

Longitudinal Data Analysis

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Repeated Measures and Longitudinal Data

- In repeated measures designs, measurements are taken repeatedly on each individual
- When these measurements are taken over time, it is called *longitudinal* design
- Typically various covariates concerning the individuals are recorded
	- 3 types of predictors
		- Time (or some notion thereof)
		- Time-variant
		- Time-invariant
- A general goal is to determine how a response depends on the covariates over time

Studying Change over Time

- Study can be experimental or observational
- Data can be collected prospectively or retrospectively
- Time can be measured in a variety of units (e.g. weeks, months, years, semesters, sessions, etc.)
- Nonetheless, need three or more waves of data
	- More waves is always better. If your data has only three waves, you must fit simpler models with stricter assumptions (e.g. linear). Additional waves allow for more flexible models with less restrictive assumptions (e.g. curvilinear).
	- There is nothing sacrosanct about equal spacing. If you expect rapid nonlinear change during some time periods, you should collect more data at those tmes.
	- The resultant data need not be balanced. In other words, each person need not have the same number of waves.

Traditional Methods have Limitations

- Traditional techniques for longitudinal data analysis, such as repeated measures ANOVA, can handle only complete data cases
- The assumptions of repeated measures ANOVA, in particular the assumption of sphericity/circularity, are often too restrictive for longitudinal data
	- Circularity refers to the conditon where the variances of the diferences between all pairs of within-subject conditons are equal
	- Departure from circularity results in infated F-ratos
- Far less flexible in handling complex data structures compared to mult-level / mixed-efects modeling

Multi-level / Mixed-effects Modeling of Longitudinal Data

- The mixed model approach to analyzing longitudinal data was commenced with the paper of Laird and Ware (1982)
- Most of the work that has been undertaken to model longitudinal data has been parametric, in the sense that the effects of continuous covariates have been modeled linearly or by using some parametric nonlinear model
- An alternative to nonlinear mixed modeling is to incorporate smoothing methods (aka semiparametric mixed models) \rightarrow We'll come back to this after learning about additive models

Example #1: Tolerance data set

- National Youth Survey (NYS; Raudenbush & Chan 1992)
- Participants filled a survey at ages 11, 12, 13, 14, and 15 to provide their tolerance of deviant behavior
- Using a 4-point scale, where 1 = very wrong, 2 = wrong, 3 = a little bit wrong, and $4 = not$ wrong at all, participants indicated whether it was wrong for someone their age to
	- i. Cheat on tests
	- ii. Purposely destroy property of others
	- iii. Use marijuana
	- iv. Steal something
	- etc.
- Response (i.e. tolerance) was computed as respondent's average score
- Additional covariates
	- Gender (1=male and 0=female)
	- Exposure, representng respondent's self-reported exposure to deviant behavior at age 11, also on a 0-4 scale

Typical Workflow

"Tidy datasets are all alike but every untidy dataset is untidy in its own way." - Hadley Wickham

person-level person-period

tidying

Typical Workflow

A Framework for Analyzing Longitudinal Data

- Despite the unique set of outcome, predictors, and research questions, each longitudinal study poses an identical pair of questions:
	- **Q1) Within-individual change**
		- **How does the outcome change over time?**
		- E.g. how is each individual's pattern of tolerance over tme? Is individual change linear or non-linear? Is it consistent over tme or does it fuctuate?
	- **Q2) Inter-individual diferences in change**
		- " Can we predict differences in these changes?
		- E.g. do boys and girls experience different patterns of tolerance? Does baseline exposure afect boys and girls in the same way?

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- To assess the nature and idiosyncrasies of each person's temporal patern of change
- **STEP1: Start with scaterplots (response vs. time) at the individual level Q1**
	- Easier to discern if sets of individuals are ploted on the same panel
	- Better to use identical axes across individuals
	- In large data sets, may have to inspect a random subset of individuals

- **STEP2: Use a trajectory to summarize each person's temporal pattern of change**
	- Nonparametric approach (e.g. smoothing spline)
		- Requires no assumptions
		- Letting the data speak for themselves
	- $-$ Parametric approach (e.g. linear, quadratic, etc.)
		- Requires assumptions
		- But instead provides numeric summaries of the trajectories
- Exploratory analysis often suggests that different people require different functions
	- $-$ Measurement error makes it difficult to discern if individual patterns are true signal or simply due to random fluctuation
	- $-$ Of course, fitting person-specific models, one individual at a time, is not the most efficient use of longitudinal data; That's why we will use mixed-efects modeling eventually

