

On Model Selections for Repeated Measurement Data in Clinical Research

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- Introduction
- Methods
- Simulation Study
- Application Example
- Discussions

Repeated Measurement Design

- Repeated Measurement Design:
 - Multiple responses of each experimental unit are collected at different time points or under different conditions

- Data Structure:

ID	Group	Time (month)	Outcome
1	0	0	2
1	0	6	4
1	0	12	9
2	1	0	5
2	1	6	3
2	1	12	4
⋮	⋮	⋮	⋮

- Applications
 - long term intervention efficacy
 - risk factors of chronic disease
- Benefits
 - experiments efficiencies
- Disadvantages
 - correlated responses
 - missing data

Repeated Measurement Design (cont.)

- Data Structure with Missing:

ID	Group	Time (month)	Outcome
1	0	0	NA
1	0	6	4
1	0	12	9
2	1	0	5
2	1	6	3
2	1	12	NA
3	1	0	6
⋮	⋮	⋮	⋮
⋮	⋮	⋮	⋮

Mixed Effects Model (Laird & Ware 1982)

- Takes longitudinal information and correlations among repeated measurements into account.
- Allow to evaluate and test the overall effect across every time point and the treatment effect at a fixed time point.
- Non-missing observations can help partially recover the lost information in the missing data.

- Full Model:

$$Y_{ijk} = \alpha_0 + \alpha_{Tt}I(i = 1) + \sum_{k=1}^{T-1} \alpha_k I(\text{Time} = k) + \sum_{k=1}^{T-1} \alpha_{k+T-1} I(i = 1) I(\text{Time} = k) + r_{ijk} + \epsilon_{ijk} \quad (1)$$

- i : treatment assignment (1 for treated and 0 for untreated)
- j : indexes an individual subject
- k : measurement time point post baseline measurement ($=1, \dots, T$)
- Y_{ijk} : response change from baseline

Mixed Effects Model (cont.)

- Full Model:

$$Y_{ijk} = \alpha_0 + \alpha_{Tt}I(i = 1) + \sum_{k=1}^{T-1} \alpha_k I(\text{Time} = k) + \\ + \sum_{k=1}^{T-1} \alpha_{k+T-1} I(i = 1) I(\text{Time} = k) + r_{ijk} + \epsilon_{ijk}$$

- α_0 : intercept.
- α_{Tt} : treatment effect (at last time point).
- $\alpha_k (k = 1, \dots, T - 1)$: time effects.
- $\alpha_{k+T-1} (k = 1, \dots, T - 1)$: time and treatment interactions.
- r_{ijk} : cluster effects, i.e. $\mathbf{r}_{ij} \sim N(\mathbf{0}, \mathbf{\Sigma})$ with $T \times T$ dimension.
- $\epsilon_{ijk} \sim N(0, \sigma^2)$: random error and $\epsilon_{ijk} \perp r_{ijk}$

- Covariance Structure:

Unstructured Covariance Matrix:

$$\Sigma = \begin{pmatrix} \sigma_1^2 & \sigma_{12} & \cdots & \sigma_{1T} \\ \sigma_{21} & \sigma_2^2 & \cdots & \sigma_{2T} \\ \vdots & \vdots & \ddots & \vdots \\ \sigma_{T1} & \sigma_{T2} & \cdots & \sigma_T^2 \end{pmatrix}$$

$\frac{T(T-1)}{2}$ covariance parameters, could be unstable

Compound Symmetry Covariance Matrix:

$$\Sigma = \sigma^2 \begin{pmatrix} 1 & \rho & \cdots & \rho \\ \rho & 1 & \cdots & \rho \\ \vdots & \vdots & \ddots & \vdots \\ \rho & \rho & \cdots & 1 \end{pmatrix}$$

2 covariance parameters, strong assumptions

- Main Effects Model:

$$Y_{ijk} = \alpha_0 + \alpha_{Trt}I(i = 1) + \sum_{k=1}^{T-1} \alpha_k I(\text{Time} = k) + r_{ijk} + \epsilon_{ijk} \quad (2)$$

