



# New Horizons in Statistical Modeling of Intraindividual Variability with Intensive Longitudinal Data

## **SBM 2023 Pre-Conference Workshop**

Genevieve F. Dunton, PhD, University of Southern California

Donald Hedeker, PhD, University of Chicago

Wei-Lin Wang, PhD, University of Southern California

# Speakers

- Genevieve F. Dunton, PhD, University of Southern California
- Donald Hedeker, PhD, University of Chicago
- Wei-Lin Wang, PhD, University of Southern California

# Workshop Agenda

- 11:00-11:15am Introduction–Overview, Agenda, How to download (Dunton)
- 11:15-11:30am Conceptual Overview and Research Applications (Dunton)
- 11:30-12:00pm Statistical Modeling of Within-Subject Variances (Hedeker)
- 12:00-12:10pm Short Break
- 12:10-12:50pm MixWILD Demonstration and usage (Wang)
- 12:50-1:00pm Closing/Q&A (Dunton, Hedeker, Wang)



# How to Download MixWILD



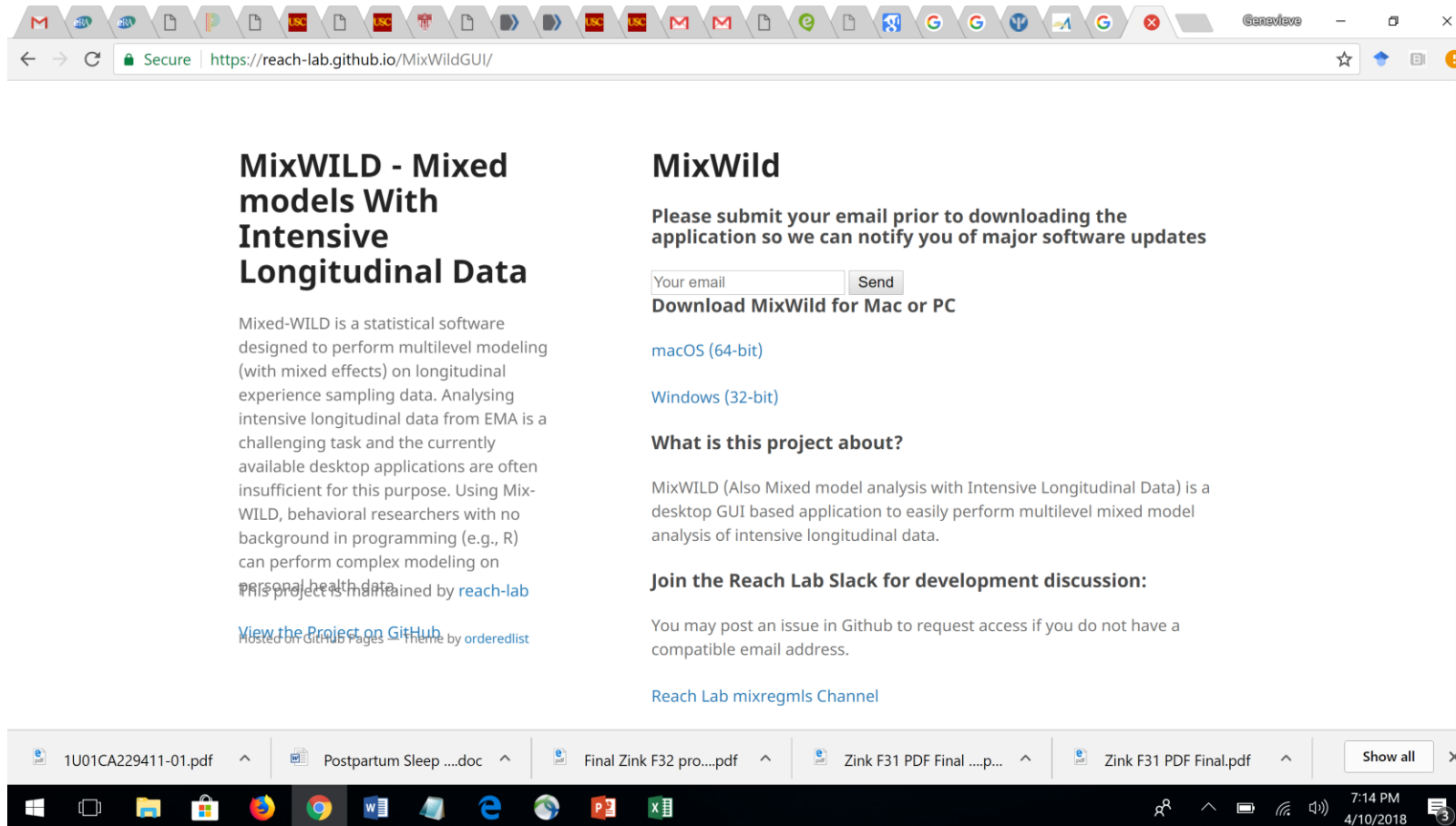
MixWILD for Windows

MixWILD for Mac

- Please visit: <https://reach-lab.github.io/MixWildGUI/>
- Please submit your email prior to downloading the application in the web page so we can notify you of major software updates.
- Click on macOS or Windows to download the program.
- Select your directory to save the program.

When finished downloading, double-click on the MixWILD icon and follow the instructions to complete installation.

# MixWILD GitHub Page



- Software download
- User guide
- Cheat sheets
- Example datasets
- Video tutorial
- Published papers
- User discussion board

# MixWILD: Conceptual Overview and Research Applications



Genevieve F. Dunton, PhD, MPH  
University of Southern California

# Why use MixWILD?

Repeated Occurrence Health Behaviors (e.g., phy. act., healthy eating)

- High frequency (e.g., daily or multiple times per day)
- Time-varying explanatory factors (e.g., context, self-control)



Limited Occurrence Health Behaviors (e.g., vaccinations, diagnostic tests)

- Low frequency (e.g., annually)
- Time-invariant explanatory factors (e.g., access to health care)



# Methodological Weaknesses in Health Behavior Research

- Behaviors measured infrequently using retrospective or summary measures
- Measures capture usual level of behavior or determinants on a typical week or month
- Not conducive to testing factors that vary frequently over micro-timescales (e.g., min, hours)



# Intensive Longitudinal Data (ILD)



- High-frequency and high-density repeated measures data
- Collected over a micro-timescale (e.g., seconds, minutes, hours, days)
- Real-world settings



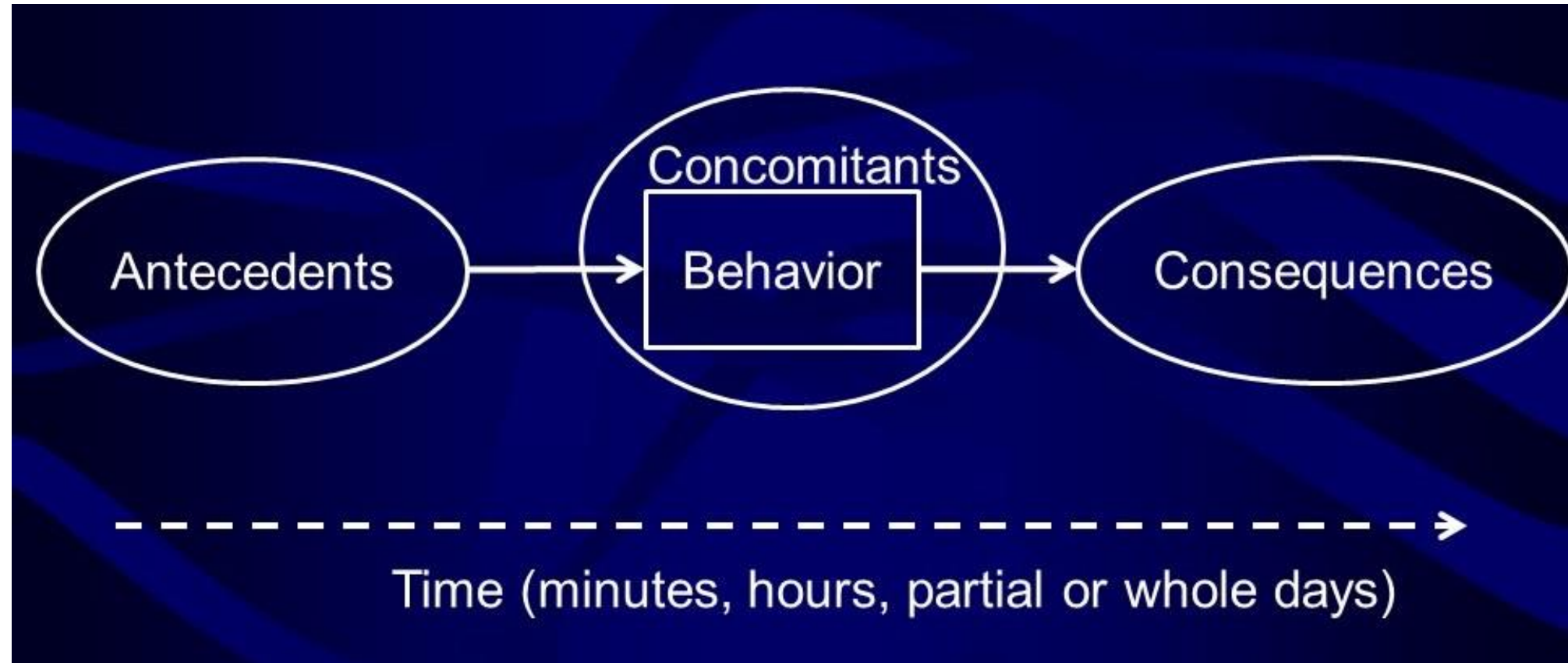
Types of ILD from mobile and sensor devices: self-report (EMA), body movement, biological responses, geographic location, phone/app use, social interactions, and communication patterns.

# Ecological Momentary Assessment (EMA)

- **Ecological**
  - ▣ Real-world environments & experiences
  - ▣ Provides ecological validity
- **Momentary**
  - ▣ Real-time assessment
  - ▣ Avoids recall bias
- **Assessment**
  - ▣ Self-report (subjective)
  - ▣ Multiple repeated measures

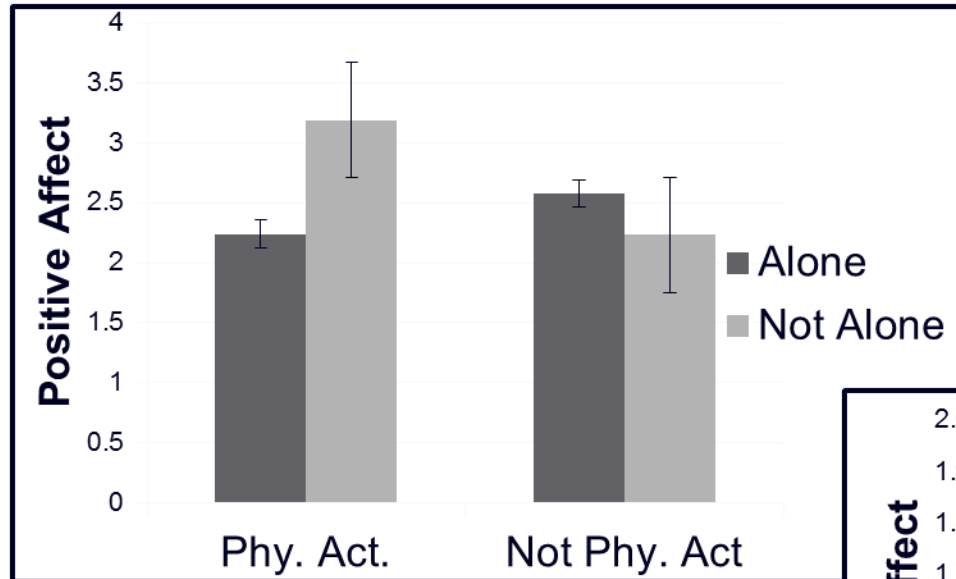


# Common Use of ILD: Examining Momentary Within-Subject Effects of Time-Varying Variables on Health Behaviors

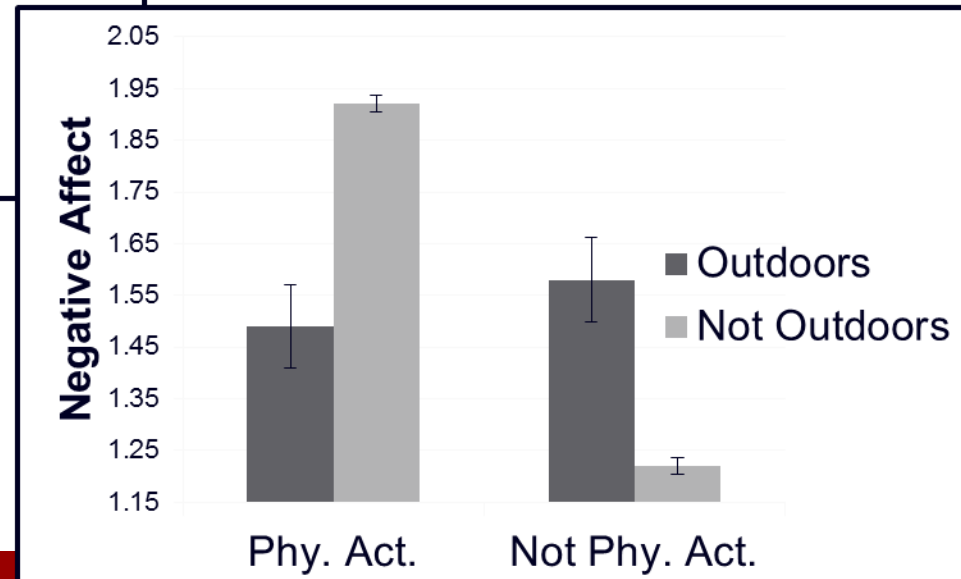


# Common Use of ILD: Examining Momentary Within-Subject Effects of Time-Varying Variables on Health Behaviors

## Contextual Influences on Affective Response During Physical Activity



Measured through 12 days  
of EMA

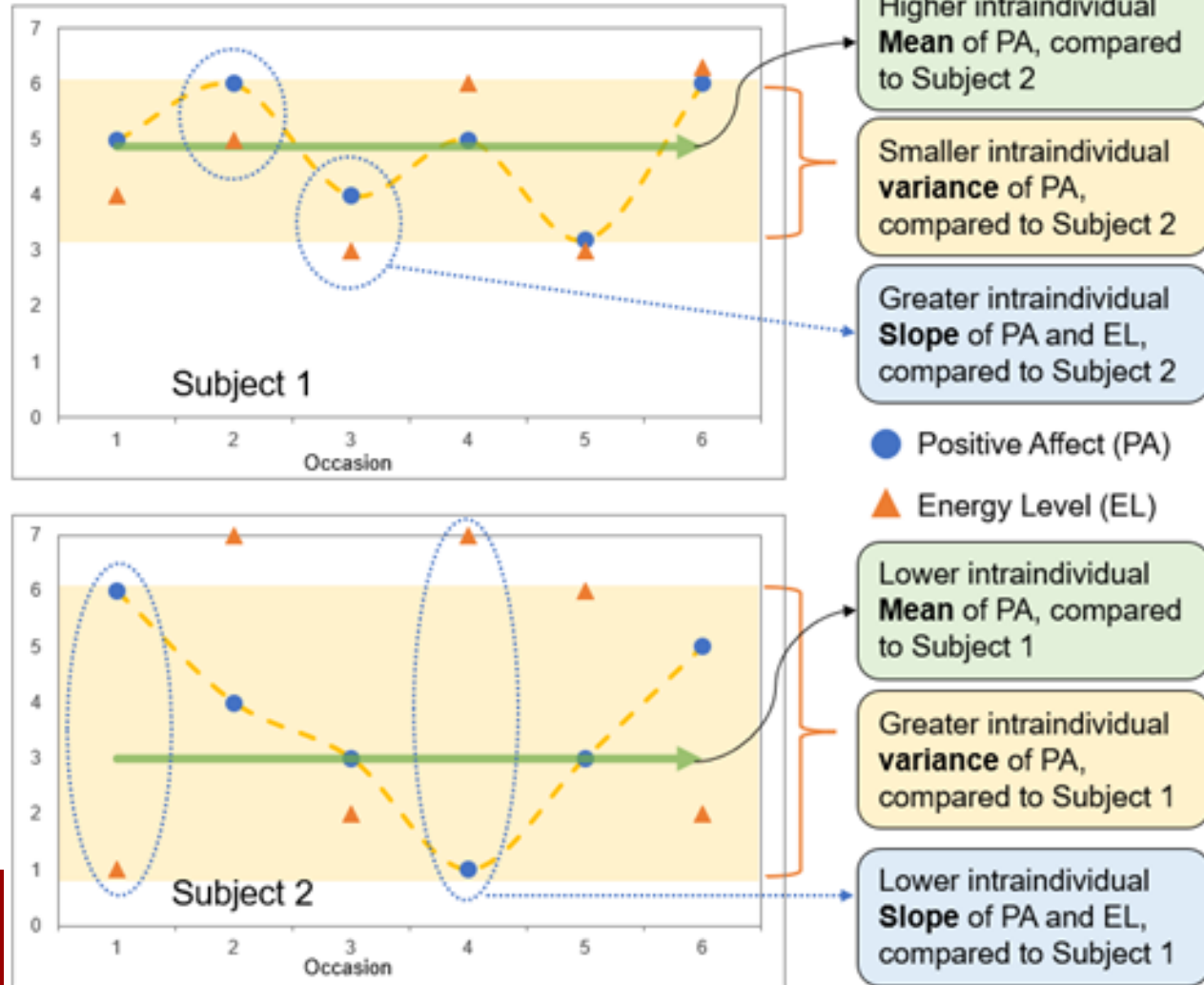


# Less Common Use of ILD: Examining Aggregated Intraindividual Effects of Within-Subject Effects of Time-Varying Variables on Health Behaviors

1. Intraindividual means (i.e., random location effect)  
*How happy is a subject, on average, across occasions?*
2. Intraindividual variances (i.e., random scale effect)  
*How erratic is a subject's mood across occasions?*
3. Intraindividual slopes (i.e., random slope effect)  
*How strongly is a subject's mood related to activity across occasions?*

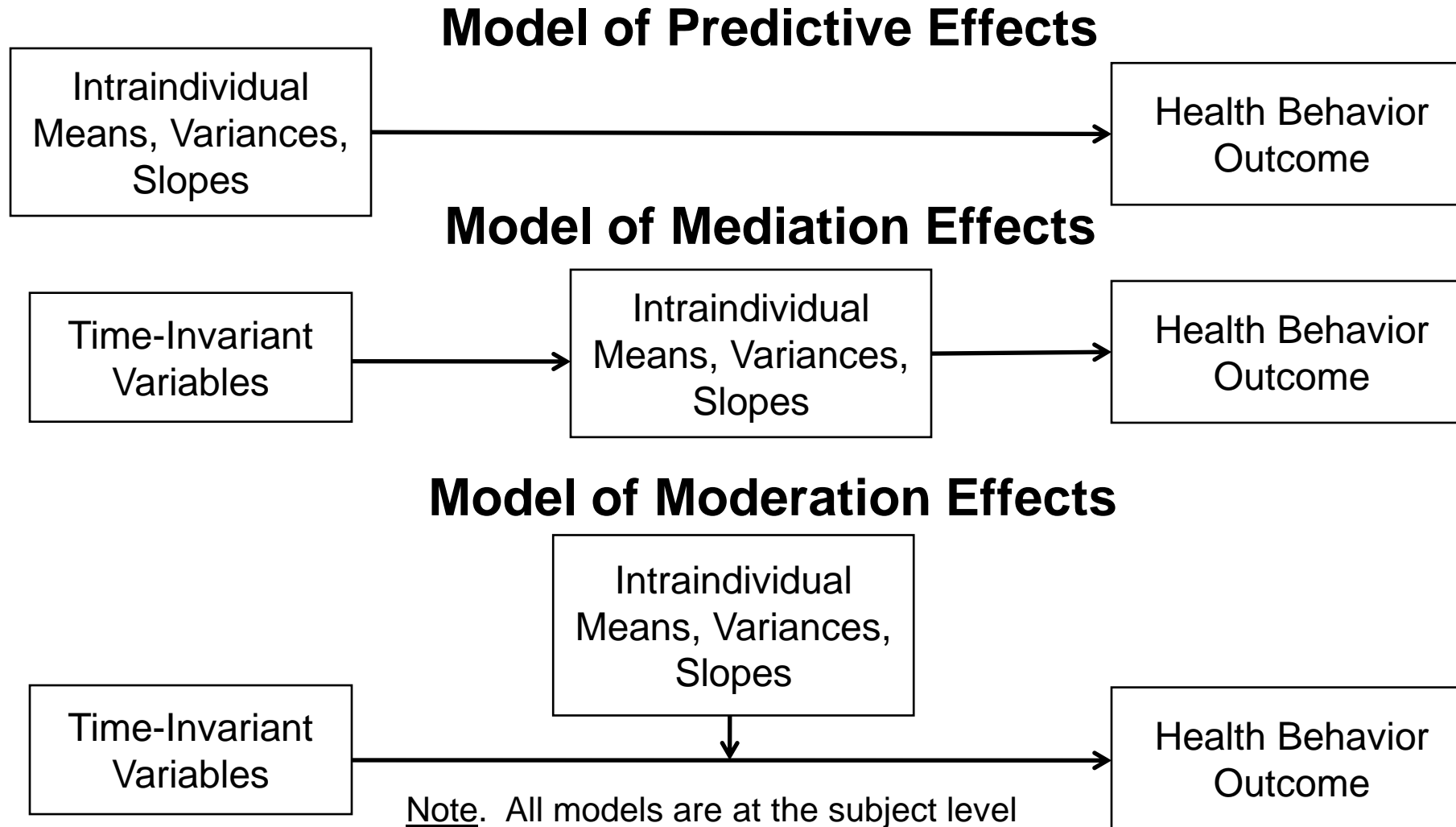
# ILD Can Examine Aggregated Intraindividual Effects of Time-Varying Variables on Health Behaviors