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- Does everyone change in the same way, or do the trajectories of change differ substantially across people? **Q2**
- STEP3: Plot, on a single graph, the entire set of individual **trajectories, and compute an average change trajectory**
	- i. Discretize time on a reasonably refined grid
	- ii. Estmate individual trajectories on the grid
	- iii. Average individual estmates for each point on the grid
	- iv. Apply the same smoothing algorithm, nonparametric or parametric, used to obtain individual trajectories
- NEVER infer the shape of the individual change trajectories from the shape of their averages
	- $-$ The only kind of trajectories for which the "average of the curves" is identical to the "curve of the averages" is one whose mathematical representation is *linear in the parameters* [Keats 1983]
	- We examine the averages simply for comparison, not to learn anything about the underlying shapes of the individual trajectories

- **STEP4: Use the results of exploratory analysis to (re-)frame questions about change**
	- $-$ Adopting a parametric model for individual change allows us to express generic questions about inter-individual diferences in change as specifc questons about the parameters of the model
	- $-$ E.g. in the case of linear fit, it is often helpful to examine
		- i. Sample means of of the estmated intercepts and slopes
		- ii. Sample variances (or SD) of the estmated intercepts and slopes
		- iii. Sample correlation between the estimated intercepts and slopes

• **STEP5: Explore the relationship between change and time-invariant predictors**

- Allows us to uncover systematc paterns in the individual change trajectories corresponding to inter-individual variaton observed in personal trajectories, e.g.
	- Examining diferences by gender allows us to assess
		- Whether boys or girls are initally more tolerant of deviant behavior
		- Whether boys and girls tend to have diferent annual rates of change
	- Examining diferences by baseline exposure allows us to assess
		- Whether a child' inital level of tolerance is associated with baseline exposure
		- Whether a child' rate of change in tolerance is associated with baseline exposure
- $-$ If a predictor is continuous you can temporarily categorize it

- What about time-variant predictors?
	- $-$ For example via loess (with multiple predictors) or thin plate splines (from mgcv). See persp() in base R graphics for plotting 3D surfaces.

 0.2 0.4

 0.8

age

age

Typical Workflow

Coffee break? ^M

Recap of Mixed Modeling: ML vs. REML

- Choice of methods for parameter estimation
	- Maximum Likelihood (ML)
	- Restricted Maximum Likelihood (REML)
- The general wisdom is that
	- ML produces more accurate estimates of the fixed parameters, whereas REML produces more accurate prediction of random variances
	- Therefore, the choice depends on whether the hypotheses are focused on the fixed parameters or random variances
	- In practice, the choice of ML or REML will make only small diferences to the parameter estmates
	- If you want to compare models, you MUST use ML

Recap of Mixed Modeling: Model Selection

- Assessing the fit and comparing multi-level models
	- $-$ Likelihood ratio test based on ML theory
		- R reports the -2*log-likelihood (-2LL)
		- Essentially the smaller the value of LL the better
	- R also produces two adjusted LL values, which can be interpreted in a similar manner
		- AIC: Akaike's Information Criterion
			- A goodness of fit measure corrected for model complexity
		- BIC: Bayesian Information Criterion
			- Comparable to AIC but slightly more conservatve (i.e. it corrects more harshly for the number of parameters)
		- Neither AIC or BIC are intrinsically interpretable; they are only useful in comparing models. Smaller values in both cases mean a better-fitting model
		- Recommended approach: start simple!

- A key assumption of standard linear regression is the assumption of independently distributed error terms for the individual observations within a sample
	- $-$ Essentially means that there are no relationships among individual observations for the dependent variable once the independent variables are accounted for
- In real world, data often has a hierarchical structure, hence the name "multi-level"
	- It simply means that some variables are *clustered* or *nested* within other variables
	- In longitudinal design, this hierarchy (in part) stems from repeated measures obtained from the same individual over tme

- When dealing with multi-level data, the assumption of independent errors is violated
	- $-$ i.e. the <u>potential</u> inter-individual correlation (correlation among repeated measures obtained from the same individual) may result in inappropriate estmate of model parameters
- Moreover, by ignoring multi-level structure of data we may miss important relatonships involving each level in the data
- Solution ^E When data has multi-level structure, we allow the parameters of the model to vary between clusters
	- $-$ Fixed vs. random coefficients (mixed-effects modeling)
	- $-$ Packages for multi-level modeling in R
		- nlme (allows for flexible modeling of the covariance structure)
		- Ime4

Allowing the parameters of the model to vary between clusters:

Either allowing the intercepts to vary among children, or allowing the slopes to vary among children, or allowing both intercepts and slopes to vary among children.

