Measuring Individual Change: A Gentle Introduction to the Pros and Cons of Modern Models

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Overview

- Organizing principles
 - > From one to many kinds of variance
- From cross-sectional to longitudinal (growth) models
 - > Multilevel/hierarchical linear/latent growth curve models
- Models for exploring "heterogeneous populations"
 - > Latent class/profile/transition/mixture models
- Models for confirming/testing heterogeneous groups
 - > Heterogeneous variance longitudinal models
 - Confirmatory longitudinal mixture models

The Two Sides of Any Model

Model for the Means:

- > Aka Fixed Effects, Structural Part of Model
- > What you are used to **caring about for testing hypotheses**
- How the expected outcome for a given observation varies as a function of values for known predictor variables

Model for the Variance:

- > Aka Random Effects and Residuals, Stochastic Part of Model
- > What you are used to **making assumptions about** instead
- ➤ How residuals are distributed and related across observations (groups, persons, time, etc.) → these relationships are called "dependency" and this is the primary way that longitudinal models differ from general linear models (e.g., regression)

An Empty Between-Person Model (i.e., Single-Level)



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Adding Within-Person Information... (i.e., to become a Multilevel Model)

Full Sample Distribution 3 People, 5 Occasions each



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Empty +Within-Person Model for y_{ti}



Start off with mean of y_{ti} as "best guess" for any value:

- = Grand Mean
- = Fixed Intercept

Can make better guess by taking advantage of repeated observations:

- = Person Mean
- → Random Intercept

Empty +Within-Person Model



y_{ti} variance \rightarrow 2 sources:

Between-Person (BP) Variance:

- → Differences from **GRAND** mean
- → **INTER**-Individual Differences

Within-Person (WP) Variance:

- → Differences from **OWN** mean
- → **INTRA**-Individual Differences
- → This part is only observable through longitudinal data.

Now we have 2 piles of variance in y_{ti} to predict.

Hypothetical Longitudinal Data (black line = sample mean)



"Error" in a BP Model for the Variance: Single-Level Model



"Error" in a +WP Model for the Variance: Multilevel Model



Empty +Within-Person Model

 \rightarrow



y_{ti} variance \rightarrow 2 sources:

<u>Level 2 Random Intercept</u> // <u>Variance</u> (of U_{0i}, as $\tau_{U_0}^2$):

- Between-Person Variance
- Differences from **GRAND** mean
- → **INTER**-Individual Differences

Level 1 Residual Variance (of e_{ti} , as σ_e^2):

- → Within-Person Variance
- → Differences from **OWN** mean
- → **INTRA**-Individual Differences

BP and +WP Conditional Models

- <u>Multiple Regression</u>, **Between-Person** ANOVA: 1 PILE
 - $> y_i = (\beta_0 + \beta_1 X_i + \beta_2 Z_i...) + e_i$
 - > e_i → ONE residual, assumed uncorrelated with equal variance across observations (here, just persons) → "**BP (all) variation**"
- <u>Repeated Measures</u>, Within-Person ANOVA: 2 PILES
 - > $y_{ti} = (\beta_0 + \beta_1 X_i + \beta_2 Z_i...) + U_{0i} + e_{ti}$
 - > U_{0i} → A random intercept for differences in person means, assumed uncorrelated with equal variance across persons → "BP (mean) variation" = $\tau_{U_0}^2$ is now "leftover" after predictors
 - ► e_{ti} → A residual that represents remaining time-to-time variation, usually assumed uncorrelated with equal variance across observations (now, persons and time) → "WP variation" = σ_e² is also now "leftover" after predictors

Repeated Measures (RM) ANOVA

- Models with a random intercept to predict a constant correlation of outcomes from the same person are also called:
 - > Hierarchical linear models, multilevel models, general linear mixed models, growth curve models, random coefficient models...
- The key to how these latter models can extend beyond traditional RM ANOVA lies in maximum likelihood estimation (ML in MLM) instead of least squares (LS in ANOVA)
 - > Options for other types of outcomes: Normal vs. not-normal
 - > Options for uncooperative participants: Missing or unbalanced data
 - > Options for extension: What if a random intercept is not enough to describe all sources of between-person differences?

Addressing Uncooperative Participants

• ML allows incomplete and unbalanced responses...