Figure 1-2: Conceptual Graphs of Intraindividual Means, Variances, and Slopes

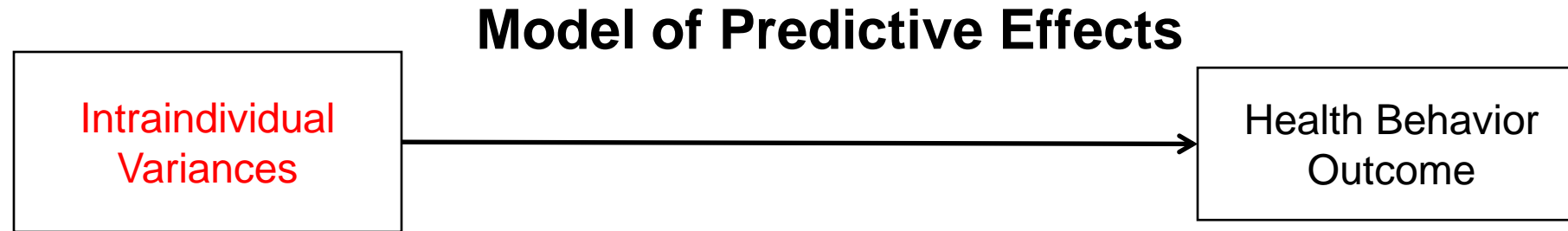




# ILD Can Examine Aggregated Intraindividual Effects of Time-Varying Variables on Health Behaviors



# ILD Can Examine Aggregated Intraindividual Effects of Time-Varying Variables on Health Behaviors



Note. All models are at the subject level



# Intraindividual Variance Predicting a Subject-Level Health Behavior Outcome

Individuals with greater variability in feelings of energy have lower odds of meeting physical activity guidelines

**Table 4**

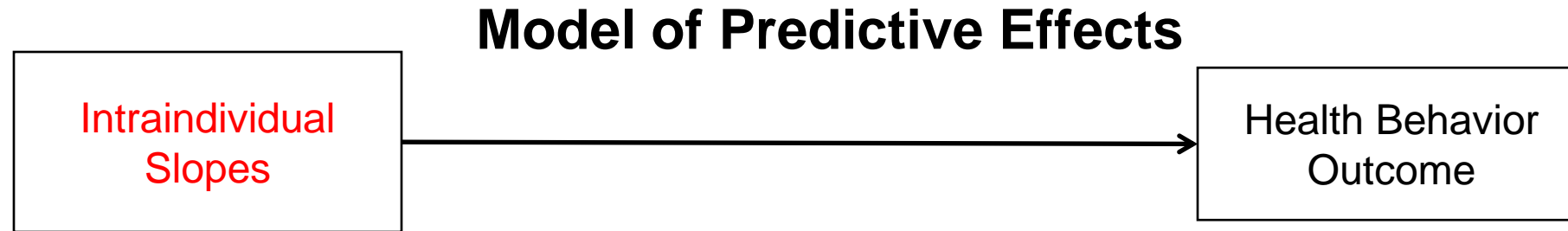
Subject-Level Regression Models predicting Odds of Meeting Physical Activity Guidelines (Logistic) and Sedentary Time (Linear).

	Predicting Odds of Meeting Physical Activity Guidelines Estimate (Standard Error)	Predicting Minutes of Sedentary Time per Valid Hour of Wear Estimate (Standard Error)
<b>Positive Affect</b>		
Intercept	0.58 (0.32)	33.74** (0.68)
Mean level of Positive Affect	0.09 (0.10)	0.33 (0.31)
Variability in Positive Affect	0.07 (0.09)	0.16 (0.23)
Sex (Female)	-0.92** (0.21)	1.04* (0.41)
Age	0.01 (0.01)	0.11** (0.01)
<b>Feelings of Energy</b>		
Intercept	0.51 (0.55)	34.93** (1.03)
Mean level of Feelings of Energy	-0.09 (0.18)	-0.26 (0.30)
Variability in Feelings of Energy	-0.43* (0.21)	0.15 (0.37)
Sex (Female)	-1.13** (0.36)	1.40* (0.58)
Age	-0.01 (0.01)	0.07** (0.02)

Note. Model using positive affect data are based on 617 participants. Model using feelings of energy data are based on 245 participants. For logistic regression models predicting odds of meeting physical activity guidelines, logit (i.e., log odds) estimates are displayed.

\* $p < 0.05$ . \*\* $p < 0.01$ .

# ILD Can Examine Aggregated Intraindividual Effects of Time-Varying Variables on Health Behaviors

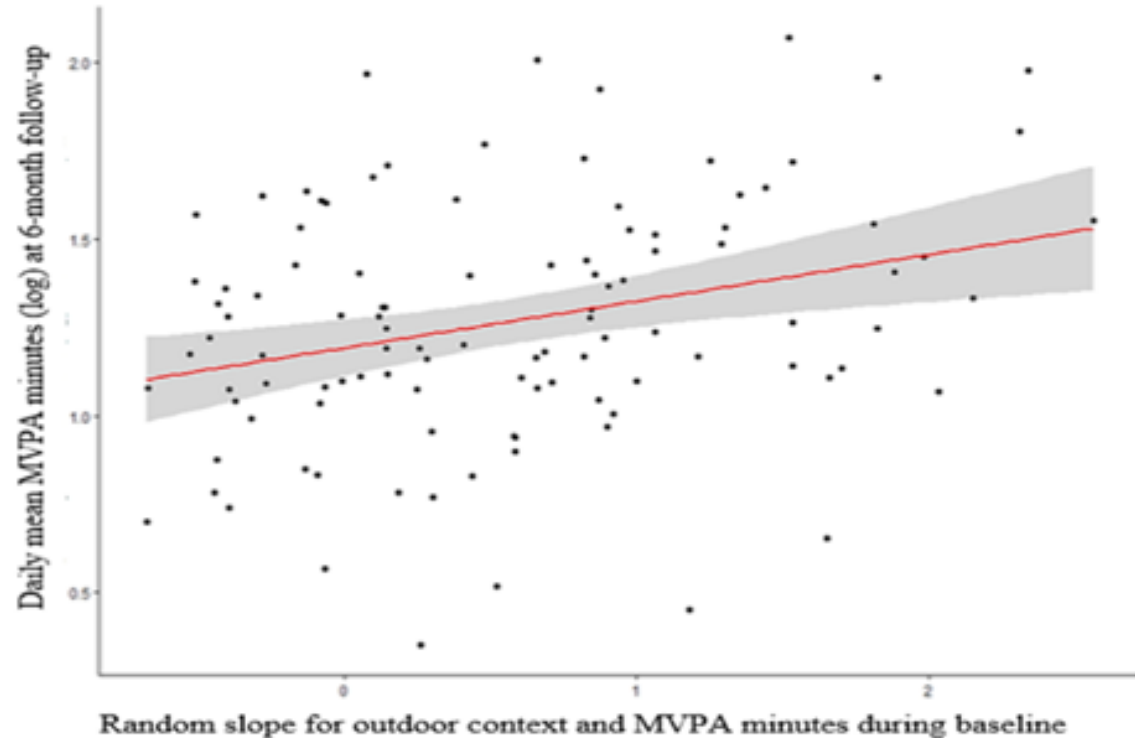


Note. All models are at the subject level

# Intraindividual Slope Predicting a Subject-Level Outcome

Intraindividual slope =  
momentary association  
between context and  
MVPA

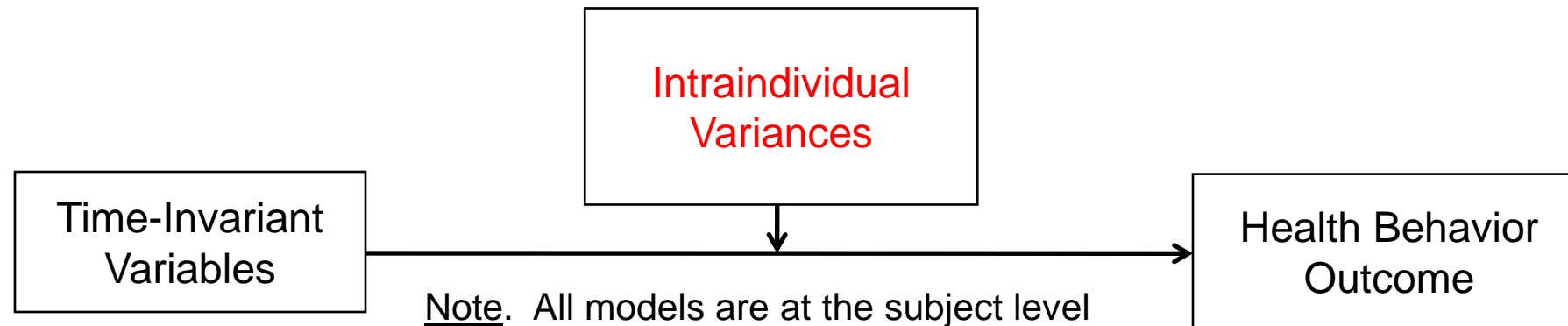
Figure 6: Intraindividual Slope (i.e., Momentary Association of Outdoor Context and MVPA) Predicting Adults' MVPA Six Months Later



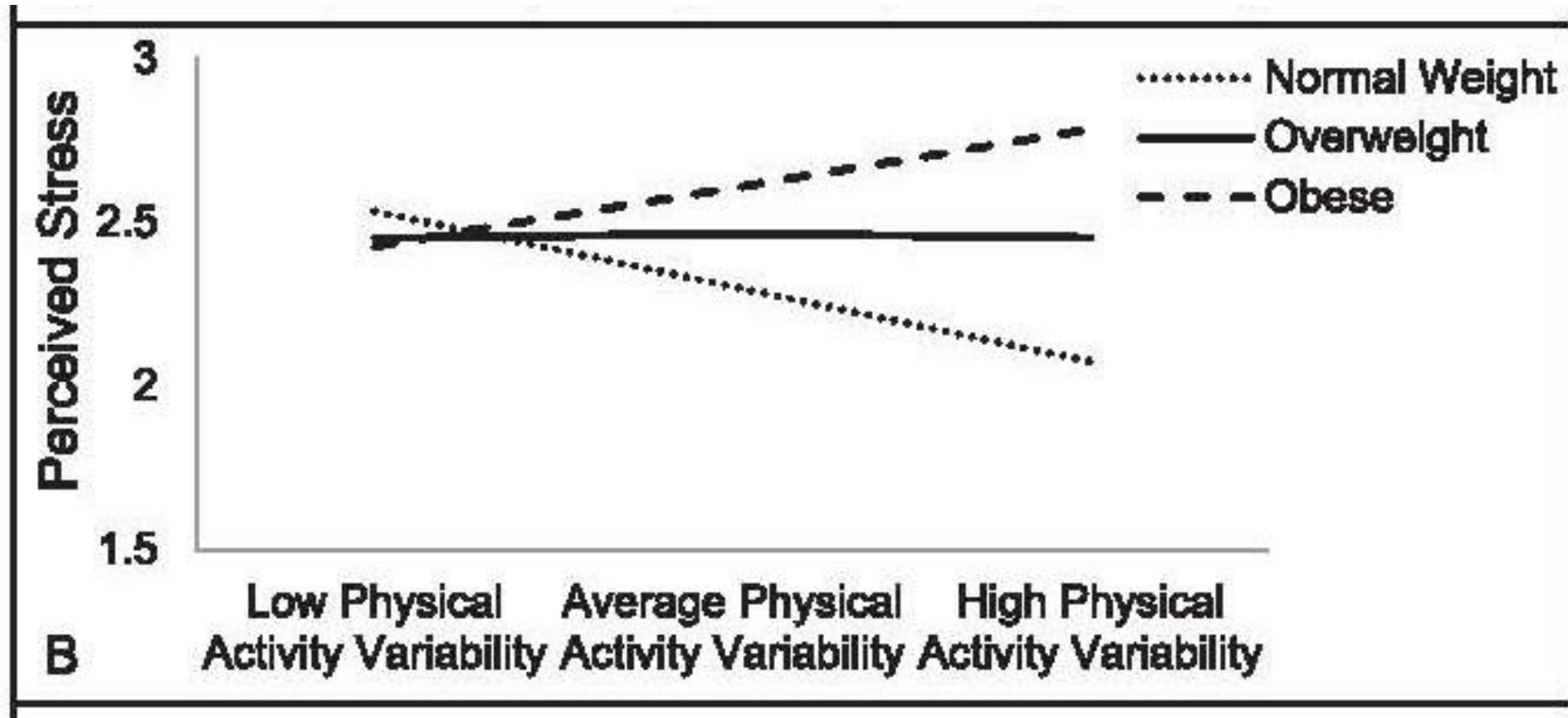
Participants, who had more momentary MVPA when outdoors (vs. indoors) during baseline (i.e., higher intraindividual slope), had higher daily MVPA six months later

# ILD Can Examine Aggregated Intraindividual Effects of Time-Varying Variables on Health Behaviors

## Model of Moderation Effects



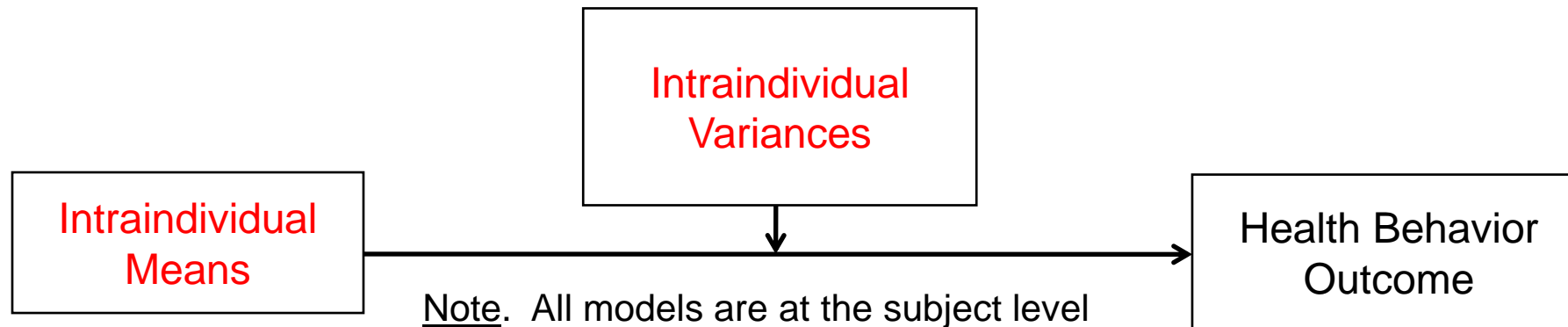
# Intraindividual Variance Moderating the Effect of a Subject-Level Factor on a Subject-Level Outcome



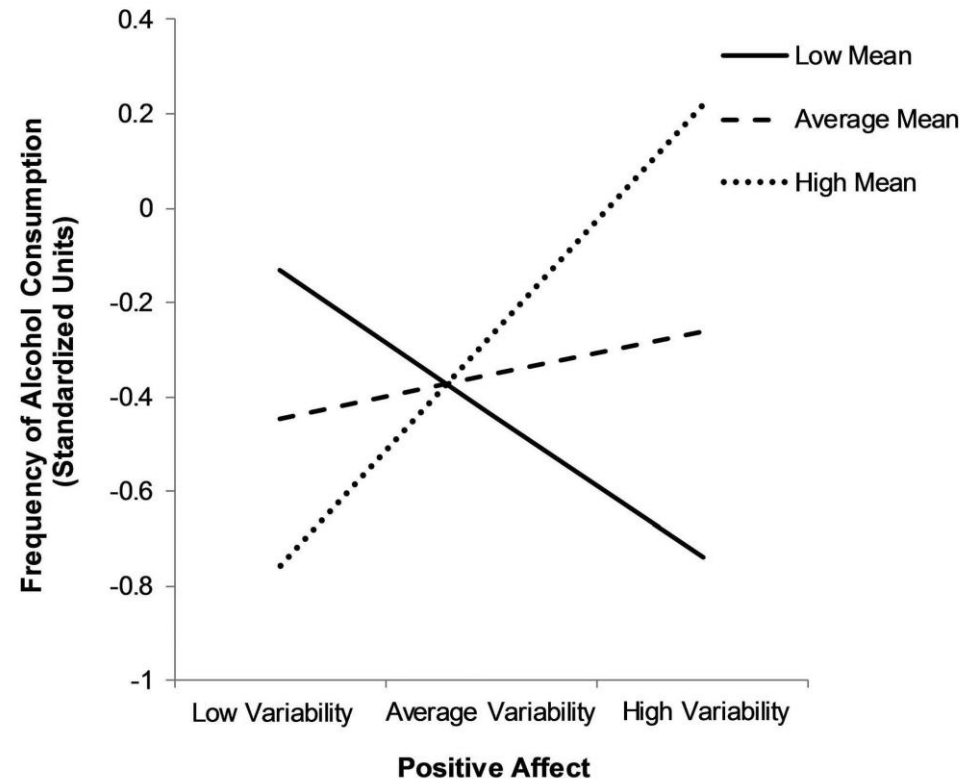
Individuals with obesity have higher levels of perceived stress, but only for those with high day-to-day variability in physical activity

# ILD Can Examine Aggregated Intraindividual Effects of Time-Varying Variables on Health Behaviors

## Model of Moderation Effects



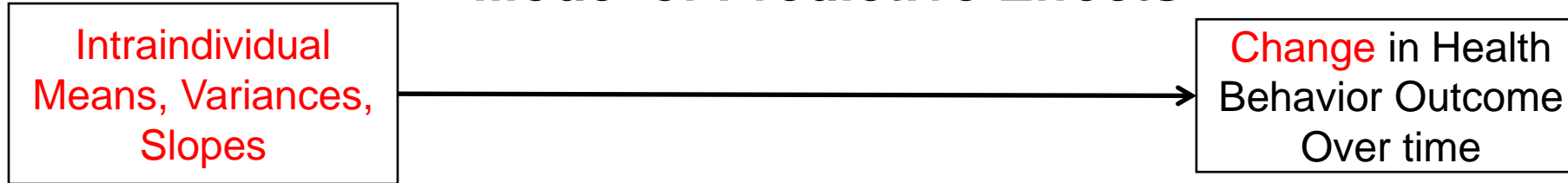
# Intraindividual Variance Moderating the Effect of a Intraindividual Mean on a Subject-Level Outcome



Individuals with low mean positive affect have higher alcohol consumption, but only for those with low variability in positive affect