- α_0 : intercept.
- α_{Trt} : treatment effect.
- $\alpha_k (k = 1, \dots, T - 1)$: time effects.
- r_{ijk} : cluster effects, i.e. $\mathbf{r}_{ij} \sim N(\mathbf{0}, \mathbf{\Sigma})$ with $T \times T$ dimension.
- $\epsilon_{ijk} \sim N(0, \sigma^2)$: random error and $\epsilon_{ijk} \perp r_{ijk}$

- Null Model:

$$Y_{ijk} = \alpha_0 + \sum_{k=1}^{T-1} \alpha_k I(\text{Time} = k) + r_{ijk} + \epsilon_{ijk} \quad (3)$$

- α_0 : intercept.
- $\alpha_k (k = 1, \dots, T - 1)$: time effects.
- r_{ijk} : cluster effects, i.e. $\mathbf{r}_{ij} \sim N(\mathbf{0}, \mathbf{\Sigma})$ with $T \times T$ dimension.
- $\epsilon_{ijk} \sim N(0, \sigma^2)$: random error and $\epsilon_{ijk} \perp r_{ijk}$

- Overall effects:

$$H_0 : \alpha_{Tt} = \alpha_T = \alpha_{T+1} = \cdots = \alpha_{2T-2} = 0 \text{ vs } H_a : \text{Otherwise.}$$

- Effects at a fixed time point (usually at last time point):

$$H_0 : \alpha_{Tt} = 0 \text{ vs } H_a : \alpha_{Tt} \neq 0$$

- Overall Effects:

- likelihood ratio test:

$$\zeta_{LRT} = -2 \log \frac{\text{likelihood of null model (3) with UN}}{\text{likelihood of full model (1) with UN}} \sim \chi^2$$

- Effects at Last Time Point:

- t-test: $\zeta_T = \frac{\hat{\alpha}_{Trt,T}}{SE(\hat{\alpha}_{Trt,T})}$

- full model (1) with UN: $\zeta_{FUN} = \frac{\hat{\alpha}_{Trt,FUN}}{SE(\hat{\alpha}_{Trt,FUN})}$

- Candidate Model Space:
 - Full and Main Effects Models
 - Covariance Structures
 - compound symmetry (CS)
 - autoregressive (AR)
 - unstructured (UN)
 - Candidate Models: full and main effects model with covariance structures
- Model Selection Criteria: BIC

Test Statistics for Treatment Effects

- Overall Effects:
 - model selection test statistics

$$\zeta_{MSA} = \begin{cases} \tilde{\alpha}' \Sigma^{-1} \tilde{\alpha} & \text{if } \hat{M} \in \text{full model set (1)} \\ (\tilde{\alpha}_{Tt} / SE(\tilde{\alpha}_{Tt}))^2 & \text{if } \hat{M} \in \text{main effects model set (2)} \end{cases}$$

$\tilde{\alpha}' = (\tilde{\alpha}_{Tt}, \tilde{\alpha}_T, \dots, \tilde{\alpha}_{2T-2})$ is the MLE of $\alpha' = (\alpha_{Tt}, \alpha_T, \dots, \alpha_{2T-2})$ from the optimal model \hat{M} if it happens to be the full model (1), while $\tilde{\alpha}_{Tt}$ is the MLE of α_{Tt} from the optimal model \hat{M} when it happens to be the main effects model (2)

Σ is the covariance matrix of $\tilde{\alpha}'$

- Effects at Last Time Point:

- model selection test statistics: $\zeta_{MSL} = \frac{\tilde{\alpha}_{Trt, \hat{M}}}{SE(\tilde{\alpha}_{Trt, \hat{M}})}$

- How to estimate Σ ?
 - Estimate of Σ is not trivial due to stochastic nature of model selection
 - Distribution of selecting a specific candidate model as the optimal model \hat{M} is unknown \Rightarrow bootstrapping

Restricted Cluster Bootstrapping Estimate of Σ

- Simple bootstrap resampling does not work
- Restricted cluster bootstrap resampling

Step 1: Perform model selection on the observed dataset, D , to get the optimal model \hat{M} .

