missing datum with a value from another individual who has the closest scores on the same variable measured at other timepoints, or the average value of four individuals who have the closest scores on the same variable at other timepoints.

Elliott P, Hawthorne G. Aust N Z J Psychiatry. 2005 Jul;39(7):575-82.

### **A final comment**

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"Analysing data that you do not have is so obviously impossible that it offers endless scope for expert advice on how to do it."

Ranald R. MacDonald, University of Stirling,UK. www.psychology.stir.ac.uk/staff/rmacdonald/Missing.htm; seen 30/8/2005