# ILD Can Examine Aggregated Intraindividual Effects of Time-Varying Variables on Health Behaviors

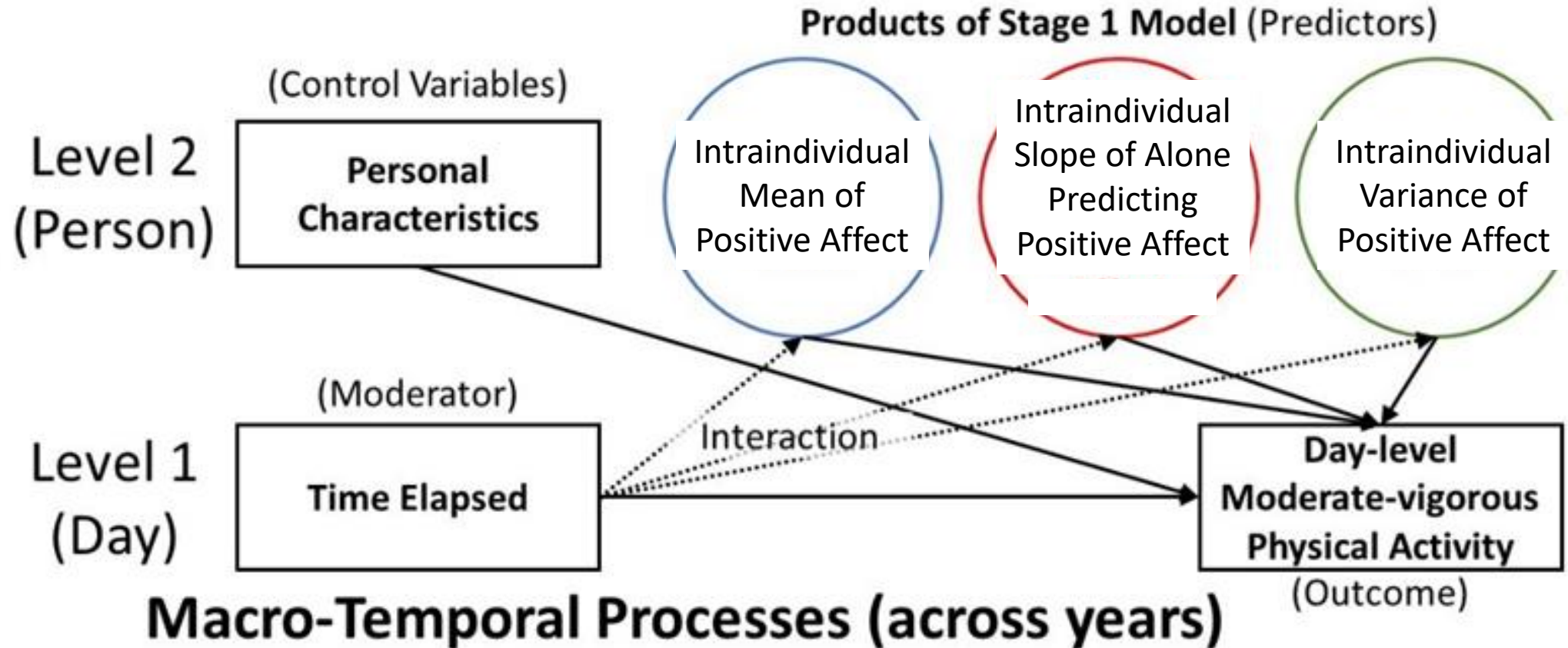
## Model of Predictive Effects



Note. Outcome is no longer at the subject level

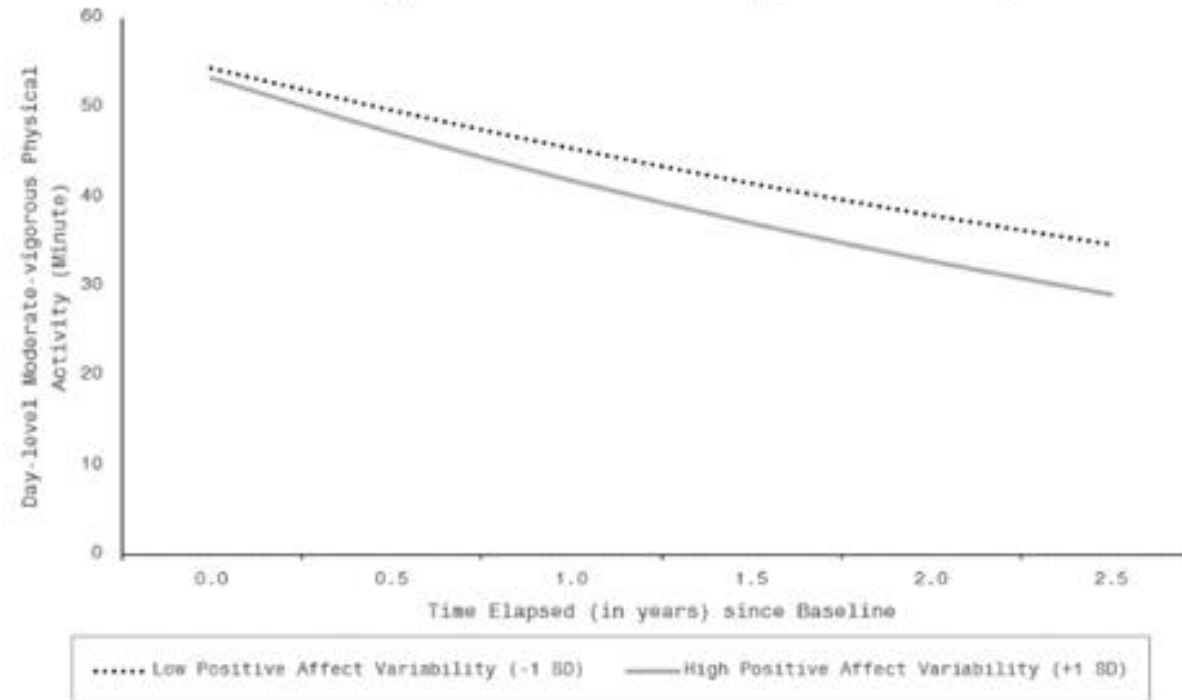


# Intraindividual Means, Variances, and Slopes Predicting Change in an Outcome over Time



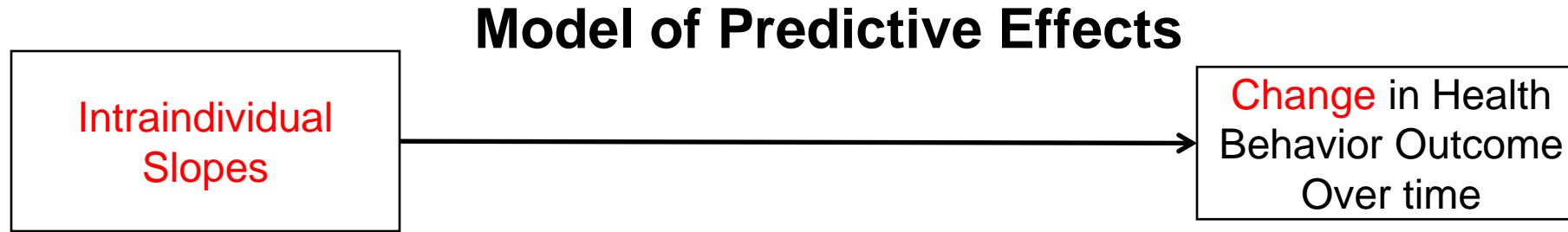
# Intraindividual Variance Predicting Change in an Outcome over Time

Figure 4: Time Elapsed (in years) Since Baseline  $\times$  Intraindividual Scale (i.e., Positive Affect Variability) Interaction Predicting Children's Day-level MVPA



Children who had greater intraindividual variance in positive affect had a faster rates of decline in physical across three years

# ILD Can Examine Aggregated Intraindividual Effects of Time-Varying Variables on Health Behaviors

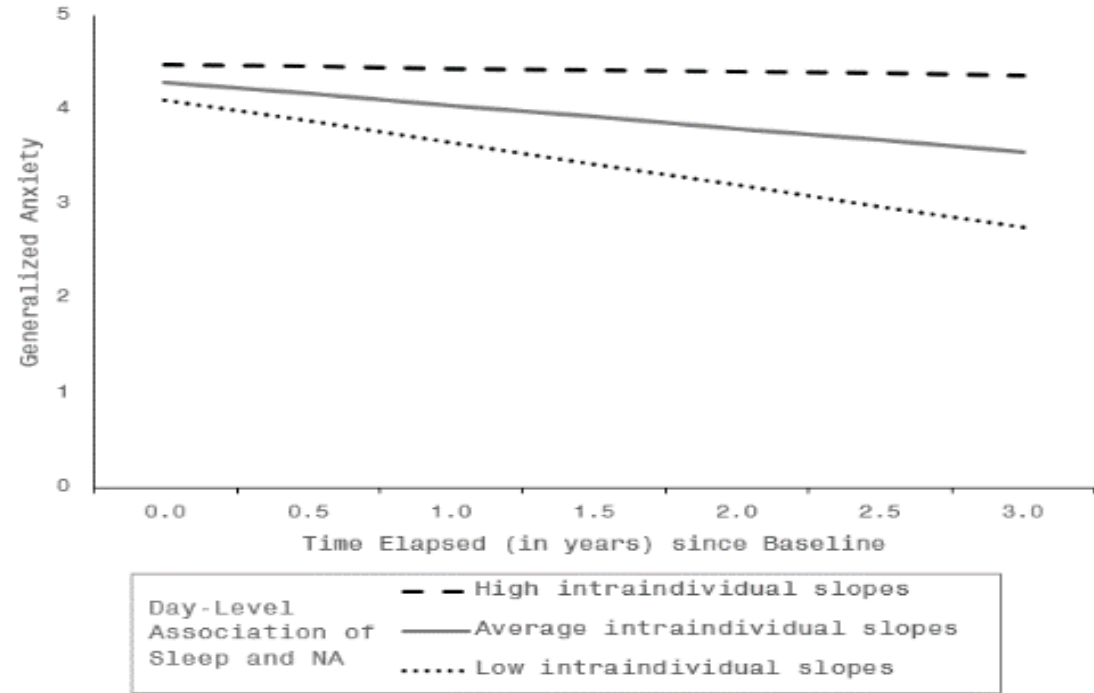


Note. Outcome is no longer at the subject level

# Intraindividual Slope Predicting Change in an Outcome over Time

Intraindividual slope =  
day-level association  
between negative affect  
and sleep

Figure 5: Time Elapsed (in years) Since Baseline x Intraindividual Slope (i.e., Day-Level Association of Sleep and NA) Interaction Predicting Children's Generalized Anxiety



Children who needed to sleep more on nights following days with higher negative affect had a slower rate of decline in generalized anxiety across three years

# Overview of MixWILD

- First Stage Model- estimates intraindividual means, variances, and slopes as random effects in mixed-effects location scale multilevel model
- Second Stage Model- uses random means, variances, and slopes from first stage as predictors of a subject-level or time-varying outcome in a single or multilevel linear or logistic regression model

**MIX{WILD}**  
Mixed Model Analysis With Intensive Longitudinal Data

# Other 2023 SBM Presentations Using MixWILD

**Do et al.** Investigating day-level associations between affective variability and physical activity using Ecological Momentary Assessment.  
**Symposium on Friday April 28, 2023 at 9:00 AM**

**Yang et al.** The mean level, between-person differences, and within-person variability of older adults' daily sleep quality and duration.  
**Paper Session 34 on Friday April 28, 2023 at 1:00 PM**

**Wang et al.** Associations of smartphone usage with average day-level and day-to-day variability of mood in emerging adults  
**Poster Session E on Sat. April 29, 2023 at 11:00 AM**

# Acknowledgments

## Collaborators

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# **A Brief Introduction to the Mixed-Effects Location Scale (MELS) Model**

Don Hedeker

Department of Public Health Sciences

University of Chicago

<https://voices.uchicago.edu/hedeker/>

Supported by National Cancer Institute grant R01 CA240713 (Hedeker & Dunton)

## **Ecological Momentary Assessment (EMA) data**

experience sampling and diary methods, intensive longitudinal data

- Subjects provide frequent reports on events and experiences of their daily lives (*e.g.*, 30-40 responses per subject collected over the course of a week or so)
- electronic diaries: cell phones, palm pilots, personal digital assistants (PDAs), interactive voice response (IVR) systems, actigraphs, web-based
- Capture particulars of experience in a way not possible with more traditional designs  
*e.g.*, allow investigation of phenomena as they happen over time
- Reports could be time-based, following a fixed-schedule, randomly triggered, event-triggered

# Data are rich and offer many modeling possibilities!

- person- and occasion-level effects on occasion-level responses  
⇒ potential influence of context and/or environment  
*e.g.*, subject response might vary when alone vs with others
- data are inherently multilevel
  - occasions (level-1) within subjects (level-2)
  - occasions (level-1) within days (level-2) within subjects (level-3)
  - occasions (level-1) within waves (level-2) within subjects (level-3)
- References for mixed model analysis of EMA data
  - Schwartz, J.E. & Stone, A. (2007). The analysis of real-time momentary data: A practical guide. In: A.A. Stone, S.S. Shiffman, A. Atienza, and L. Nebeling, editors, *The science of real-time data capture: Self-report in health research*. Oxford, England: Oxford University Press, p. 76-113.
  - Walls, T.A., Jung, H., & Schwartz, J.E. (2006). Multilevel models for intensive longitudinal data. In: Walls, T.A. and Schafer, J.L., editors, *Models for intensive longitudinal data*. New York: Oxford University Press, p. 3-37

# Learning Objectives

- Using mixed-effects location scale (MELS) models, examine why subjects differ in mean level *as well as variability*
  - Between-subjects variance
    - e.g.*, subject heterogeneity can vary by gender, age, or context
    - \* modeling of between-subjects variance in terms of covariates
    - \* inclusion of random subject intercepts and slopes
  - Within-subjects variance
    - e.g.*, subject inconsistency can vary by gender, age, or context
    - \* modeling of within-subjects variance in terms of covariates, including random subject scale
- MixWILD freeware program example

Carroll (2003) Variances are not always nuisance parameters, *Biometrics*.

## Mixed-Effects Location Scale Models for EMA data

- Hedeker, Mermelstein, & Demirtas (2008). An application of a mixed-effects location scale model for analysis of Ecological Momentary Assessment (EMA) data. *Biometrics*, 64, 627-634.
- Hedeker, D., Mermelstein, R.J., & Demirtas, H. (2012). Modeling between- and within-subject variance in EMA data using mixed-effects location scale models. *Statistics in Medicine*, 31 3328-3336.

**Multilevel (mixed-effects regression) model** for measurement  $y$  of subject  $i$  ( $i = 1, 2, \dots, N$ ) on occasion  $j$  ( $j = 1, 2, \dots, n_i$ )

$$y_{ij} = \mathbf{x}_{ij}'\boldsymbol{\beta} + v_i + \epsilon_{ij}$$

$\mathbf{x}_{ij} = p \times 1$  vector of regressors (including a column of ones)

$\boldsymbol{\beta} = p \times 1$  vector of regression coefficients

$v_i \sim N(0, \sigma_v^2)$  BS variance; how homogeneous/heterogeneous are subjects?

$\epsilon_{ij} \sim N(0, \sigma_\epsilon^2)$  WS variance; how consistent/erratic are the data within subjects?

Model with no covariates:  $y_{ij} = \beta_0 + v_i + \epsilon_{ij}$

- $v_i$  is subject's mean (deviation from  $\beta_0$ )
  - if subjects are alike,  $v_i \approx 0$  and  $\sigma_v^2$  will approach 0
  - if subjects are different,  $v_i \neq 0$  and  $\sigma_v^2$  will increase from 0

$\Rightarrow$  magnitude of  $\sigma_v^2$  indicates how different subjects are from each other (homogeneity/heterogeneity)
- $\epsilon_{ij}$  is subject  $i$ 's error at time  $j$  (deviations from their mean)
  - if subjects are all well-fit,  $\epsilon_{ij} \approx 0$  and  $\sigma_\epsilon^2$  will approach 0
  - if subjects are not well-fit,  $\epsilon_{ij} \neq 0$  and  $\sigma_\epsilon^2$  will increase from 0

$\Rightarrow$  magnitude of  $\sigma_\epsilon^2$  indicates how data vary within subjects (consistency/erraticism)

## Log-linear models for variances

BS variance  $\sigma_{v_{ij}}^2 = \exp(\mathbf{u}_{ij}'\boldsymbol{\alpha})$  or  $\log(\sigma_{v_{ij}}^2) = \mathbf{u}_{ij}'\boldsymbol{\alpha}$

WS variance  $\sigma_{\epsilon_{ij}}^2 = \exp(\mathbf{w}_{ij}'\boldsymbol{\tau})$  or  $\log(\sigma_{\epsilon_{ij}}^2) = \mathbf{w}_{ij}'\boldsymbol{\tau}$

- $\mathbf{u}_{ij}$  and  $\mathbf{w}_{ij}$  include covariates (and  $\mathbf{1}$ )
- subscripts  $i$  and  $j$  on variances indicate that these change depending on covariates  $\mathbf{u}_{ij}$  and  $\mathbf{w}_{ij}$  (and their coefficients)
- exp function ensures a positive multiplicative factor, and so resulting variances are positive



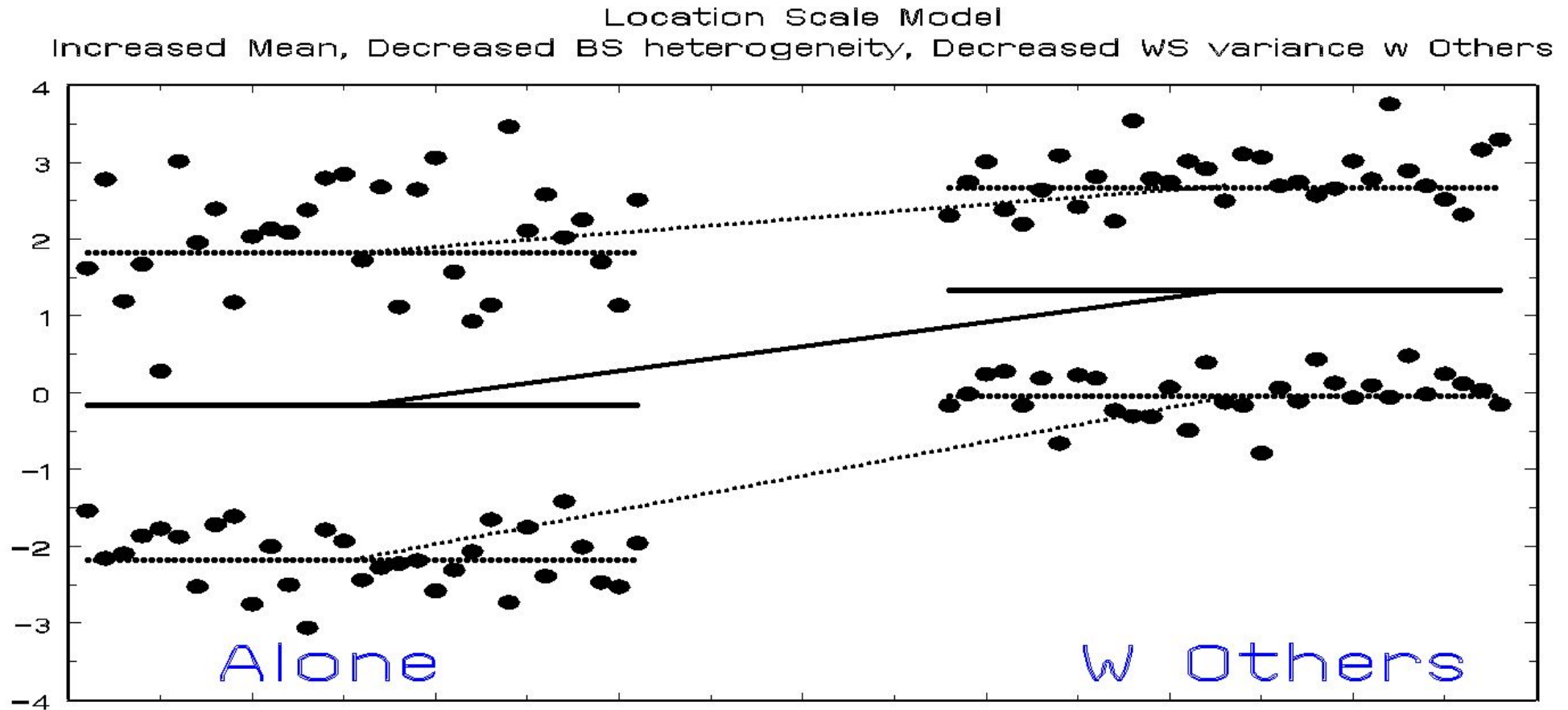
## How can WS variables influence BS variance?

$$\sigma_{v_{ij}}^2 = \exp(\mathbf{u}_{ij}'\boldsymbol{\alpha})$$

- Do rainy days and Mondays get everyone down?
- Is Tuesday just as bad as Stormy Monday for all?
- Are all kids happy on the last day of school?

Example: strong positive effect of being alone on BS variance of positive and negative mood

⇒ being alone increases subject heterogeneity (or, subjects report more similar mood when with others)



- Means are increased with others
- Subjects are more similar to each other when with others (BS var)
- Within-subject data are more consistent with others (WS var)

## WS variance varies across subjects

$$\sigma_{\epsilon_{ij}}^2 = \exp(\mathbf{w}_{ij}'\boldsymbol{\tau} + \omega_i) \quad \text{where} \quad \omega_i \sim N(0, \sigma_\omega^2)$$

$$\log(\sigma_{\epsilon_{ij}}^2) = \mathbf{w}_{ij}'\boldsymbol{\tau} + \omega_i$$

- $\omega_i$  are log-normal subject-specific perturbations of WS variance
- $\omega_i$  are “scale” random effects - how does a subject differ in terms of the variation in their data
- $v_i$  are “location” random effects - how does a subject differ in terms of the mean of their data

## Multilevel model of WS variance

$$\log(\sigma_{\epsilon_{ij}}^2) = \mathbf{w}_{ij}'\boldsymbol{\tau} + \omega_i$$

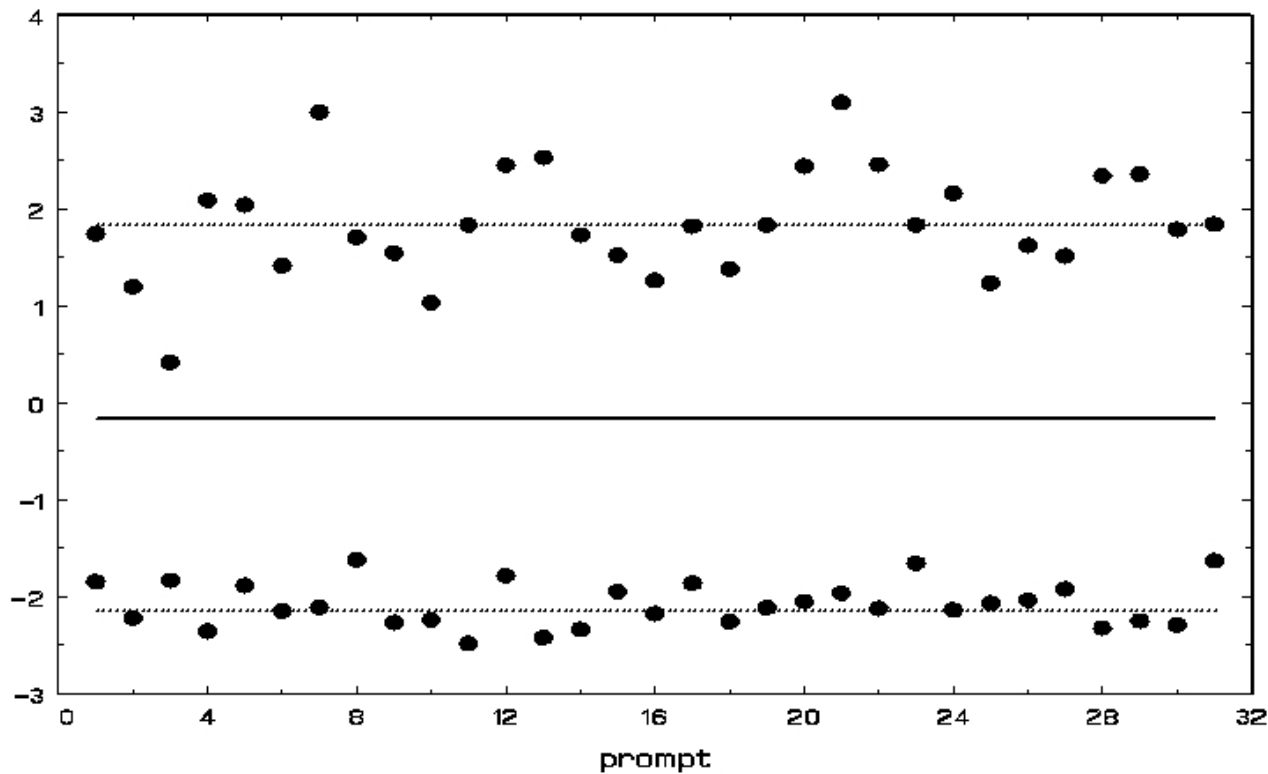
Why not use some summary statistic per subject (say, calculated subject standard deviation  $S_{y_i}$ ) in a second-stage model?

$$S_{y_i} = \mathbf{x}_i'\boldsymbol{\beta} + \epsilon_i$$

latter approach

- treats all standard deviations as if they are equally precise (but some might be based on 2 prompts or 40 prompts)
- does not recognize that these are estimated quantities (underestimation of sources of variation)
- does not allow occasion-varying predictors

⇒ We use multilevel models for mean response, why not for variance?