Step 2: Conduct **cluster level** resampling with replacement to get a resampled dataset and perform model selection on the resampled dataset D^b . Obtain the selected optimal model \hat{M}^b . If $\hat{M}^b = \hat{M}$, then the MLE of α , i.e. $\tilde{\alpha}^b$, is used for the calculation of $\hat{\Sigma}_B$.

Step 3: Repeat Step 2 B times and calculate $\hat{\Sigma}_B$, i.e. the post-model selection covariance matrix estimate of $\tilde{\alpha}$.

$$\hat{\Sigma}_B = \frac{1}{B^* - 1} \sum_{b=1}^B (\tilde{\alpha}^b - \bar{\alpha}^b) \otimes (\tilde{\alpha}^b - \bar{\alpha}^b)' I(\hat{M}^b = \hat{M})$$

$$\bar{\alpha}^b = \frac{1}{B^*} \sum_{b=1}^B \tilde{\alpha}^b I(\hat{M}^b = \hat{M}).$$

$$B^* = \sum_{b=1}^B I(\hat{M}^b = \hat{M}).$$

- data model (no missing)

$$\begin{aligned}y_{ijk} = & -70 + 0.3 * I(i = 1) + 4 * I(k = 1) + 3 * I(k = 2) \\ & + 2 * I(k = 3) + I(k = 4) + \alpha_1 * I(i = 1)I(k = 1) \\ & + \alpha_2 * I(i = 1)I(k = 2) + \alpha_3 * I(i = 1)I(k = 3) \\ & + \alpha_4 * I(i = 1)I(k = 4) + r_{ijk} + \epsilon_{ijk}\end{aligned}$$

- strong: $\alpha_1 = 0.45, \alpha_2 = 0.42, \alpha_3 = 0.39, \alpha_4 = 0.36$
- moderate: $\alpha_1 = 0.24, \alpha_2 = 0.27, \alpha_3 = 0.33, \alpha_4 = 0.36$
- weak: $\alpha_1 = 0.1, \alpha_2 = 0.075, \alpha_3 = 0.05, \alpha_4 = 0.025$
- no interaction
- random cluster effect $r_{ij} \sim N(0, \mathbf{\Omega})$
- $\epsilon_{ijk} \sim N(0, 1)$

Covariance Structure (CS):

$$\Sigma = \begin{pmatrix} 1 & 0.4 & 0.4 & 0.4 & 0.4 \\ & 1 & 0.4 & 0.4 & 0.4 \\ & & 1 & 0.4 & 0.4 \\ & & & 1 & 0.4 \\ & & & & 1 \end{pmatrix}$$

Table 1: Overall Treatment Effects Testing^a

Data Model	Interaction	Statistics ^b	Power	Type I Error
Full (CS)	Strong	ζ_{LRT}	0.996	0.044
		ζ_{MSA}	1.000	0.048
	Moderate	ζ_{LRT}	0.967	0.044
		ζ_{MSA}	0.995	0.048
	Weak	ζ_{LRT}	0.621	0.044
		ζ_{MSA}	0.834	0.048
Main Effects (CS)	None	ζ_{LRT}	0.479	0.044
		ζ_{MSA}	0.710	0.048

a: results based on 1000 simulations with 100 subjects at each treatment arm

b: model selection via BIC with 200 bootstrap resampling

ζ_{LRT} : mixed effects model based likelihood ratio test for overall effects

ζ_{MSA} : model selection statistics for testing overall treatment effects

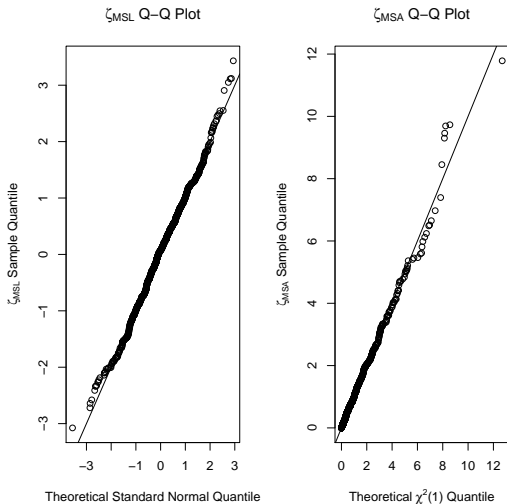
Simulation Results (Cont.)