<u>RM ANOVA via LS:</u> uses					ses	<u>MLM via ML:</u>	ID	Sex	Time	Y
multivariate (wide) data					uala	uses stacked	100	0	1	5
structure:						(long) data	100	0	2	6
ID	Sex	T1	T2	Т3	T4	structure:	100	0	3	8
100	0	5	6	8	12	Only <u>occasions</u>	100	0	4	12
101	1	Л	7		11	with missing data	101	1	1	4
Dooi		+	are excluded					1	2	7
excluded (data from ID 101					LO1	ID 100 uses 4 cases	101	1	3	•
are not included at all)						ID 101 uses 3 cases	101	1	4	11

Time can also be **unbalanced** across people such that each person can have his or her own measurement schedule: Time "0.9" "1.4" "3.5" "4.2"...

Moving Beyond a Random Intercept

• 2 questions about the possible effects of *time*:

1. Is there an effect of time on average?

- > If the line describing the sample means not flat?
- > Significant FIXED effect of time
- 2. Does the average effect of time vary across individuals?
 - > Does each individual need his or her own line?
 - > Significant RANDOM effect of time

Fixed and Random Effects of Time

(Note: The intercept is random in every figure)



A "Random Linear Time" Model



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MLM "Handles" Dependency

• Where does each kind of person dependency go? Into a new random effects variance component (or "pile" of variance):



Piles of Variance (as Random Effects)

- By adding a random slope, we **carve up** our total variance into 3 piles:
 - > BP (error) variance around intercept
 - > BP (error) variance around slope
 - WP (error) residual variance

These 2 piles are just 1 pile of "error variance" in Univariate RM ANOVA

• But making piles does NOT make error variance go away...



Options for Longitudinal Models

• Although models and software are separate, longitudinal data can be analyzed via multiple analytic frameworks:

» "Multilevel/Mixed Models"

- Dependency over time, persons, groups, etc. are modeled via random effects (multivariate → univariate through "levels" of stacked/long data)
- Builds on GLM, generalizes more easily to additional levels of analysis and crossed sampling (e.g., if people change groups over time)

» "Structural Equation Models"

- Dependency over time *only* is modeled via latent variables (single-level analysis using multivariate/wide data)
- Generalizes easier to broader analysis of latent constructs, mediation, and multivariate multilevel models in general (aka, "Multilevel SEM")
- Because random effects and latent variables are the same thing, many longitudinal models can be specified/estimated either way

- BP model: e_{ti}-only model for the variance
 - $> y_{ti} = \gamma_{00} + e_{ti}$



<u>Mean of the intercept factor</u> = fixed intercept γ₀₀

<u>Loadings of intercept factor</u> = 1 (all occasions contribute equally)

<u>Item intercepts = 0</u> (always)

Variance of intercept factor = 0 so far

<u>Residual variance (e)</u> is assumed to be equal across occasions

> After controlling for the *fixed* intercept, residuals are assumed uncorrelated

- +WP model: U_{0i} + e_{ti} model for the variance
 - $> y_{ti} = \gamma_{00} + U_{0i} + e_{ti}$



<u>Mean of the intercept factor</u> = fixed intercept γ₀₀

Loadings of intercept factor = 1 (all occasions contribute equally)

Variance of intercept factor = random intercept variance

<u>Residual variance (e)</u> is assumed to be equal across occasions

After controlling for the *random* intercept, residuals are assumed uncorrelated

- Fixed linear time, random intercept model:
 - $> y_{ti} = \gamma_{00} + (\gamma_{10} \text{Time}_{ti}) + U_{0i} + e_{ti}$



> After controlling for the *fixed linear slope and random intercept*, residuals are assumed uncorrelated

- Random linear time model:
 - > $y_{ti} = \gamma_{00} + (\gamma_{10} \text{Time}_{ti}) + U_{0i} + (U_{1i} \text{Time}_{ti}) + e_{ti}$



> After controlling for the random linear slope and random intercept, residuals are assumed uncorrelated

Intermediate Summary

- Longitudinal models use random effects/latent variables to quantify and predict sources of variability:
 - > Between persons (BP) in intercept and (aspects of) change over time
 - Why do people start at different places?
 - Why do people change at different rates?
 - > Within persons (WP) after controlling for individual change
 - Why are you off your line today?
- Individuals are conceptualized as continuously varying from one another in each of the between-person dimensions
 - > If so, then one set of variances describes the entire sample
 - > What if that's not the case?
 - > Enter the "heterogeneous population model" variants...