Model allows covariates to influence

- mean: level of solid line
- BS variance: dispersion of dotted lines
- WS variance: dispersion of points

additional random subject effects on: mean and WS variance

# MixWILD: Mixed-effects models With Intensive Longitudinal Data

Dzubur, E., Ponnada, A., Nordgren, R., Yang, C.-H., Intille, S., Dunton, G., & Hedeker, D. (2020). MixWILD: A program for examining the effects of variance and slope of time-varying variables in intensive longitudinal data. *Behavior Research Methods*, 52:1403–1427.

<https://reach-lab.github.io/MixWildGUI>

## Example: a MELS model using MixWILD

Data are from: <https://dataverse.harvard.edu/dataverse/harvard>

“How health behaviors relate to academic performance via affect: An intensive longitudinal study”  
by Flueckiger L, Lieb R, Meyer AH, Mata J

ID	Subject number
Day	Survey day
Sex	Participants' sex
Age	Participants' age
Sem	Semester: Number of semesters studied
SQ	Sleep quality 1 (very bad) to 4 (very good)
PhysAct	Physical activity: Number of minutes engaged in mild, moderate and strenuous exercise weighted by metabolic equivalents and then summed to produce a total daily leisure activity score
PA	Positive affect 1 (not at all) to 7 (extremely)
NA	Negative affect 1 (not at all) to 7 (extremely)
LGA	Learning goal achievement 0 (not at all) to 4 (completely)
Exam	Examination success 0 (fail) 1 (pass)
HSG	High school grades 1 (lowest grade) to 6 (highest grade)
BDI	Beck Depression Inventory 1(not) 2 (mild to moderate) 3 (clinically relevant symptoms)
Added variable:	
Day_c	centered and scaled version of day ( -2.2143 to 2.2143; 1 unit = 1 week)
<b>-99</b>	<b>Missing value</b>

# Dataset\_HealthBehavAcadPerfAffect.csv

Dataset_HealthBehavAcadPerfAffect - Excel															
File Home Insert Page Layout Formulas Data Review View Tell me what you want to do...															
Clipboard Font Alignment Number Styles Cells															
A1 ID															
	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O
1	ID	Day	Sex	Age	Sem	SQ	PhysAct	PA	NA	LGA	Exam	HSG	BDI	Day_c	
2	1	1	1	22	2	3	30	2.666667	3.666667	2	1	4.6	2	-2.21429	
3	1	2	1	22	2	3	60	3.333333	3	1	1	4.6	2	-2.07143	
4	1	3	1	22	2	3	360	2.666667	4	0	1	4.6	2	-1.92857	
5	1	4	1	22	2	3	780	1.333333	5	1	1	4.6	2	-1.78571	
6	1	6	1	22	2	3	210	3	4.666667	2	1	4.6	2	-1.5	
7	1	8	1	22	2	3	310	4	3.666667	2	1	4.6	2	-1.21429	
8	1	9	1	22	2	3	90	3.333333	3.333333	0	1	4.6	2	-1.07143	
9	1	10	1	22	2	3	405	3.333333	2.666667	2	1	4.6	2	-0.92857	
10	1	11	1	22	2	3	405	2.333333	2.666667	2	1	4.6	2	-0.78571	
11	1	12	1	22	2	3	360	3	4	2	1	4.6	2	-0.64286	
12	1	13	1	22	2	3	270	1.666667	5.666667	1	1	4.6	2	-0.5	
13	1	14	1	22	2	3	30	3.333333	2.333333	2	1	4.6	2	-0.35714	
14	1	15	1	22	2	3	540	2.666667	2	1	1	4.6	2	-0.21429	
15	1	17	1	22	2	2	405	4.333333	3.333333	1	1	4.6	2	0.071429	
16	1	18	1	22	2	3	405	2.666667	3	1	1	4.6	2	0.214286	
17	1	19	1	22	2	3	540	2.666667	3.333333	1	1	4.6	2	0.357143	
18	1	21	1	22	2	3	60	3.666667	2.666667	2	1	4.6	2	0.642857	
19	1	22	1	22	2	3	405	3.333333	3	1	1	4.6	2	0.785714	
20	1	23	1	22	2	3	405	3.333333	3.333333	2	1	4.6	2	0.928571	
21	1	24	1	22	2	2	60	1.666667	4.666667	2	1	4.6	2	1.071429	
22	1	25	1	22	2	3	405	5	2.666667	2	1	4.6	2	1.214286	



## MELS model of PA

outcome: PA (positive affect)

regressor: Day\_c (centered and scaled version of day)

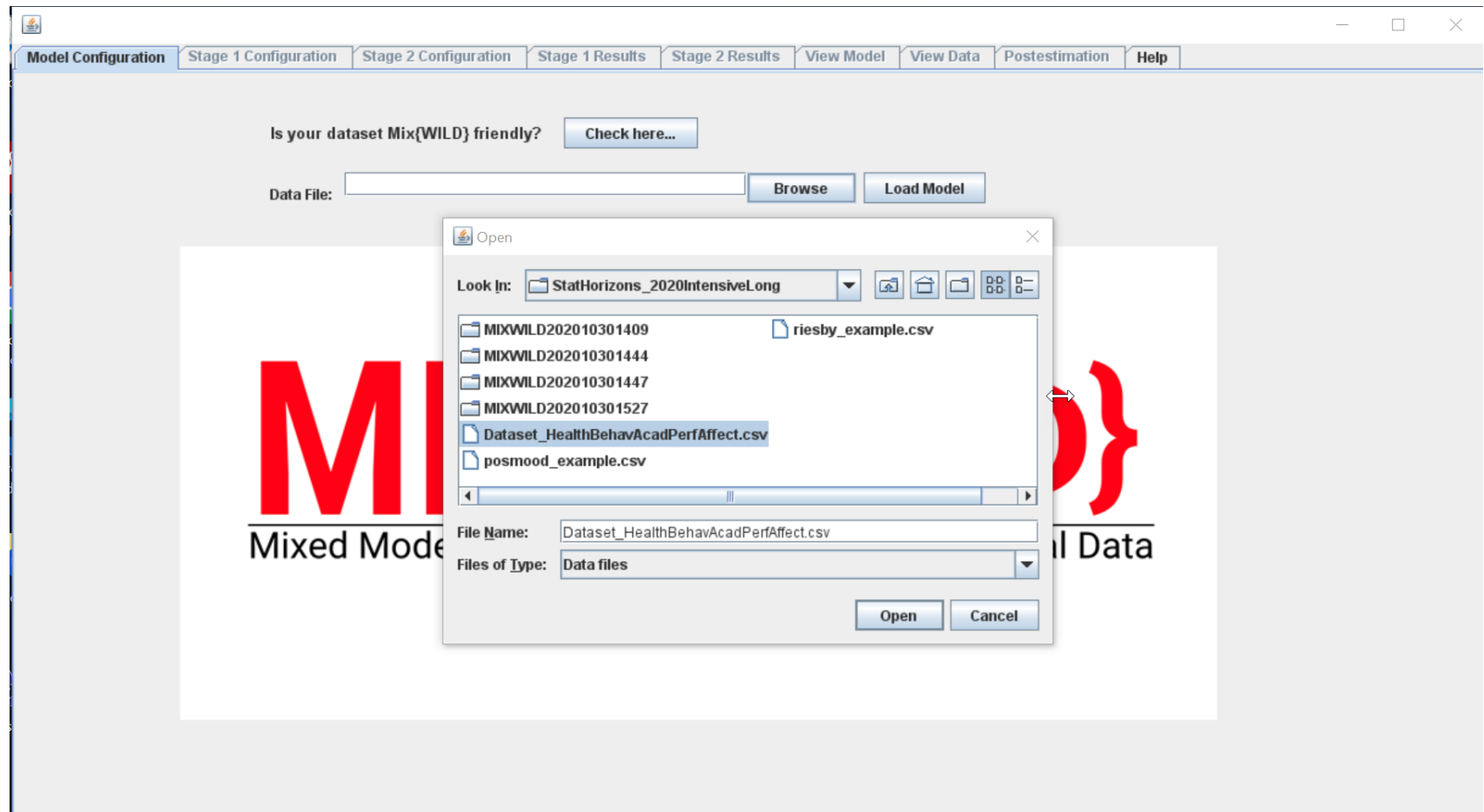
$$PA_{ij} = \beta_0 + \beta_1 Day\_c_{ij} + v_i + \epsilon_{ij}$$

$$\sigma_{v_{ij}}^2 = \exp(\alpha_0 + \alpha_1 Day\_c_{ij})$$

$$\sigma_{\epsilon_{ij}}^2 = \exp(\tau_0 + \tau_1 Day\_c_{ij} + \tau_v v_i + \omega_i)$$

Here,  $v_i$  is the random subject location effect, and  $\omega_i$  is the random subject scale effect, both normally distributed

Browse for Dataset\_HealthBehavAcadPerfAffect.csv



Provide a title and make selections, then click on Submit

The screenshot shows the 'Model Configuration' tab of the MIX{WILD} software. The interface includes a navigation bar with tabs: Model Configuration, Stage 1 Configuration, Stage 2 Configuration, Stage 1 Results, Stage 2 Results, View Model, View Data, Postestimation, and Help. The main content area is divided into sections for dataset information, Stage 1 model settings, and Stage 2 model settings. At the bottom, there is a logo for MIX{WILD} and three buttons: Save Model, Reset, and Submit.

Is your dataset Mix{WILD} friendly? [Check here...](#)

Data File:  [Browse](#)

Title:

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as?

---

Stage 1 outcome: ☒ Continuous ☐ Dichotomous ☐ Ordinal [?](#)

Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

---

Include Stage 2 model: ☐ Yes ☒ No [?](#)

**MIX{WILD}**  
Mixed Model Analysis With Intensive Longitudinal Data

[Save Model](#) [Reset](#) [Submit](#)

Select PA as the Stage 1 Outcome, select linear association

**Selected model configuration:**  
Random location effects: Intercept  
Stage 2 outcome: None

ID Variable:  
ID

Stage 1 Outcome:  
PA

Configure Stage 1 Regressors ...

Options ...

Specify the relationship between the mean and WS variance.

☐ No Association  
☒ Linear Association  
☐ Quadratic Association

**MIX {WILD}**  
Mixed-Effects Modeling Software for Windows and Mac

**Stage 1 Regressors**

	Mean	BS Variance	WS Variance
Level-1			
Level-2			

Save Model Clear Stage 1 Run Stage 1

Select Day\_c as PA as a time-varying predictor, click Submit

The screenshot displays the MIX{WILD} software interface. The main window is titled "MIX{WILD} Mixed Model Analysis With Intensive Longitudinal Data". The "Stage 1 Configuration" tab is active. The "Selected model configuration:" section shows "Random location effects: Intercept" and "Stage 2 outcome: None". The "ID Variable:" is set to "ID". The "Stage 1 Outcome:" is set to "PA". The "Specify the relationship between the mean and WS variance." section has three radio buttons: "No Association", "Linear Association" (selected), and "Quadratic Association". The "Add Stage 1 Regressors" dialog box is open, showing a list of variables: Day, Sex, Age, Sem, SQ, PhysAct, NA, LGA, Exam, HSG, and BDI. The "Level-1 (Time Varying)" pane contains "Day\_c". The "Level-2 (Time Invariant)" pane is empty. The dialog box has "Add" and "Remove" buttons for each pane, and "Cancel", "Reset", and "Submit" buttons at the bottom. The background window has buttons for "Configure Stage 1 Regressors ...", "Options ...", "Save Model", "Clear Stage 1", and "Run Stage 1".

Model Configuration Stage 1 Configuration Stage 2 Configuration Stage 1 Results Stage 2 Results View Model View Data Postestimation Help

Selected model configuration:  
Random location effects: Intercept  
Stage 2 outcome: None

ID Variable:  
ID

Stage 1 Outcome:  
PA

Configure Stage 1 Regressors ...

Options ...

Specify the relationship between the mean and WS variance.

☐ No Association  
☒ Linear Association  
☐ Quadratic Association

MIX{WILD}  
Mixed Model Analysis With Intensive Longitudinal Data

Add Stage 1 Regressors

Variables

Day  
Sex  
Age  
Sem  
SQ  
PhysAct  
NA  
LGA  
Exam  
HSG  
BDI

Add  
Remove

Level-1 (Time Varying)  
Day\_c

Level-2 (Time Invariant)

Add  
Remove

Cancel Reset Submit

Save Model Clear Stage 1 Run Stage 1

Mean	BS Variance	WS Variance
------	-------------	-------------

Model Configuration Stage 1 Configuration Stage 2 Configuration Stage 1 Results Stage 2 Results View Model View Data Postestimation Help

Selected model configuration:  
Random location effects: Intercept  
Stage 2 outcome: None

ID Variable:  
ID

Stage 1 Outcome:  
PA

Configure Stage 1 Regressors ...

Options ...

Specify the relationship between the mean and WS variance.

☐ No Association

☒ Linear Association

☐ Quadratic Association

**MIX {WILD}**  
Mixed Model Analysis With Interactive Randomized Trials

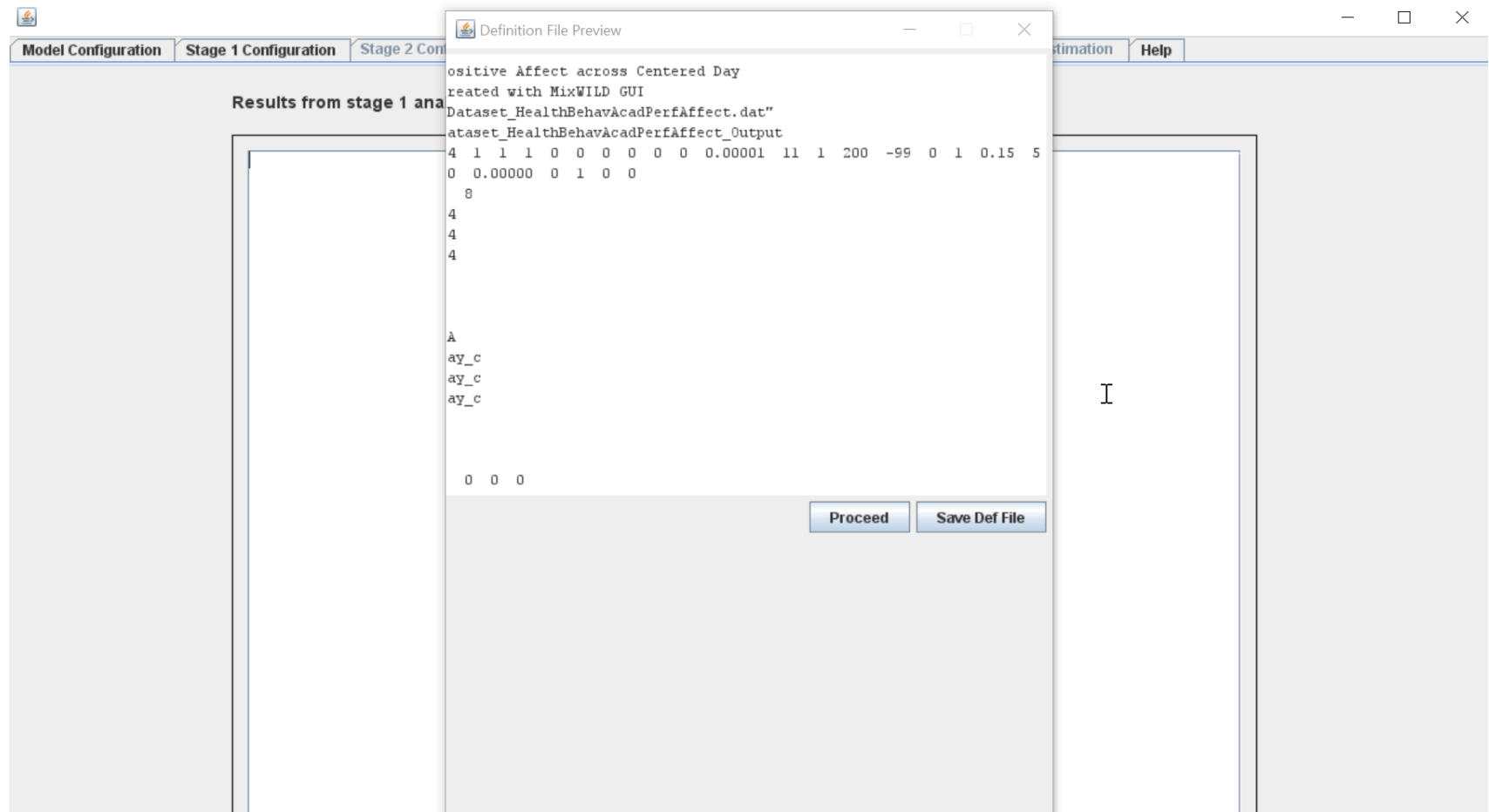
### Stage 1 Regressors

	Mean	BS Variance	WS Variance
<b>Level-1</b>			
Day_c	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Disaggregate?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
<b>Level-2</b>			

Save Model Clear Stage 1 Run Stage 1

Add Day\_c to the mean, BS variance, and WS variance models; click on Run Stage 1

DEF file is created, click on Proceed



MIXREGLS: Mixed-effects Location Scale Model with BS and WS variance models

-----  
mixREGLS.DEF specifications  
-----

Positive Affect across Centered Day  
Created with MixWILD GUI

data and output files:  
Dataset\_HealthBehavAcadPerfAffect\_Output  
Dataset\_HealthBehavAcadPerfAffect\_Output\_1.out\_1.out

CONVERGENCE CRITERION = 0.00001000  
RIDGEIN = 0.1500  
NQ = 11  
QUADRATURE = 1 (0=non-adaptive, 1=adaptive)  
MAXIT = 200

-----  
Descriptives  
-----

Number of level-1 observations = 2109

Number of level-2 clusters = 72



Number of level-1 observations for each level-2 cluster

27	32	32	32	32	31	31	32	32	30	20	32	32
31	30	32	31	32	27	32	29	32	32	31	32	26
22	29	27	32	32	32	27	32	32	8	17	32	32
30	20	32	32	30	28	32	32	32	29	32	32	32
28	30	32	32	31	32	32	32	32	11	26	30	32
31	32	19	30	16	30	32						

Dependent variable

	mean	min	max	std dev
PA	4.1841	1.0000	7.0000	1.6107

Mean model covariates

	mean	min	max	std dev
Intercept	1.0000	1.0000	1.0000	0.0000
Day_c	0.0160	-2.2143	2.2143	1.3061

BS variance model covariates

	mean	min	max	std dev
Intercept	1.0000	1.0000	1.0000	0.0000
Day_c	0.0160	-2.2143	2.2143	1.3061

WS variance model covariates

	mean	min	max	std dev
Intercept	1.0000	1.0000	1.0000	0.0000
Day_c	0.0160	-2.2143	2.2143	1.3061

```

-----
Model WITH RANDOM Scale
-----

```

==> BAD NR ITERATION      9 with NEW ridge =    0.4500

Total Iterations = 15

Final Ridge value = 0.0

Log Likelihood                      =     -3099.477

Akaike's Information Criterion =     -3107.477

Schwarz's Bayesian Criterion    =     -3116.583

==> multiplied by -2

Log Likelihood                      =     6198.953

Akaike's Information Criterion =     6214.953

Schwarz's Bayesian Criterion    =     6233.167

Variable	Estimate	AsymStdError	z-value	p-value
-----				
BETA (regression coefficients)				
Intercept	4.13067	0.14192	29.10653	0.00000
Day_c	-0.10365	0.01829	-5.66789	0.00000
ALPHA (BS variance parameters: log-linear model)				
Intercept	0.33310	0.17051	1.95351	0.05076
Day_c	0.11952	0.01981	6.03332	0.00000
TAU (WS variance parameters: log-linear model)				
Intercept	-0.09005	0.10237	-0.87967	0.37904
Day_c	0.13537	0.02615	5.17739	0.00000
Random scale standard deviation				

Std Dev	0.73614	0.07047	10.44576	0.00000
Random location (mean) effect on WS variance				
Loc Eff	-0.36749	0.09955	-3.69164	0.00022

#### BS variance ratios and 95% CIs

Variable	Ratio	Lower	Upper
ALPHA (BS variance parameters: log-linear model)			
Intercept	1.39529	0.99890	1.94898
Day_c	1.12695	1.08404	1.17157

#### WS variance ratios and 95% CIs

Variable	Ratio	Lower	Upper
TAU (WS variance parameters: log-linear model)			
Intercept	0.91388	0.74774	1.11694
Day_c	1.14496	1.08776	1.20516
Random location (mean) effect on WS variance			
Location Effect	0.69247	0.56972	0.84166
Random scale standard deviation			
Std Dev	2.08785	1.81850	2.39710

## Interpretation of Results: Model with RANDOM scale

- Using the centered and scaled version Day\_c, the intercepts represent the average across days and the slopes for day represent change per week.
- Mean model: the mean PA is estimated to be a bit over 4, and the slope is negative and significant ( $\hat{\beta} = -0.10365, p = 0.00001$ ). PA decreases by approximately one-tenth of a point per week.
- BS variance model: this is a log-linear model, so in addition to the estimates from the log-linear model, the program provides exponentiated estimates as well. From these, the BS variance is estimated to be 1.39529 with a 95% confidence interval of 0.99890 to 1.94898. The effect of Day\_c is positive and significant ( $\hat{\alpha} = 0.11952, p = 0.00001$ ). The exponentiated slope is 1.12695 with a 95% confidence interval of 1.08404 to 1.17157. The exponentiated slope represents a variance ratio (ratio of BS variance comparing values one week apart, for example the BS variance at week 2 divided by the BS variance at week 1). From the estimate of 1.13, we can conclude that the BS variance increases by a factor of 13% per week; thus, subjects become more heterogeneous over time.
- WS variance model: this is also a log-linear model, so in addition to the estimates from the log-linear model, the program provides exponentiated estimates. From these, the WS variance is estimated to be 0.91388 with a 95% confidence interval of 0.74774 to 1.11694. The effect of Day\_c is positive and significant ( $\hat{\tau} = 0.13537, p = 0.00001$ ). The exponentiated slope is 1.14496 with a 95% confidence interval of 1.08776 to 1.20516. The exponentiated slope represents a variance ratio (ratio of WS variance comparing values one week apart, for example the WS variance at week 2 divided by the WS variance at week 1). From the estimate of 1.15, we can conclude that the WS variance increases by a factor of 15% per week; thus, subjects exhibit more erraticism (less consistency) over time.
- The standard deviation of the random scale effect is estimated to be 0.73614, and this is a highly significant effect. Thus, subjects vary considerably in terms of how consistent/erratic they are in their PA reports. The relationship between the random location and scale effects is negative and significant ( $\hat{\tau} = -0.36749, p = 0.00022$ ) indicating that subjects with higher average PA are also more consistent, and subjects with lower average PA are more erratic.