Instead of estmatng 16 intercepts and 16 slopes, we will estmate one fxed intercept, one fixed slope, one random intercept and one random slope.

 $y_{ij} = \alpha + \varepsilon_{ij}$ for individual i at time j

 $y_{ij} = \alpha_0 + \varepsilon_{ij}$ (Model 00) α_0 is a fixed variable

 $y_{ij} = (\alpha_0 + \alpha_1) + \varepsilon_{ij}$ (Model 01)

 α_0 is a fixed variable whereas α_1 is a random variable

> anova(fit.00, fit.01) Model df logLik Test L.Ratio p-value **AIC BIC** fit.00 2 117.7198 122.4839 -56.85992 -51.51093 1 vs 2 10.698 116.1679 0.0011

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y_{ij} = \alpha + \varepsilon_{ij} \quad \text{for individual } i \text{ at time } j
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y_{ij} = (\alpha_0 + \alpha_1) + \varepsilon_{ij} \quad \text{(Model 01)}
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\alpha \text{ fixed } \text{as a single line } \alpha \text{ is a same domain } b \text{ and } c \text{ is a single line.}
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 α_0 is a fixed variable whereas α_1 is a random variable

$$
y_{ij} = \alpha + \varepsilon_{ij} \quad \text{for individual } i \text{ at time } j
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y_{ij} = (\alpha_0 + \alpha_1) + \varepsilon_{ij} \quad \text{(Model 01)}
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 α_0 is a fixed variable whereas α_1 is a random variable

Multi-level Modeling Approach to Longitudinal Data $y_{ij} = \alpha + \beta t_j + \varepsilon_{ij}$ for individual i at time j $y_{ij} = \alpha_0 + \beta_0 t + \varepsilon_{ij}$ (Model 03) α_0 and β_0 are fixed variables $y_{ij} = \alpha_0 + (\beta_0 + \beta_1)t_i + \varepsilon_{ij}$ (Model 04)

 α_0 and β_0 are fixed variables whereas β_1 is a random variable

> anova(fit.03,fit.04) Model df **AIC BIC** logLik Test L.Ratio p-value 3 107.55383 114.69991 -50.77692 $fit.03$ 1 -33.46633 1 vs $-.0001$ 2 34.62117 74 'nб

$$
y_{ij} = \alpha + \beta t_j + \varepsilon_{ij} \quad \text{for individual } i \text{ at time } j
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$$
y_{ij} = \alpha_0 + (\beta_0 + \beta_1)t_j + \varepsilon_{ij} \quad \text{(Model 04)}
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Summary of a Typical Workflow

- Assumptions about the covariance structure of the data
	- Variance components
	- Diagonal
	- $-$ AR(1) \equiv typically used when data is measured over time components
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		- $\text{-} \quad \text{corAR1}(...)$
		- corCAR1(…)
		- corARMA(…)

We

– Unstructured

Working Exercise #1: "Beat the Blues" clinical trial

- Depression is a major clinical problem
- Cognitive Behavioral Therapy (CBT) aka "talking therapy" seems to be as effective as drugs
- Problem is, there is a shortage of skilled therapists
- A clinical trial of an interactive multimedia program known as "Beat the Blues", aimed to assess the efects of the program compared to "Treatment as Usual" (TAU)
- More details in [Proudfoot et al. 2003, Psychological Medicine]
- Longitudinal design
	- –100 individuals
	- –Outcome measurements obtained at 5 tme points (pre, 2m, 3m, 5m, and 8m)
	- * Outcome recorded according to Beck Depression Inventory (BDI), a [0,63] score based on a 21-queston multple choice self-report questionnaire
	- –Covariates of interest
		- Taking antidepressant drugs (drug, YES or NO)
		- Duration of current episode of depression (length, less or more than 6 months)

Working Exercise #2: Bone Data

References

- Applied Longitudinal Data Analysis; by Judith D. Singer and John B. Willet; Oxford University Press
- Semiparametric Regression; by David Ruppert, M.P. Wand, and R.J. Carroll; Cambridge University Press
- Generalized Additive Models; by T.J. Hastie and R.J. Tibshirani; Chapman & Hall/CRC
- Generalized Additive Models An Introduction with R (2nd Editon); by Simon N. Wood; CRC Press

Lifetime Warranty "

If you have follow-up questions, need help with your future analysis, or simply want to stay in touch, feel free to contact me at:

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