Models for Finding "Hidden" Groups

- Related to traditional cluster analysis (using least squares)
 - > ML variants were popularized by Lazarsfeld and Henry (1968)
- Instead of continuous individual variation, models postulate existence of qualitatively different latent (hidden) subgroups
- More generally known as "finite mixture models," specific model names depend on type of outcomes to be classified:
 - > Categorical, cross-sectional outcomes? "Latent class analysis"
 - > Continuous, cross-sectional outcomes? "Latent profile analysis"
 - > Change in group status over time? "Latent transition analysis"
 - > Change in longitudinal outcomes over time? "Growth mixture models"
 - > All have similar limitations, but we'll focus on **growth mixture models**

Growth Mixture Models (GMM)



 Latent Class = categorical unobserved variable that predicts probabilistic membership in *c* classes

Left: typical depiction of a GMM

GMMs are advertised being able to detect differing latent trajectories across people, but as used in practice, they have significant limitations:

- 1. Completely exploratory
- 2. Sensitive to non-normality
- 3. Distort individual variability
- 4. Classes can only predict existing random effects
- 5. Classes are not needed to examine prediction

1. GMMs are exploratory

How many classes? *????*

- Programs provide relative goodness of fit info, but simulation results suggest these are problematic in practice
 - Information criteria (AIC, BIC) are inadequate for determining # classes
 - Entropy based on classification is only valid if the model fits...

• What are the classes? How should they differ? *???*

- > Nature of the classes is determined entirely by the program
- Get probability of membership to each class for each person, but this will likely change after predicting class membership
 - Should NEVER use the most likely class as an observed variable!

2. GMMs predict non-normality



- A lesser-known but statistically indistinguishable purpose of GMMs is to approximate a non-normal overall distribution
 - So if you fit a GMM erroneously assuming conditional normality, you WILL find two or more latent classes for that reason alone

3. GMMs distort individual variability

- What about individual differences within classes? Well, that depends on the program, too:
 - > SAS PROC TRAJ: What variability?
 - Mplus: Variability is equal across classes, which is likely to be logically impossible... (but freeing this constraint leads to estimation problems)



4. GMMs can only predict model-specified random effects

- Latent classes serve to categorize existing intercepts and slopes...
 - For example, given the specification of a random linear time slope model, latent groups may only differ in level and kinds of linear change...





... just as people *already* do in the random linear growth model!

Latent classes just get in the way.

5. GMMs are not needed to examine prediction



 After fitting a GMM, it is often of interest to then predict class membership from covariates...

5. GMMs are not needed to examine prediction



• ...but the covariates should directly predict the random intercept and slopes themselves instead!

So what should we do instead?

- Before fitting a typical GMMs, specify the most appropriate conditional outcome distribution
 - > Account for floor/ceiling effects of observed measures
- Determine if groups are really necessary to answer your questions... for prediction of differences, probably not!
- If differences due to known predictors are of interest, consider **location-scale longitudinal models** instead (i.e., heterogeneous variance models; see Don Hedeker's work)
 - Allows for prediction of mean differences in intercepts and slopes, as well as prediction of differences in their **amount of variability**
 - > Also very useful to intra-individual variability (IIV) designs

So what should we do instead?

- Specify a confirmatory, hypothesis-driven model that defines the different group trajectories a priori
- Most useful given **qualitatively different** kinds of change
 - > Fit different model of change within each group
 - > Constrain parameters as needed to ensure order/interpretability



Conclusions

- Longitudinal models with random effects/latent variables expand on traditional RM ANOVA via ML:
 - Multiple sources of between-person differences (from random intercepts only to random slopes for change)
 - > Individuals vary continuously from another in growth terms
- Typical uses of growth mixture models try to describe these continuous differences via latent groups instead
 - But are completely exploratory, sensitive to violations of distributional assumptions, inflexible with respect to forms of change, and get in the way of predicting individual differences
 - Confirmatory models may remedy these problems but are seldom used in longitudinal applications