## MELS summary (Stage 1)

- Mixed models (aka multilevel or hierarchical linear models) useful for analysis of intensive longitudinal data.
- Intensive longitudinal data and the mixed-effects location scale model allow one to consider modeling of the between-subjects and within-subjects variances in terms of covariates.
  - What subject and/or contextual variables associated with subject homogeneity/heterogeneity?
  - What subject and/or contextual variables associated with within-subject consistency/erraticism?
- Model and software can also allow for multiple random subject effects of location (intercept and slope).
- Random effects can be considered as predictors of stage-2 subject-level outcomes (continuous, binary, ordinal, nominal, count) using plausible values replications.

## Stage 2 analysis (optional, but hopefully useful)

Stage 1 random subject effect estimates (e.g., intercept  $\hat{v}_{0i}$ , slope  $\hat{v}_{1i}$ , scale  $\hat{\omega}_i$ ) and other subject-level variables  $\mathbf{x}_i$  can be used as regressors and interaction terms to predict a Stage 2 subject-level outcome  $y_i$

- Multiple regression for continuous subject-level outcome

$$y_i = \beta_0 + \beta_1 \hat{v}_{0i} + \beta_2 \hat{v}_{1i} + \beta_3 \hat{\omega}_i + \mathbf{x}_i' \boldsymbol{\beta} + \varepsilon_i$$

- Logistic regression for binary/ordinal/nominal subject-level outcome; Poisson regression for subject-level count outcome
- Multilevel (random-intercept) Stage 2 model is also possible

Since the random subject effects are estimates with estimated uncertainty, “plausible value” replications of the the random effects are performed (Mislevy, 1991, *Psychometrika*); akin to multiple imputation for missing values.

Yes Include Stage 2 model, no separate data file, single level, dichotomous outcome

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

① CSV file path: iSets\Dataset\_HealthBehavAcadPerfAffect.csv **Change Dataset**

Title (optional): MELS to Stage 2 model

① Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as? -99

**Stage 1 Model**

① Stage 1 outcome: ☒ Continuous ☐ Dichotomous ☐ Ordinal

① Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

① Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

① Include Stage 2 model: ☒ Yes ☐ No

Include separate Stage 2 data file: ☐ Yes ☒ No

Stage 2 CSV file path: **Import Dataset**

① Stage 2 model type: ☒ Single level ☐ Multilevel

① Stage 2 outcome: ☐ Continuous ☒ Dichotomous/Ordinal ☐ Count ☐ Multinomial

① Set a seed for Stage 2 resampling (optional): 23041

**Save Model** **Reset** **Continue**

All as before; click on Configure Stage 2

The screenshot shows the MixWILD-2.0 software interface with the 'Stage 1 Configuration' tab selected. The interface is divided into several sections:

- Model Configuration:** Includes tabs for 'Model Configuration', 'Stage 1 Configuration', 'Stage 2 Configuration', 'View Data', and 'Help'.
- Selected Model Configuration:** Displays 'Stage 1 model: Intercept Only' and 'State 1 outcome: Continuous'.
- ID Variable:** A dropdown menu showing 'ID'.
- Stage 1 Outcome:** A dropdown menu showing 'PA'.
- Buttons:** 'Configure Stage 1 Regressors ...' and 'Options ...'.
- Relationship Selection:** Radio buttons for 'No Association', 'Linear Association' (selected), and 'Quadratic Association'.
- Stage 1 Regressors:** A table with columns 'Mean', 'BS Variance', and 'WS Variance' for 'Level-1' and 'Level-2'.
- Bottom Buttons:** 'Save Model', 'Clear Stage 1', and 'Configure Stage 2'.

**Stage 1 Regressors Table:**

	Mean	BS Variance	WS Variance
<b>Level-1</b>			
Day_c	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Disaggregate?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
<b>Level-2</b>			



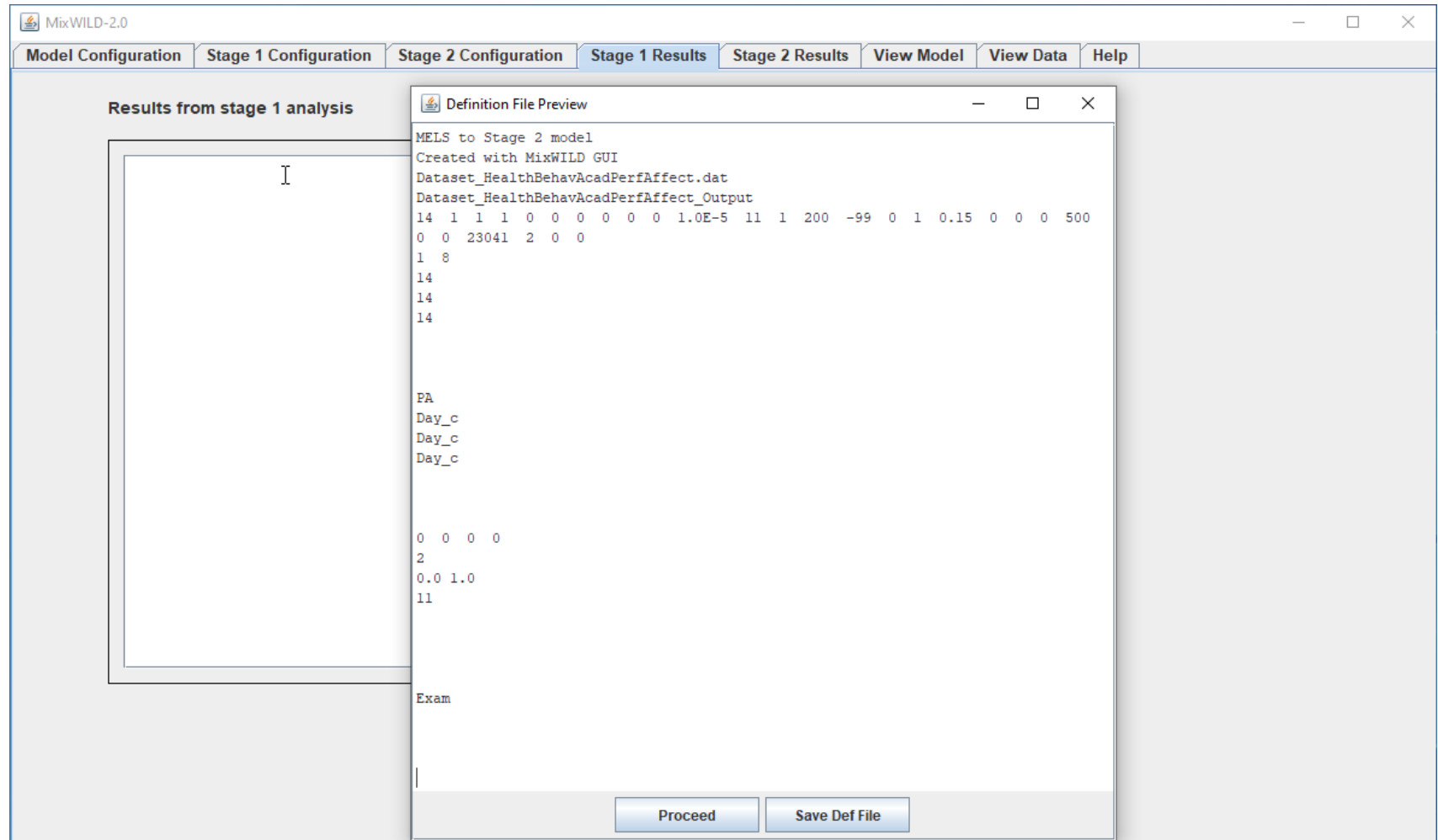
Select Exam as Stage 2 Outcome; Run Stage 1 and 2

The screenshot displays the MixWILD-2.0 software interface, specifically the Stage 2 Configuration tab. The window has a title bar with the text "MixWILD-2.0" and standard window controls. Below the title bar is a menu bar with options: "Model Configuration", "Stage 1 Configuration", "Stage 2 Configuration" (which is the active tab), "View Data", and "Help".

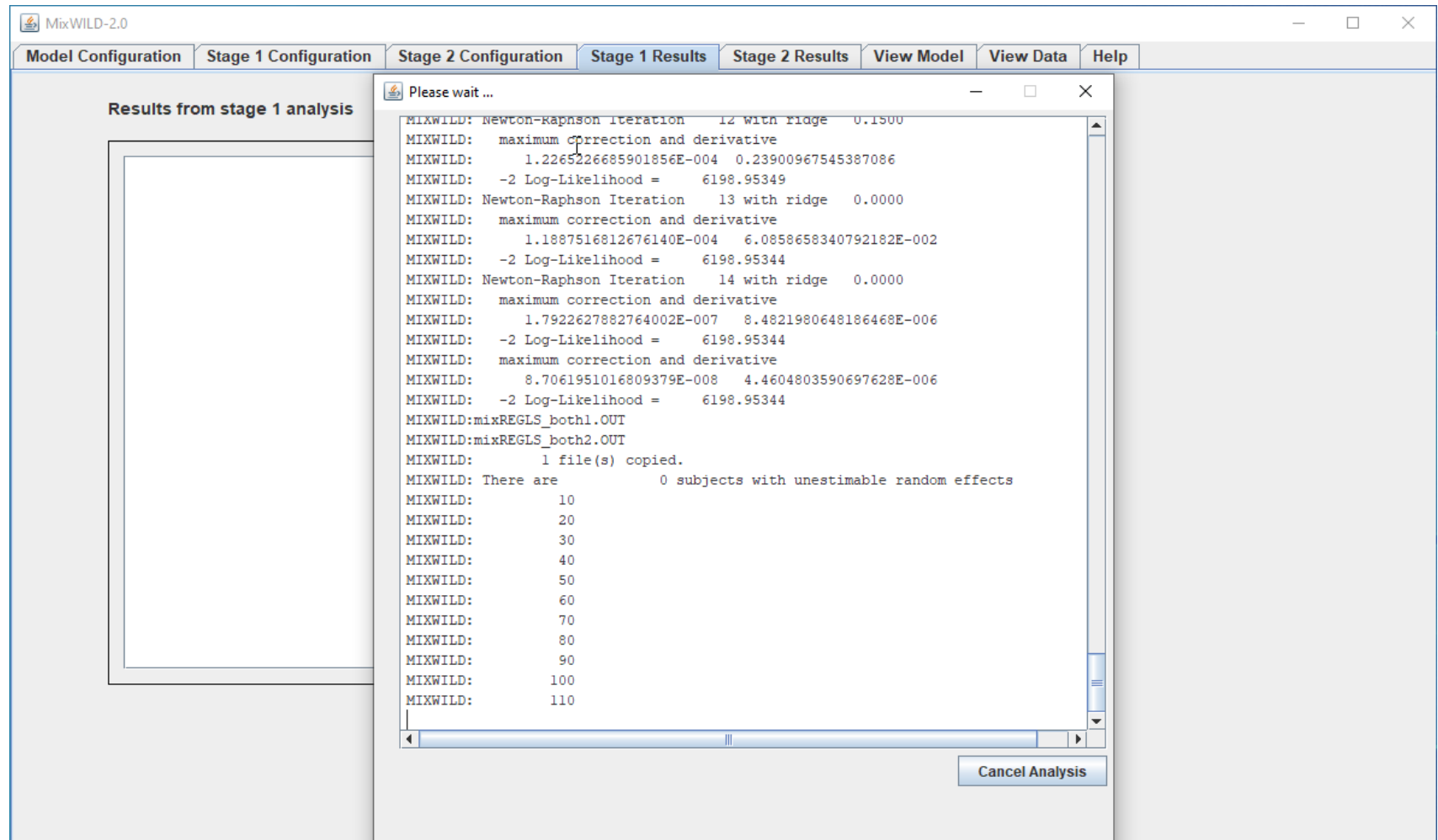
The main content area is divided into two sections. On the left, under the heading "Selected Model Configuration", the following settings are listed: "Stage 1 model: Intercept Only", "Stage 1 outcome: Continuous", "Stage 2 model type: Single-level", "Stage 2 outcome: Dichot/Ord", and "Number of resamples (stage 2): 500". Below this list, there is a section for "Stage 2 Outcome:" with a dropdown menu currently set to "Exam". Underneath the dropdown are two buttons: "Configure Stage 2 Regressors..." and "Check outcome categories".

On the right side of the window, the "Stage 2 Interactions" section is visible. It contains two large, empty rectangular boxes labeled "Level-1" and "Level-2". Above each box are four tabs: "Main Effects", "Random Location", "Random Scale", and "Location X Scale". Below the "Level-2" box, there is a checkbox labeled "Suppress 2-way Location X Scale Interaction" which is currently unchecked. At the bottom right of the window, there are three buttons: "Save Model", "Clear Stage 2", and "Run Stage 1 and 2".

Click on Proceed



After completing Stage 1, it performs 500 logistic regressions



## Stage 2 Results: descriptive statistics

The screenshot displays the MixWILD-2.0 software window with the 'Stage 2 Results' tab selected. The main content area shows the results of a stage 2 analysis, including descriptive statistics and final model results. A 'Save Results As ...' button is located at the bottom right of the window.

**Results from stage 2 analysis**

Level 2 units = 72

-----  
Descriptives  
-----

Categories of the Dependent Variable

Category	Frequency	Proportion
0.00	38.00	0.52778
1.00	34.00	0.47222

Random Location and Scale EB mean estimates

	mean	min	max	std dev
Locat_1	0.0146	-1.7555	2.0856	0.9921
Scale	-0.0000	-3.3078	1.8765	0.9385
Locat_1*Scale	-0.0131	-5.3343	2.3261	1.0731

There are 0 subjects with unestimable random effect values

Number of replications = 500

-----  
Final Results  
-----

Average Log Likelihood	=	-47.107 (sd= 1.038)
Akaike's Information Criterion	=	-51.107
Schwarz's Bayesian Criterion	=	-55.660

Save Results As ...

## Stage 2 Results: no significant effects on exam success

MixWILD-2.0

Model Configuration Stage 1 Configuration Stage 2 Configuration Stage 1 Results Stage 2 Results View Model View Data Help

**Results from stage 2 analysis**

Locat_1	0.0146	-1.7555	2.0856	0.9921
Scale	-0.0000	-3.3078	1.8765	0.9385
Locat_1*Scale	-0.0131	-5.3343	2.3261	1.0731

There are 0 subjects with unestimable random effect values

Number of replications = 500

-----  
Final Results  
-----

Average Log Likelihood = -47.107 (sd= 1.038)  
Akaike's Information Criterion = -51.107  
Schwarz's Bayesian Criterion = -55.660

==> multiplied by -2  
Log Likelihood = 94.214  
Akaike's Information Criterion = 102.214  
Schwarz's Bayesian Criterion = 111.320

Variable	Estimate	AsymStdError	z-value	p-value
Intercept	-0.12156	0.24941	-0.48739	0.62599
Locat_1	0.00813	0.29077	0.02798	0.97768
Scale_1	0.32521	0.32502	1.00059	0.31702
Locat_1*Scale	-0.54475	0.34826	-1.56420	0.11777

Save Results As ...

## MELS Model References - some are available at <https://voices.uchicago.edu/hedeker>

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# Running Mixed-Effects Location Scale (MELS) Models on MixWILD

Wei-Lin Wang

2023.04.26 @ SBM Workshop

Supported by National Cancer Institute grant R01CA240713 (Hedeker & Dunton)



# Learning Objects

- Understand what MixWILD is and why we need MixWILD
- Learn the basic settings of MixWILD software
- Utilize MixWILD to build your own models to address your research questions and interpret results
  - Stage 1 model
  - Two-stage model approach (Stage 1 & Stage 2 Models)
  - Try your own model (Optional)
- Learn some handy ways of MixWILD troubleshooting

# What is MixWILD?





# What is MixWILD

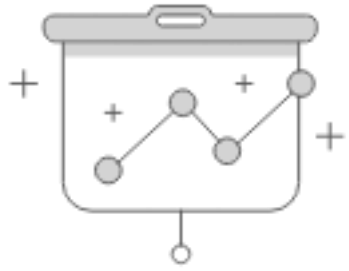
- **M**ixed model analysis **W**ith **I**ntensive **L**ongitudinal **D**ata (MixWILD)
- It is a standalone and user-friendly statistical software program, using a Java platform for Windows and Mac.
- The software consists of a front-end GUI (i.e., view and controller) and backend data processing (i.e., model).



# What can MixWILD do

- MixWILD allows one to examine the effects of subject-level parameters (intercept, slope(s), and scale) comprised of time-varying variables on a subject-level outcome or an outcome nested within time or clusters.
- This is specifically in the context of studies using intensive sampling methods, such as ecological momentary assessment (EMA).

# Why do we need MixWILD?



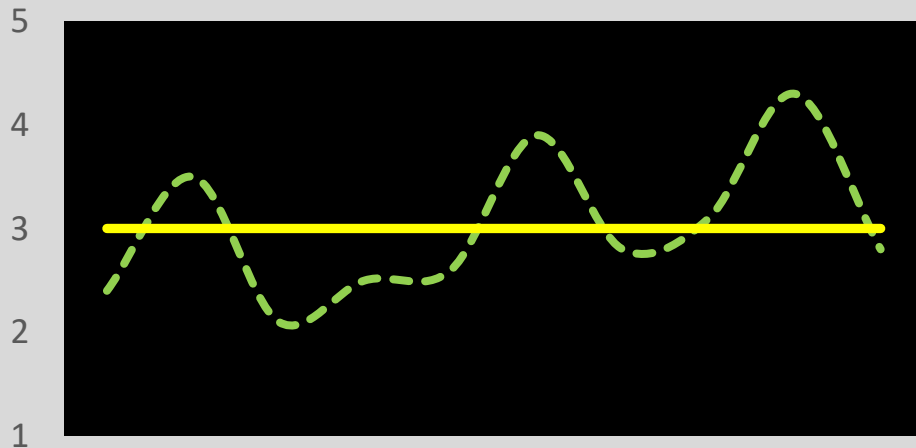
# Novel Stat Project

Novel Statistical Models for EMA Studies of Physical Activity  
1R01HL121330 & R01CA240713  
(Dunton and Hedeker, Pls)

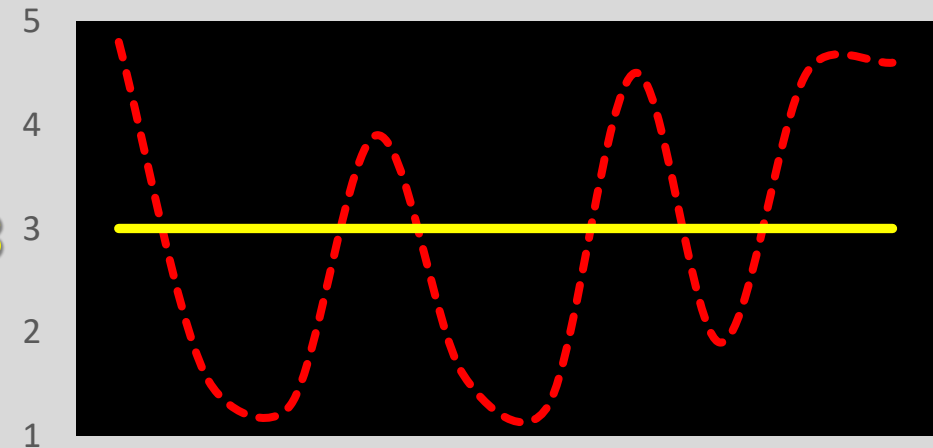
MixWILD is a handy and free standalone application to run Mixed-effects location and scale models.