Table 2: Last Time Point Treatment Effect Testing^a

Data Model	Interaction	Statistics ^b	Power	Type I Error
Full (CS)	Strong	ζ_T	0.336	0.044
		ζ_{FUN}	0.343	0.046
		ζ_{MSL}	0.981	0.048
	Moderate	ζ_T	0.336	0.044
		ζ_{FUN}	0.343	0.046
		ζ_{MSL}	0.987	0.048
	Weak	ζ_T	0.336	0.044
		ζ_{FUN}	0.343	0.046
		ζ_{MSL}	0.834	0.048
Main Effects (CS)	None	ζ_T	0.336	0.044
		ζ_{FUN}	0.343	0.046
		ζ_{MSL}	0.710	0.048

Simulation Results (Cont.)

Figure 1: Model Selection Test Statistics Distributions Under the Null



- Data Model (missing)

$$\begin{aligned}y_{ijk} = & -70 + 0.3 * I(i = 1) + 0.3 * u_{ij} + 4 * I(k = 1) \\ & + 3 * I(k = 2) + 2 * I(k = 3) + I(k = 4) \\ & + \alpha_1 * I(i = 1)I(k = 1) + \alpha_2 * I(i = 1)I(k = 2) \\ & + \alpha_3 * I(i = 1)I(k = 3) + \alpha_4 * I(i = 1)I(k = 4) \\ & + r_{ijk} + \epsilon_{ijk}\end{aligned}$$

- $u_{ij} \sim Unif(-1, 1)$.
- moderate: $\alpha_1 = 0.24, \alpha_2 = 0.27, \alpha_3 = 0.33, \alpha_4 = 0.36$
- no interaction
- random cluster effect $\mathbf{r}_{ij} \sim N(0, \mathbf{\Omega})$
- $\epsilon_{ijk} \sim N(0, 1)$

Simulation Studies (cont.)

- Missing Mechanism:
 - to mimic controlled trial scenario: e.g. older age subjects in the treated group may miss measurements more frequently due to drug side-effects while younger subjects in the placebo group may miss measurement more frequently due to inefficacy of treatment, etc.
 - for the treatment arm 0 and one of the 1st four time points k ($k = 1, 2, 3, 4$), five subjects with $u_{ij} > 0$ are randomly selected to have their observations set to missing, while for the treatment arm 1, 5 subjects with $u_{ij} < 0$ are randomly selected to create missing observations. For the last time point, we randomly select 10 subjects in the treatment arm 0 with $u_{ij} > 0.5$ and another 10 subjects from the treatment arm 1 with $u_{ij} < -0.5$ and set their observations to missing.
 - instead of imputing missing data, we analyze the simulated data with models (1), (3) and (2) except that we add baseline u_{ij} as a covariate into these models.

Covariance Structure (CS):

$$\Sigma = \begin{pmatrix} 1 & 0.4 & 0.4 & 0.4 & 0.4 \\ & 1 & 0.4 & 0.4 & 0.4 \\ & & 1 & 0.4 & 0.4 \\ & & & 1 & 0.4 \\ & & & & 1 \end{pmatrix}$$

- As a further comparison, when testing the last time point effect, we include the ANCOVA model test statistics $\zeta_{ANCOVA} = \frac{\hat{\alpha}_{Trt}}{SE(\hat{\alpha}_{Trt})}$ where $\hat{\alpha}_{Trt}$ is the MLE from the following ANCOVA model: $y_{ijT} = \alpha_0 + \alpha_{Trt}I(i = 1) + \alpha_u u_{ij} + \epsilon_{ij}$.