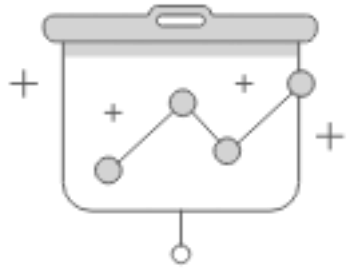
Subject A



Subject B



Mean = 3

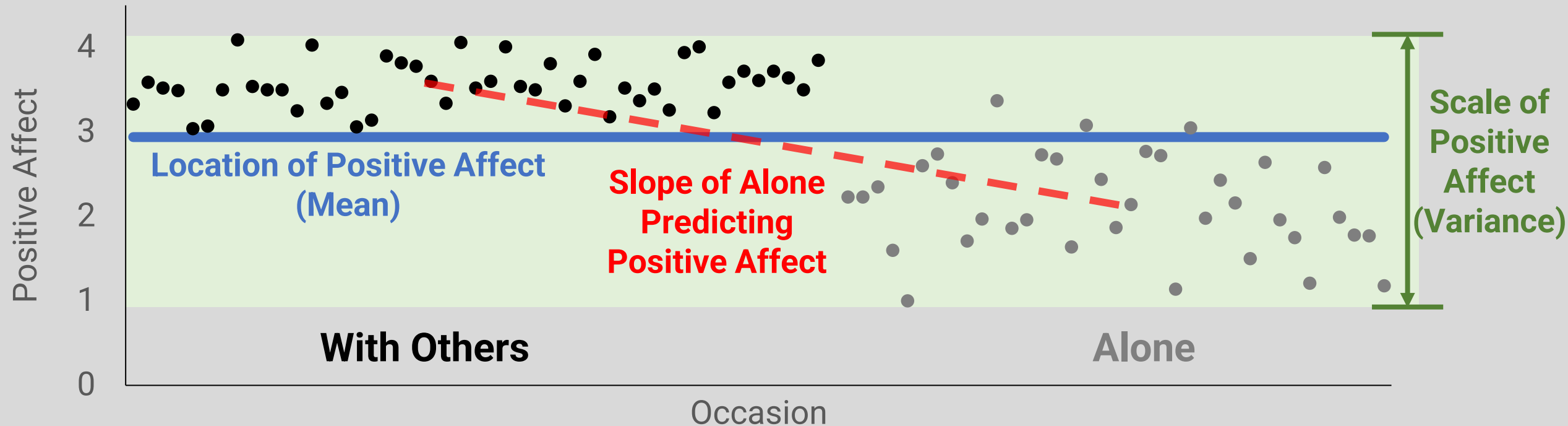


# Novel Stat Project

Novel Statistical Models for EMA Studies of Physical Activity  
1R01HL121330 & R01CA240713  
(Dunton and Hedeker, Pls)

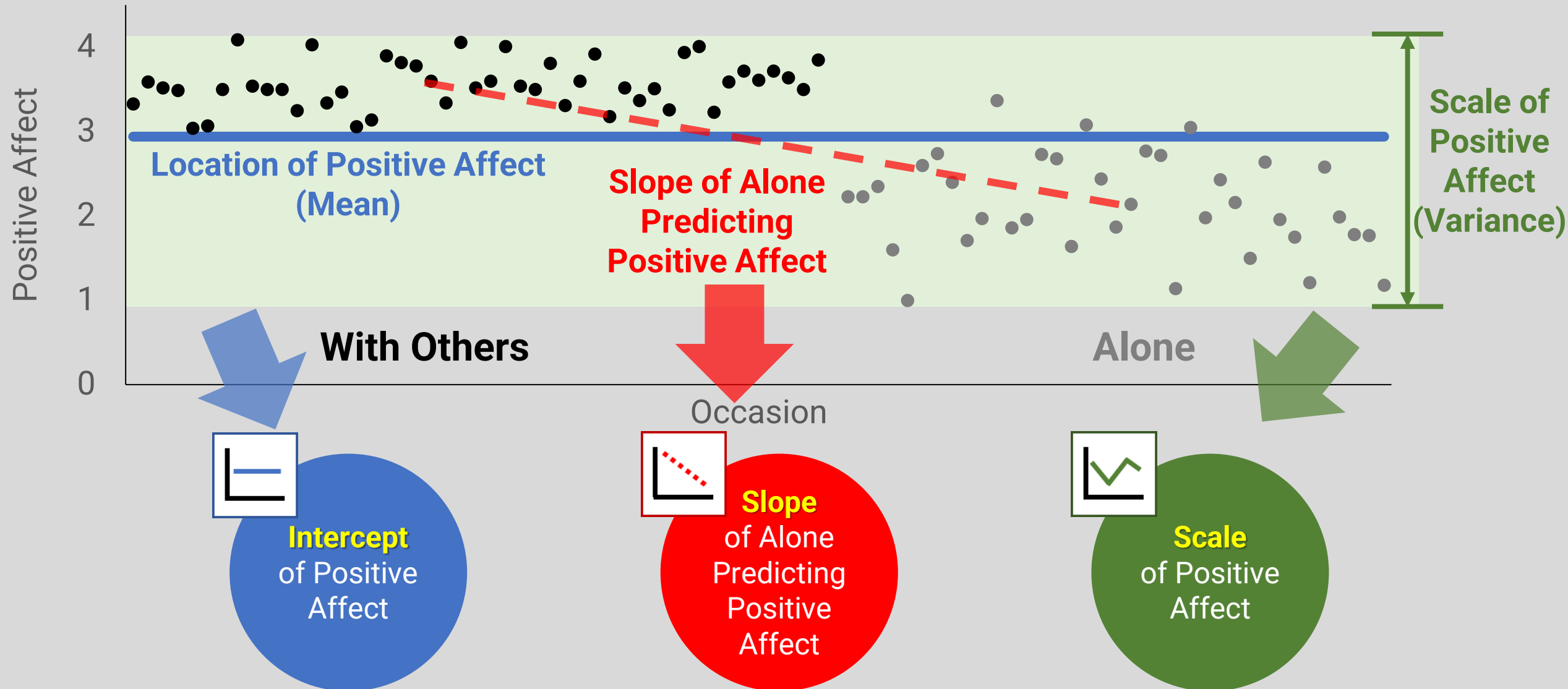
## MixWILD (Stage 1 Model)

- Go above and beyond the effects of mean!



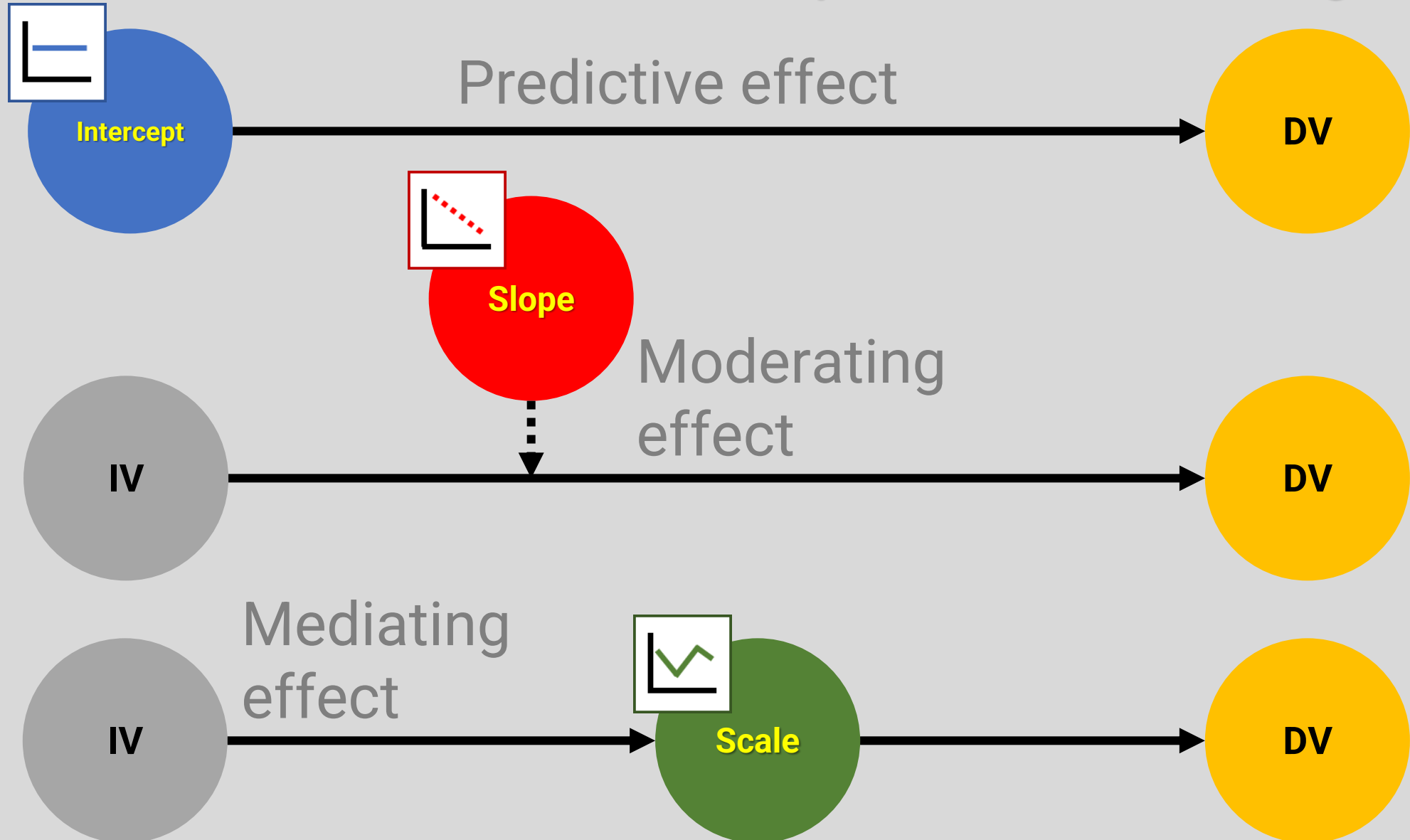
# MixWILD (Stage 2 Model)

- Extract the estimated random effects as regressors.



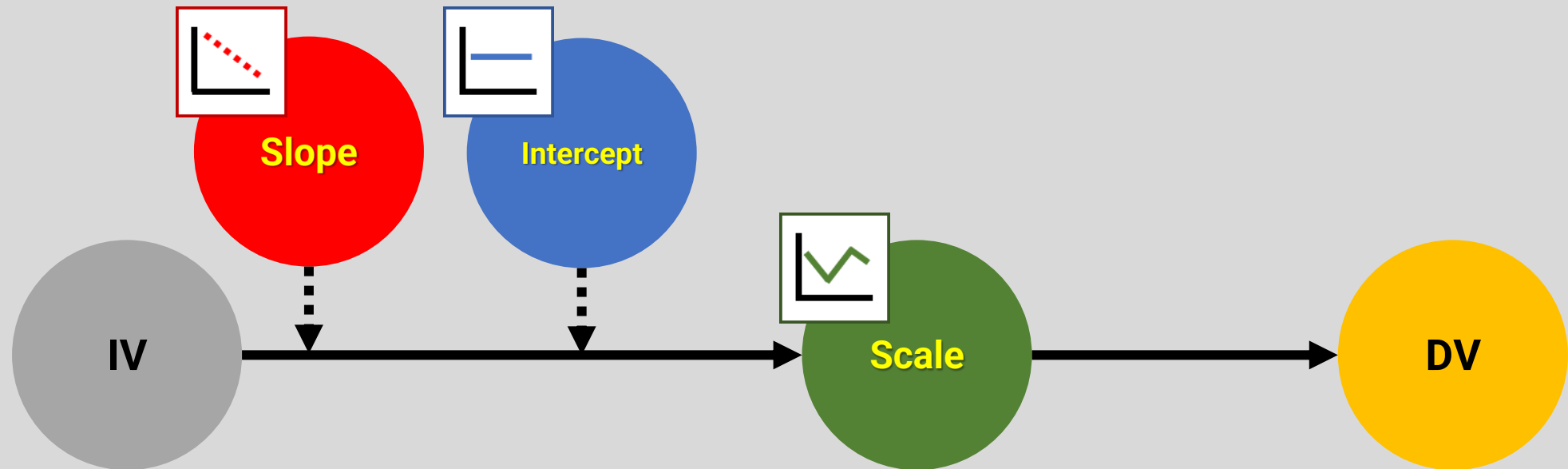
# MixWILD (Stage 2 Model)

- Test if the random effects can predict DV in Stage 2



# MixWILD (Stage 2 Model)

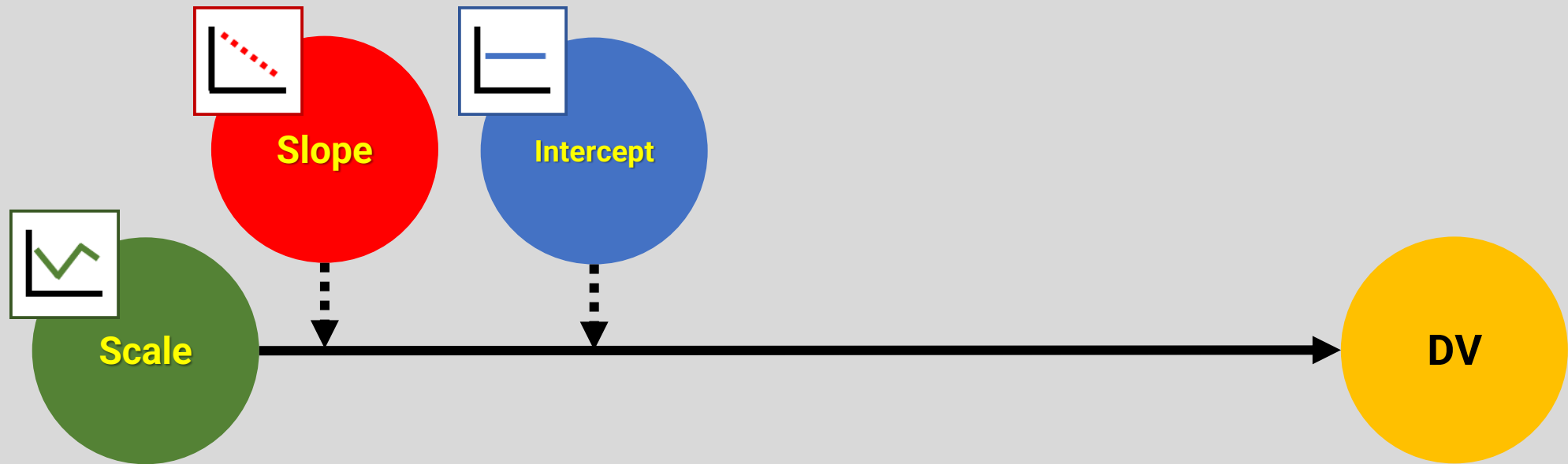
- Design your model and test associations  
(Use multiple random effects in the same model)





# MixWILD (Stage 2 Model)

- Design your model and test associations  
(Even with interactions among random effects!!!)



# Basic Settings



# Basic Settings

MixWILD-2.0

Model ConfigurationView DataHelp

Dataset

CSV file path:

:s\MixWILD\SBM\_MixWILD\_Example\_Data.csv

Change Dataset

Title (optional):

example

Does your data contain missing values?

☒ Yes ☐ No

What is your missing data coded as?

-99

Stage 1 Model

Stage 1 outcome:

☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type:

☐ Probit ☒ Logistic

Specify random location effects:

☒ Intercept only ☐ Intercept and slope(s)

Include estimates of random scale:

☒ Yes ☐ No

Stage 2 Model

Include Stage 2 model:

☐ Yes ☐ No

Save Model

Reset



# Basic Settings

Open a new .CSV file  
Load the file from your  
local address. Folder name  
**CANNOT** have **any blank**  
**SPACES**

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

① CSV file path: s:\MixWILD\SBM\_MixWILD\_Example\_Data.csv **Change Dataset**

Title (optional): example

① Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as? -99

**Stage 1 Model**

① Stage 1 outcome: ☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type: ☐ Probit ☒ Logistic

① Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

① Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

① Include Stage 2 model: ☐ Yes ☐ No

**Save Model** **Reset**



# Basic Settings

## Format missing value

Click on missing values if there are any in your dataset; specify the missing value code in the box.

Example: Missing = -99

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path: s:\MixWILD\SBM\_MixWILD\_Example\_Data.csv Change Dataset

Title (optional): example

**Does your data contain missing values?** ☒ Yes ☐ No

What is your missing data coded as? -99

**Stage 1 Model**

Stage 1 outcome: ☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type: ☐ Probit ☒ Logistic

Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

Include Stage 2 model: ☐ Yes ☐ No

Save Model Reset



# Basic Settings

## Select Stage 1 outcome

Select **Continuous**, **Dichotomous**, or **Ordinal** for Stage 1 outcome.

- **Continuous**: Weight;
- **Dichotomous**: Yes or No;
- **Ordinal**: Preference level

Choose between **Probit** or **Logistic** model if your Stage 1 outcome is **Dichotomous/Ordinal**.

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path: s\MixWILD\SBM\_MixWILD\_Example\_Data.csv **Change Dataset**

Title (optional): example

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as? -99

**Stage 1 Model**

Stage 1 outcome: ☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type: ☐ Probit ☒ Logistic

Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

Include Stage 2 model: ☐ Yes ☒ No

**Save Model** **Reset**



# Basic Settings

## Specify random location

Select "**Intercept only**" and the model includes a random subject intercept.

Select "**Intercept and slope(s)**" and the model includes a random subject intercept and random slope(s).

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path: s:\MixWILD\SBM\_MixWILD\_Example\_Data.csv [Change Dataset](#)

Title (optional): example

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as? -99

**Stage 1 Model**

Stage 1 outcome: ☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type: ☐ Probit ☒ Logistic

**Specify random location effects:** ☒ Intercept only ☐ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

Include Stage 2 model: ☐ Yes ☐ No

[Save Model](#) [Reset](#)



# Basic Settings

## Select random scale

Select “Yes” if the model includes **random subject scale** (allowing subjects to have individual within-subject variance effects); otherwise “No”.

The screenshot shows the 'Model Configuration' window of MixWILD-2.0. The window has three tabs: 'Model Configuration', 'View Data', and 'Help'. The 'Model Configuration' tab is active. The configuration is organized into three sections: 'Dataset', 'Stage 1 Model', and 'Stage 2 Model'. In the 'Dataset' section, the 'CSV file path' is set to 's:\MixWILD\SBM\_MixWILD\_Example\_Data.csv' and the 'Title (optional)' is 'example'. A 'Change Dataset' button is next to the file path. In the 'Stage 1 Model' section, 'Does your data contain missing values?' is set to 'Yes' with a missing data code of '-99'. 'Stage 1 outcome' is set to 'Ordinal', and 'Stage 1 regression type' is set to 'Logistic'. 'Specify random location effects' is set to 'Intercept only'. The 'Include estimates of random scale' option is highlighted with a red box and is set to 'Yes'. In the 'Stage 2 Model' section, 'Include Stage 2 model' is set to 'No'. At the bottom, there are 'Save Model' and 'Reset' buttons.

Section	Setting	Value
Dataset	CSV file path:	s:\MixWILD\SBM_MixWILD_Example_Data.csv
	Title (optional):	example
	Does your data contain missing values?	Yes
Stage 1 Model	What is your missing data coded as?	-99
	Stage 1 outcome:	Ordinal
	Stage 1 regression type:	Logistic
	Specify random location effects:	Intercept only
	Include estimates of random scale:	Yes
Stage 2 Model	Include Stage 2 model:	No





# Basic Settings

Select separate Stage 2 data Select “Yes” when your stage 2 data file is separate (need **ID** to link with stage 1).

Import Dataset for stage 2 separate data.

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path: s:\MixWILD\SBM\_MixWILD\_Example\_Data.csv **Change Dataset**

Title (optional): example

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as? -99

**Stage 1 Model**

Stage 1 outcome: ☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type: ☐ Probit ☒ Logistic

Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

Include Stage 2 model: ☒ Yes ☐ No

Include separate Stage 2 data file: ☐ Yes ☒ No

Stage 2 CSV file path: **Import Dataset**

Stage 2 model type: ☐ Single level ☒ Multilevel

Stage 2 outcome: ☒ Continuous ☐ Dichotomous/Ordinal ☐ Count ☐ Multinomial

Set a seed for Stage 2 resampling (optional): 777

**Save Model** **Reset** **Continue**



# Basic Settings

## Select stage 2 model

The stage-2 outcome can be single- or multilevel.

## Select stage 2 outcome

- **Continuous:** Weight;
- **Dichotomous:** Yes or No;
- **Count:** Times of having snacks/exercise per day
- **Nominal:** Types of physical activities

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path: s:\MixWILD\SBM\_MixWILD\_Example\_Data.csv **Change Dataset**

Title (optional): example

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as? -99

**Stage 1 Model**

Stage 1 outcome: ☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type: ☐ Probit ☒ Logistic

Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

Include Stage 2 model: ☒ Yes ☐ No

Include separate Stage 2 data file: ☐ Yes ☒ No

Stage 2 CSV file path: **Import Dataset**

Stage 2 model type: ☐ Single level ☒ Multilevel

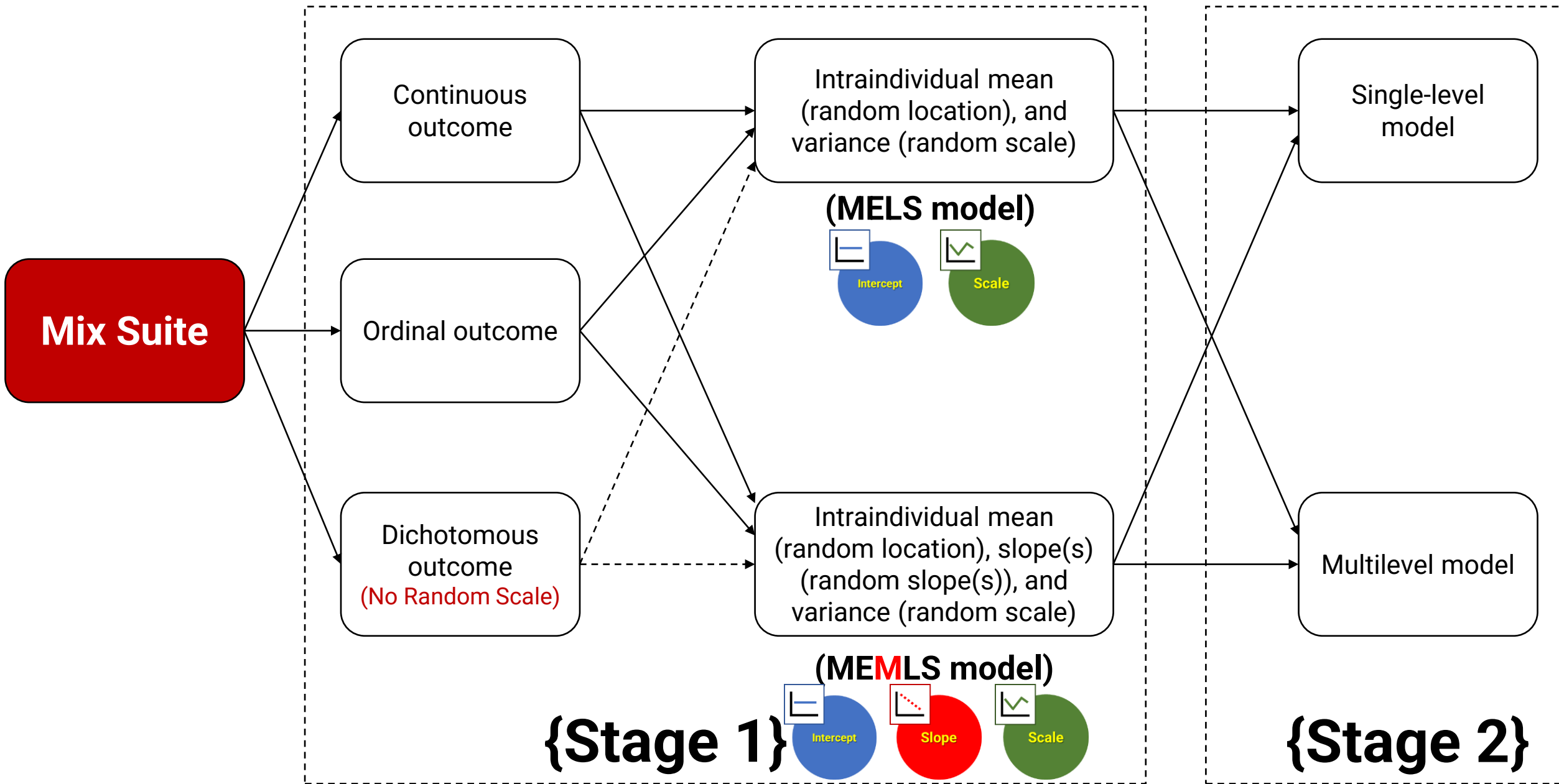
Stage 2 outcome: ☒ Continuous ☐ Dichotomous/Ordinal ☐ Count ☐ Multinomial

Set a seed for Stage 2 resampling (optional): 777

**Save Model** **Reset** **Continue**



# Basic Settings



# Exercise



# Data

Variable	Level	Description
ID	2	Subject number (1 to 82)
Day_C	1	Centered and scaled version of day (-2.21 to 2.21; 1 unit = 1 week); day-level
Sex_F	2	Dummy coded sex (0=M; 1=F)
Age_C	2	Participants' age, centered age (-6 to 28, mean=0, sd=9.15) Original scale is from 17 to 51 yrs.
Sem	2	Semester: Number of semesters studied (subject-level variable; 1 to 10, mean = 2.93, sd=1.89)
Exam	2	Examination success (0=fail, 1=pass); subject-level variable
HSG	2	High school grades 1 (lowest grade) to 6 (highest grade) Subject-level variable; (3.4 to 5.6; mean=4.68, sd=0.45)
HSG_Rank	2	Ranked version of HSG (good for stage-2 subject-level count outcome); subject-level variable
BDI	2	Beck Depression Inventory 1(not) 2 (mild to moderate) 3 (clinically relevant symptoms); subject-level variable
SQ	1	Sleep quality 1 (very bad) to 4 (very good) (day-level variable; mean = 3)
PhysAct	1	Physical activity: Number of minutes engaged in mild, moderate and strenuous exercise weighted by metabolic equivalents and then summed to produce a total daily leisure activity score (day-level variable; 0 to 3960, sd=413.68)
PhysAct_LN	1	Physical activity (Log term) (day-level variable; mean=3.92, sd=4.85)
PA	1	Positive affect 1 (not at all) to 7 (extremely); day-level
PA_D	1	Positive affect (Dichotomous) (Coded as 1 when PA > 4)
PA_Ord	1	Rounded version of PA (good for stage-1 ordinal outcome); day-level
NA	1	Negative affect 1 (not at all) to 7 (extremely); day-level
NA_D	1	Negative affect 1 (not at all) to 7 (extremely); day-level (Dichotomous) (Coded as 1 when NA > 4)
NA_Mean	2	Average negative affect per subject; subject-level variable (mean=2.65; sd=0.98)
LGA	1	Learning goal achievement 0 (not at all) to 4 (completely); day-level variable

**Missing value = -99**

Data are from: <https://dataverse.harvard.edu/dataverse/harvard>

“How health behaviors relate to academic performance via aect: An intensive longitudinal study” by Flueckiger L, Lieb R, Meyer AH, Mata J

## {Stage 1 Model}

Does the number of days in the study influence one's positive affect (PA)?



## {Stage 1 Model}

Does the number of days in the survey influence one's positive affect (PA)?

- (Mean model) Does positive affect change across days?
- (BSV model) Does the sample become more heterogeneous in PA as day passes?
- (WSV model) Do a subject's PA become more erratic as day passes?



Q1: Does the number of days in the study influence one's positive affect (PA)?

**Mix Suite**

Continuous outcome

Ordinal outcome

Dichotomous outcome  
(No Random Scale)

Intraindividual mean  
(random location), and  
variance (random scale)

**(MELS model)**



Intercept



Scale

Intraindividual mean  
(random location), slope(s)  
(random slope(s)), and  
variance (random scale)

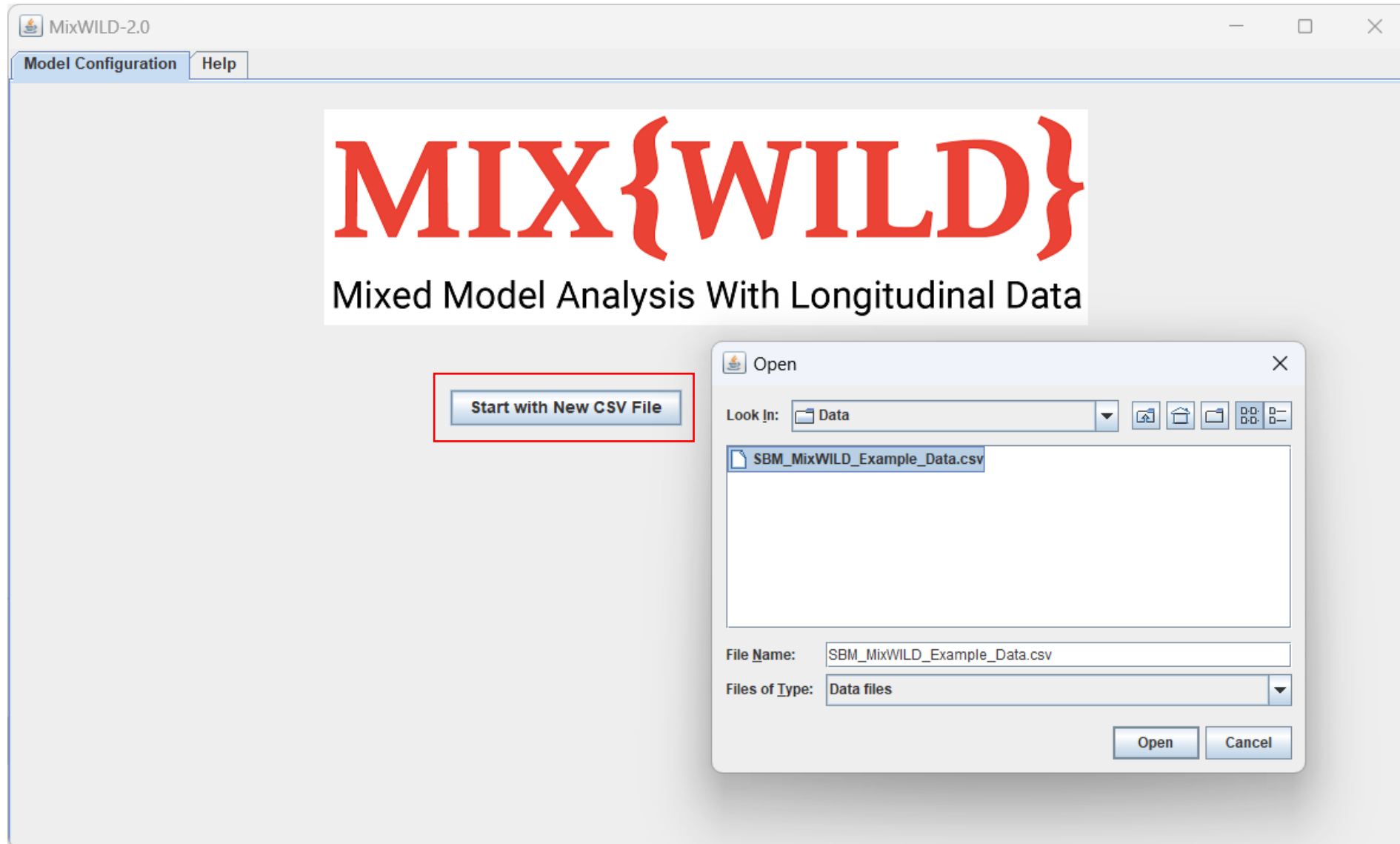
**(MEMLS model)**

**{Stage 1}**





Q1: Does the number of days in the study influence one's positive affect (PA)?



Start with “New CSV File” and locate the MixWILD example dataset



# Q1: Does the number of days in the study influence one's positive affect (PA)?

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path: :s\MixWILD\SBM\_MixWILD\_Example\_Data.csv [Change Dataset](#)

Title (optional): exercise1

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as? -99

**Stage 1 Model**

Stage 1 outcome: ☒ Continuous ☐ Dichotomous ☐ Ordinal

Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

Include Stage 2 model: ☐ Yes ☒ No

[Save Model](#) [Reset](#) [Continue](#)

Provide a title and make selections (Don't forget set up missing value)



# Q1: Does the number of days in the study influence one's positive affect (PA)?

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

① CSV file path: :s\MixWILD\SBM\_MixWILD\_Example\_Data.csv [Change Dataset](#)

Title (optional): exercise1

① Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as? -99

**Stage 1 Model**

① Stage 1 outcome: ☒ Continuous ☐ Dichotomous ☐ Ordinal

① Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

① Include estimates of random scale: ☒ Yes ☐ No

① Include Stage 2 model: ☐ Yes ☒ No

[Save Model](#) [Reset](#) [Continue](#)

Specify random effects (Select “Intercept only” and include “Random scale”)



# Q1: Does the number of days in the study influence one's positive affect (PA)?

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path:  [Change Dataset](#)

Title (optional):

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as?

**Stage 1 Model**

Stage 1 outcome: ☒ Continuous ☐ Dichotomous ☐ Ordinal

Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

Include Stage 2 model: ☐ Yes ☒ No

[Save Model](#) [Reset](#) [Continue](#)

Select “Continue” to enter the next page



# Q1: Does the number of days in the study influence one's positive affect (PA)?

MixWILD-2.0

Model Configuration Stage 1 Configuration **View Data** Help

Imported data file: SBM\_MixWILD\_Example\_Data.csv

ID	Day_C	Sex_F	Age_C	Sem	Exam	HSG	HSG_Rank	BDI	SQ	PhysAct	PhysAct_L
1	-2.21	1	-1	2	1	4.6	12	2	3	30	3.4
1	-2.07	1	-1	2	1	4.6	12	2	3	60	4.09
1	-1.93	1	-1	2	1	4.6	12	2	3	360	5.89
1	-1.79	1	-1	2	1	4.6	12	2	3	780	6.66
1	-1.5	1	-1	2	1	4.6	12	2	3	210	5.35
1	-1.21	1	-1	2	1	4.6	12	2	3	310	5.74
1	-1.07	1	-1	2	1	4.6	12	2	3	90	4.5
1	-0.93	1	-1	2	1	4.6	12	2	3	405	6
1	-0.79	1	-1	2	1	4.6	12	2	3	405	6
1	-0.64	1	-1	2	1	4.6	12	2	3	360	5.89
1	-0.5	1	-1	2	1	4.6	12	2	3	270	5.6
1	-0.36	1	-1	2	1	4.6	12	2	3	30	3.4
1	-0.21	1	-1	2	1	4.6	12	2	3	540	6.29
1	0.07	1	-1	2	1	4.6	12	2	2	405	6
1	0.21	1	-1	2	1	4.6	12	2	3	405	6
1	0.36	1	-1	2	1	4.6	12	2	3	540	6.29
1	0.64	1	-1	2	1	4.6	12	2	3	60	4.09
1	0.79	1	-1	2	1	4.6	12	2	3	405	6
1	0.93	1	-1	2	1	4.6	12	2	3	405	6
1	1.07	1	-1	2	1	4.6	12	2	2	60	4.09
1	1.21	1	-1	2	1	4.6	12	2	3	405	6
1	1.36	1	-1	2	1	4.6	12	2	3	330	5.8
1	1.5	1	-1	2	1	4.6	12	2	3	435	6.08
1	1.79	1	-1	2	1	4.6	12	2	3	525	6.26
1	1.93	1	-1	2	1	4.6	12	2	3	270	5.6
1	2.07	1	-1	2	1	4.6	12	2	3	405	6
1	2.21	1	-1	2	1	4.6	12	2	2	390	5.97
2	-2.21	1	-2	2	0	4.9	15	1	3	190	5.25

Review your data in “View Data” page



Q1: Does the number of days in the study influence one's positive affect (PA)?

MixWILD-2.0

Model Configuration Stage 1 Configuration View Data Help

**Selected Model Configuration**  
Stage 1 model: **Intercept Only**  
State 1 outcome: **Continuous**

ID Variable:  
ID

**Stage 1 Outcome:**  
PA

**Configure Stage 1 Regressors ...**

Options ...

Specify the relationship between the mean and WS variance.

☐ No Association  
☒ Linear Association  
☐ Quadratic Association

**Stage 1 Regressors**

Add Stage 1 Regressors

**Variables**

- Sex\_F
- Age\_C
- Sem
- Exam
- HSG
- HSG\_Rank
- BDI
- SQ
- PhysAct
- PhysAct\_LN
- PA\_D
- PA\_Ord
- NA
- NA\_D
- NA\_Mean
- LGA

**Level-1 (Time Varying)**

- Day\_C

**Level-2 (Time Invariant)**

Cancel Reset Submit

Select “PA” as the Stage 1 Outcome and “Day\_C” as a time-varying predictor



Q1: Does the number of days in the study influence one's positive affect (PA)?

MixWILD-2.0

Model Configuration Stage 1 Configuration View Data Help

Selected Model Configuration  
Stage 1 model: **Intercept Only**  
State 1 outcome: **Continuous**

ID Variable:  
ID

Stage 1 Outcome:  
PA

Configure Stage 1 Regressors ...

Options ...

Specify the relationship between the mean and WS variance.

☐ No Association  
☒ Linear Association  
☐ Quadratic Association

Stage 1 Regressors

	Mean	BS Variance	WS Variance
Level-1			
Day_C	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Disaggregate?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Level-2			

Save Model Clear Stage 1 Run Stage 1

Specify the relationship between the mean and WS variance, select “Linear”



# Q1: Does the number of days in the study influence one's positive affect (PA)?

MixWILD-2.0

Model Configuration Stage 1 Configuration View Data Help

**Selected Model Configuration**  
Stage 1 model: **Intercept Only**  
State 1 outcome: **Continuous**

**ID Variable:**  
ID

**Stage 1 Outcome:**  
PA

**Configure Stage 1 Regressors ...**  
**Options ...**

**Specify the relationship between the mean and WS variance.**

☐ No Association  
☒ Linear Association  
☐ Quadratic Association

**Stage 1 Regressors**

	Mean	BS Variance	WS Variance
<b>Level-1</b>			
Day_C	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Disaggregate?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
<b>Level-2</b>			

**Save Model** **Clear Stage 1** **Run Stage 1**

Specify the regressors in Stage 1 Models (Mean, BSV, and WSV)





Q1: Does the number of days in the study influence one's positive affect (PA)?

MixWILD-2.0

Model Configuration Stage 1 Configuration View Data Help

**Selected Model Configuration**  
Stage 1 model: **Intercept Only**  
State 1 outcome: **Continuous**

**ID Variable:**  
ID

**Stage 1 Outcome:**  
PA

**Configure Stage 1 Regressors ...**  
**Options ...**

**Specify the relationship between the mean and WS variance.**

☐ No Association  
☒ Linear Association  
☐ Quadratic Association

**Stage 1 Regressors**

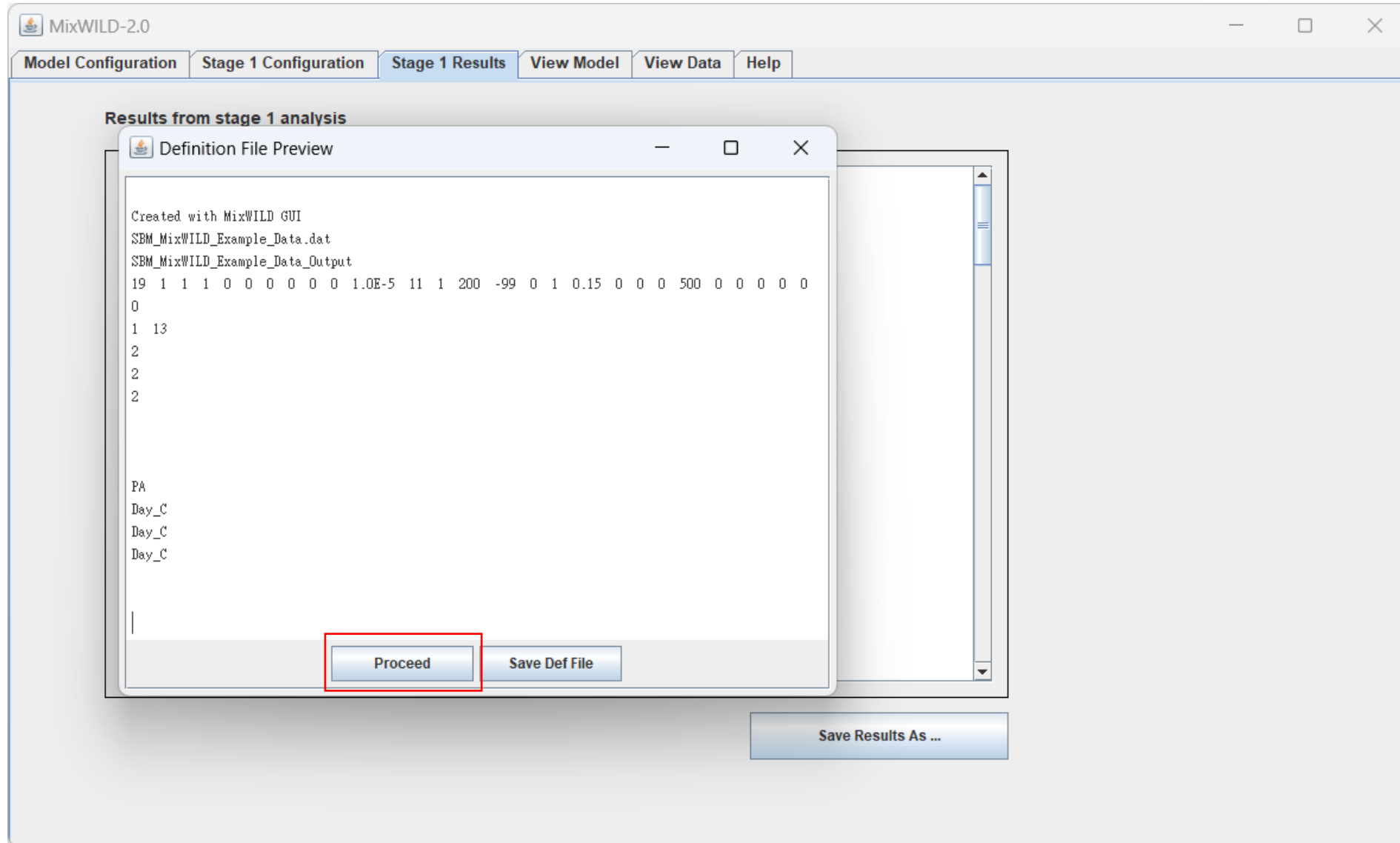
	Mean	BS Variance	WS Variance
<b>Level-1</b>			
Day_C	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Disaggregate?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
<b>Level-2</b>			

**Save Model** **Clear Stage 1** **Run Stage 1**

Click on “Run Stage 1”



Q1: Does the number of days in the study influence one's positive affect (PA)?