Table 3: Simulation Results with Missing^a

Data Model	Interaction	Statistics ^b	Hypothesis Test	Power ^c	Type I Error ^c
Full (CS)	Moderate	ζ_T	Last Time Point	0.193(0.328)	0.060(0.047)
		ζ_{ANCOVA}		0.285(0.327)	0.064(0.044)
		ζ_{FUN}		0.310(0.343)	0.049(0.049)
		ζ_{MSL}		0.989(0.987)	0.054(0.046)
		ζ_{LRT}	Overall Effect	0.968(0.965)	0.054(0.044)
		ζ_{MSA}		0.995(0.995)	0.054(0.046)

a: results based on 1000 simulations with 100 subjects at each treatment arm

b: model selection via BIC with 200 bootstrap resampling

c: values inside parentheses are based on no missing scenario

ζ_T : t-test for last time point effect

ζ_{ANCOVA} : test last time point effect via ANCOVA model

ζ_{FUN} : test last time point effect via full mixed effect model with UN covariance

ζ_{MSL} : model selection test statistics for testing last time point effect

ζ_{LRT} : mixed effects model based likelihood ratio test for overall effects

ζ_{MSA} : model selection statistics for testing overall effects

Simulation Results (Cont.)

Table 4: Simulation Results with Missing^a

Data Model	Interaction	Statistics ^b	Hypothesis Test	Power ^c	Type I Error ^c
Main Effects (CS)	None	ζ_T	Last Time Point	0.176(0.325)	0.060(0.047)
		ζ_{ANCOVA}		0.265(0.331)	0.064(0.044)
		ζ_{FUN}		0.303(0.338)	0.049(0.049)
		ζ_{MSL}	0.689(0.710)	0.054(0.046)	
		ζ_{LRT}	Overall Effect	0.463(0.486)	0.054(0.044)
		ζ_{MSA}		0.689(0.710)	0.054(0.046)

a: results based on 1000 simulations with 100 subjects at each treatment arm

b: model selection via BIC with 200 bootstrap resampling

c: values inside parentheses are based on no missing scenario

ζ_T : t-test for last time point effect

ζ_{ANCOVA} : test last time point effect via ANCOVA model

ζ_{FUN} : test last time point effect via full mixed effect model with UN covariance

ζ_{MSL} : model selection test statistics for testing last time point effect

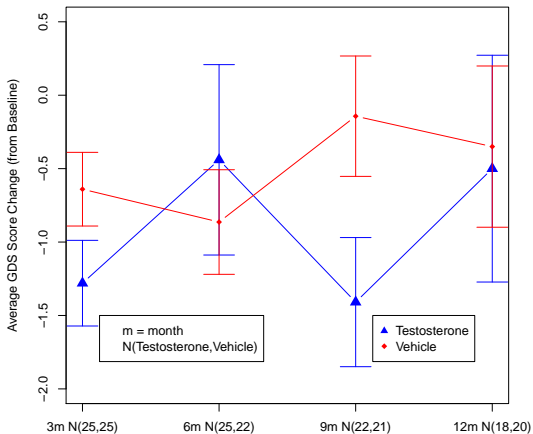
ζ_{LRT} : mixed effects model based likelihood ratio test for overall effects

ζ_{MSA} : model selection statistics for testing overall effects

- Testosterone's Cognitive Effect (Borst et. al. 2014)
 - 60 subjects
 - two treatment arms: testosterone vs vehicle
 - subject randomly assigned
 - outcome: geriatric depression scale (GDS: 0 ~ 30)
 - higher score means more depressed
 - repeatedly measured at baseline, 3, 6, 9, and 12 months

Real Application Example (Cont.)

Figure 2: Average GDS Score Change (from Baseline) Comparison



Real Application Example (Cont.)

- time by treatment interaction strong but not significant (p-value=0.10)
- overall effects test by ζ_{LRT} : p-value=0.04
- overall effects test by ζ_{MSA} :
 - model selection \Rightarrow optimal model = full model with unstructured covariance
 - p-value=0.04
- tested by ζ_T with last time point info only: p-value=0.87

- Proposed statistics
 - make use of information from optimal model deemed for the data \Rightarrow powerful for repeated measurement data.
 - utilize the information across each measurement time point \Rightarrow more robust and flexible for missing data.
- Restricted cluster bootstrapping \Rightarrow valid covariance estimate \Rightarrow valid statistical inference.
- Extension to other data types deserves further investigations.

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THANK YOU