Click on “Proceed” and it will run the model automatically



Q1: Does the number of days in the study influence one's positive affect (PA)?

MixWILD-2.0

Model Configuration Stage 1 Configuration Stage 1 Results View Model View Data Help

Results from stage 1 analysis

```
MIXREGLS_both: Mixed-effects Location Scale Model
-----
mixREGLS_both.DEF specifications
-----

Created with MixWILD GUI

data and output files:
SBM_MixWILD_Example_Data.dat
SBM_MixWILD_Example_Data_Output_stagel.out

MULTIPLE LOCATION EFFECTS = F
SCALE EFFECT = T
CONVERGENCE CRITERION = 0.00001000
RIDGEIN = 0.1500
NQ = 11
QUADRATURE = 1 (0=non-adaptive, 1=adaptive)
MAXIT = 200

-----
Descriptives
-----

Number of level-1 observations = 2109

Number of level-2 clusters = 72

Number of level-1 observations for each level-2 cluster
27 32 32 32 32 31 31 32 32 30 20 32 32
```

Save Results As ...

Check the Stage 1 Results



## Q1: Does the number of days in the study influence one's positive affect (PA)?

Variable	Estimate	AsymStdError	z-value	p-value
-----	-----	-----	-----	-----
BETA (regression coefficients)				
intercept	4.13072	0.14192	29.10578	0.00000
Day_C	-0.10383	0.01829	-5.67717	0.00000
ALPHA (BS variance parameters: log-linear model)				
intercept	0.33320	0.17051	1.95409	0.05069
Day_C	0.11952	0.01981	6.03246	0.00000
TAU (WS variance parameters: log-linear model)				
intercept	-0.09010	0.10237	-0.88010	0.37881
Day_C	0.13547	0.02616	5.17843	0.00000
Random scale standard deviation				
Std Dev	0.73612	0.07047	10.44548	0.00000
Random location (mean) effect on WS variance				
Loc Eff	-0.36756	0.09955	-3.69233	0.00022

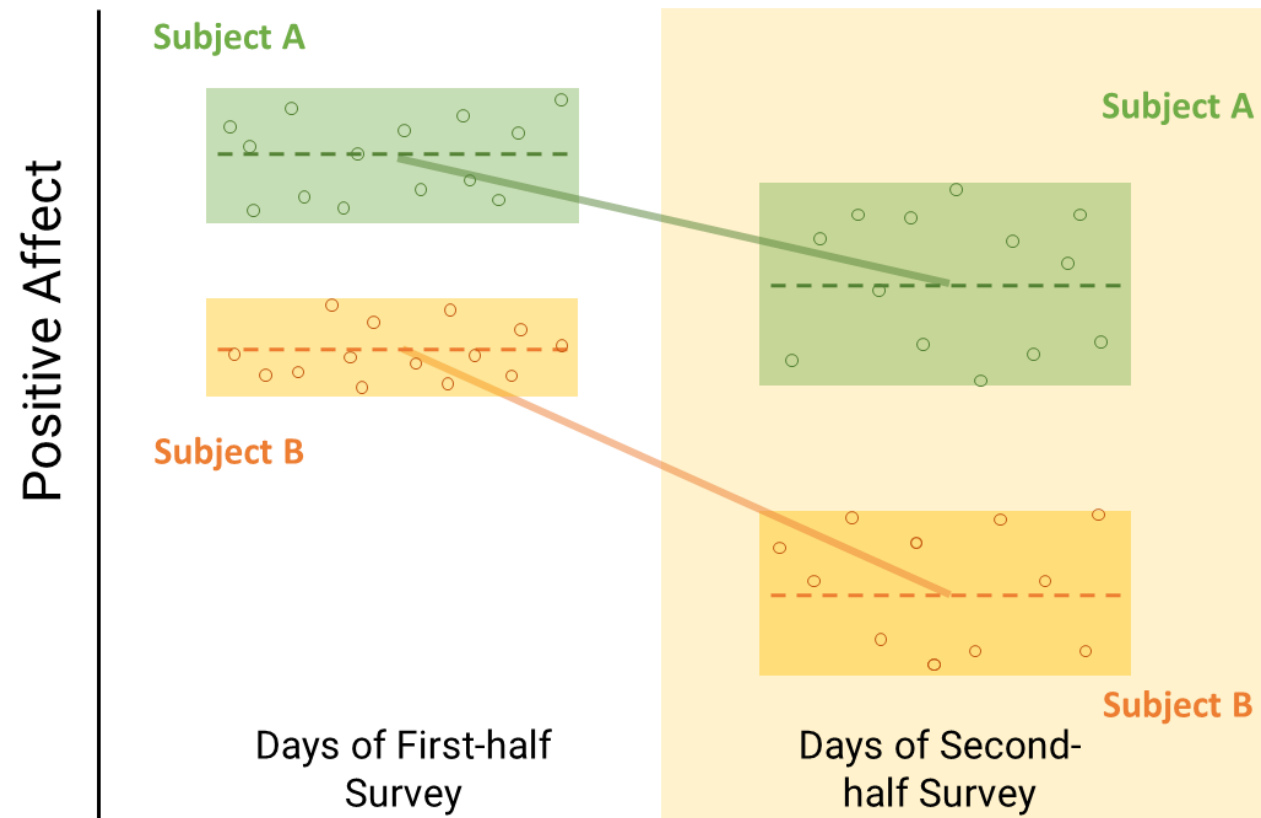
**Mean model (BETA):** the mean PA is estimated to be a bit over 4, and the slope is negative and significant (beta =  $-0.10383$ ;  $p < 0.001$ ). PA decreases by approximately one-tenth of a point per week.



## Q1: Does the number of days in the study influence one's positive affect (PA)?

(Mean model) Does positive affect change across days?

– the slope is negative and significant, and it shows PA decreases over time.





## Q1: Does the number of days in the study influence one's positive affect (PA)?

Variable	Estimate	AsymStdError	z-value	p-value
-----	-----	-----	-----	-----
BETA (regression coefficients)				
intercept	4.13072	0.14192	29.10578	0.00000
Day_C	-0.10383	0.01829	-5.67717	0.00000
ALPHA (BS variance parameters: log-linear model)				
intercept	0.33320	0.17051	1.95409	0.05069
Day_C	0.11952	0.01981	6.03246	0.00000
TAU (WS variance parameters: log-linear model)				
intercept	-0.09010	0.10237	-0.88010	0.37881
Day_C	0.13547	0.02616	5.17843	0.00000
Random scale standard deviation				
Std Dev	0.73612	0.07047	10.44548	0.00000
Random location (mean) effect on WS variance				
Loc Eff	-0.36756	0.09955	-3.69233	0.00022

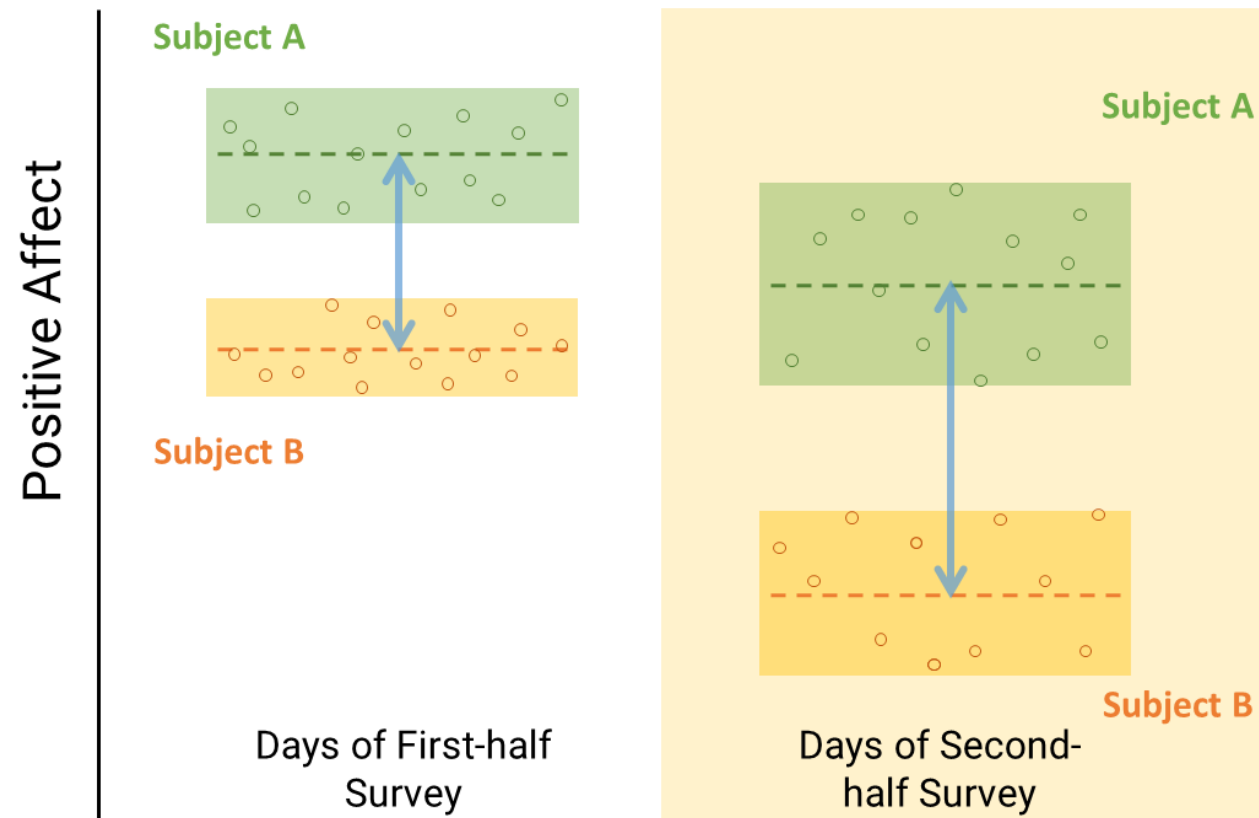
**BS variance model (ALPHA):** the effect of Day\_C is positive and significant ( $\alpha = 0.11952$ ;  $p < 0.001$ ). The exponentiated slope is 1.12696. From the estimate of 1.13, we can conclude that the BS variance increases by a factor of 13% per week; thus, subjects become more heterogeneous over time.



## Q1: Does the number of days in the study influence one's positive affect (PA)?

(BSV model) Does the sample become more heterogeneous in PA as day passes?

- The effect of Day\_C is positive and significant, and it indicates that subjects become more heterogeneous over time.





## Q1: Does the number of days in the study influence one's positive affect (PA)?

Variable	Estimate	AsymStdError	z-value	p-value
-----	-----	-----	-----	-----
BETA (regression coefficients)				
intercept	4.13072	0.14192	29.10578	0.00000
Day_C	-0.10383	0.01829	-5.67717	0.00000
ALPHA (BS variance parameters: log-linear model)				
intercept	0.33320	0.17051	1.95409	0.05069
Day_C	0.11952	0.01981	6.03246	0.00000
TAU (WS variance parameters: log-linear model)				
intercept	-0.09010	0.10237	-0.88010	0.37881
Day_C	0.13547	0.02616	5.17843	0.00000
Random scale standard deviation				
Std Dev	0.73612	0.07047	10.44548	0.00000
Random location (mean) effect on WS variance				
Loc Eff	-0.36756	0.09955	-3.69233	0.00022

**WS variance model (TAU):** the effect of Day\_C is positive and significant ( $\tau = 0.13547$ ;  $p < 0.001$ ). The exponentiated slope is 1.14507. From the estimate of 1.15, we can conclude that the WS variance increases by a factor of 15% per week; thus, subjects exhibit more erraticism (less consistency) over time.

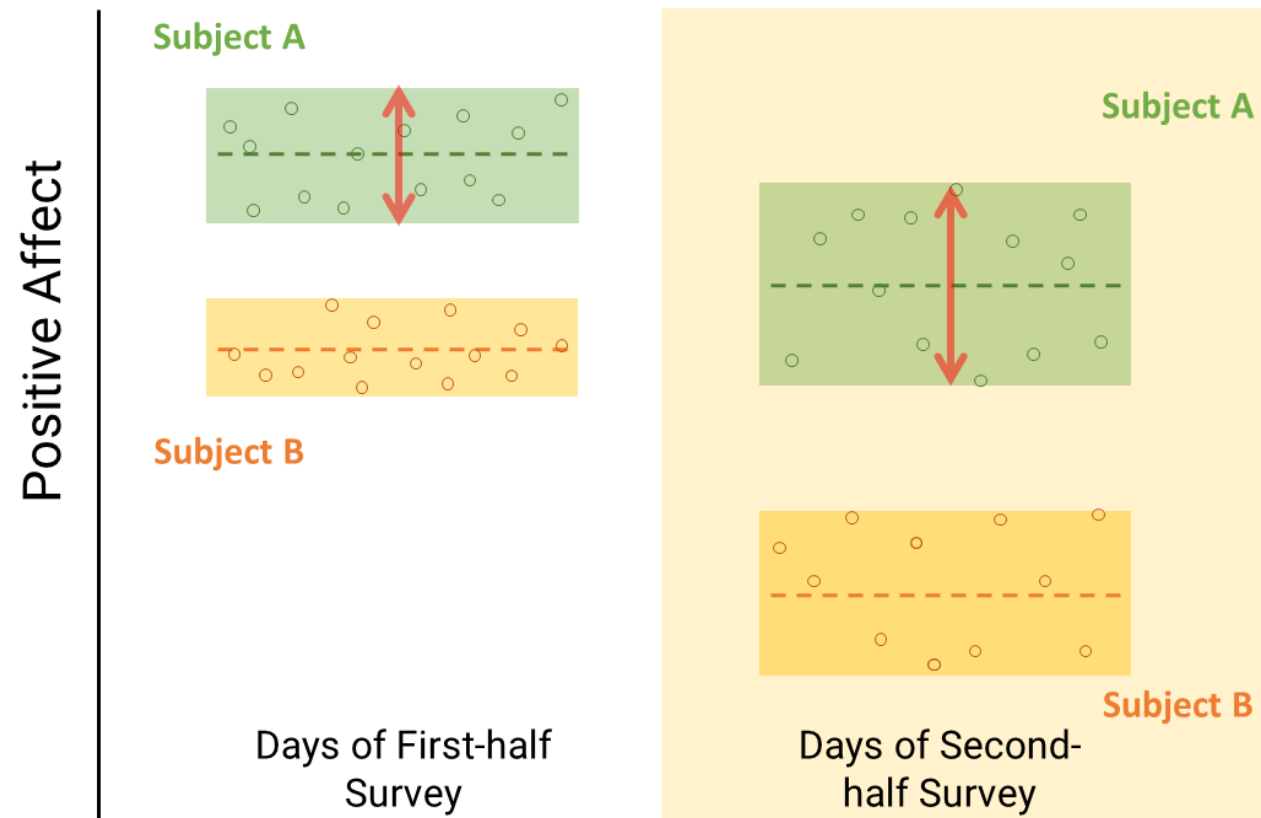




## Q1: Does the number of days in the study influence one's positive affect (PA)?

(WSV model) Do a subject's PA become more erratic as day passes?

- The effect of Day\_C is positive and significant, and it suggests subjects exhibit more erraticism (less consistency) over time.





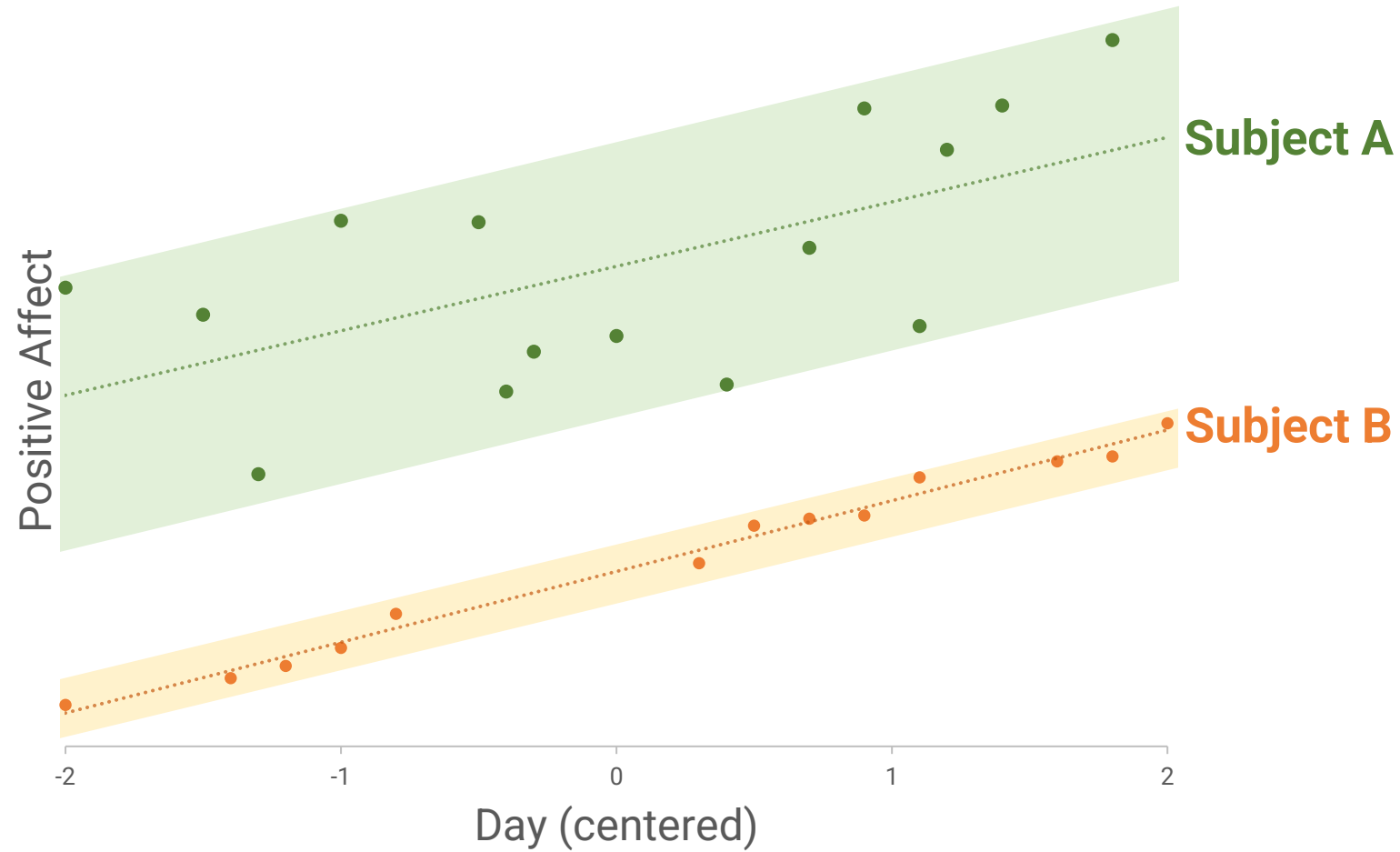
## Q1: Does the number of days in the study influence one's positive affect (PA)?

Variable	Estimate	AsymStdError	z-value	p-value
-----	-----	-----	-----	-----
BETA (regression coefficients)				
intercept	4.13072	0.14192	29.10578	0.00000
Day_C	-0.10383	0.01829	-5.67717	0.00000
ALPHA (BS variance parameters: log-linear model)				
intercept	0.33320	0.17051	1.95409	0.05069
Day_C	0.11952	0.01981	6.03246	0.00000
TAU (WS variance parameters: log-linear model)				
intercept	-0.09010	0.10237	-0.88010	0.37881
Day_C	0.13547	0.02616	5.17843	0.00000
Random scale standard deviation				
Std Dev	0.73612	0.07047	10.44548	0.00000
Random location (mean) effect on WS variance				
Loc Eff	-0.36756	0.09955	-3.69233	0.00022

**The standard deviation of the random scale effect** is estimated to be 0.73614, and this is a highly significant effect. Thus, subjects vary considerably in terms of how consistent/erratic they are in their PA reports.



Q1: Does the number of days in the study influence one's positive affect (PA)?



**The standard deviation of the random scale effect** is estimated to be 0.73614, and this is a highly significant effect. Thus, subjects vary considerably in terms of how consistent/erratic they are in their PA reports.



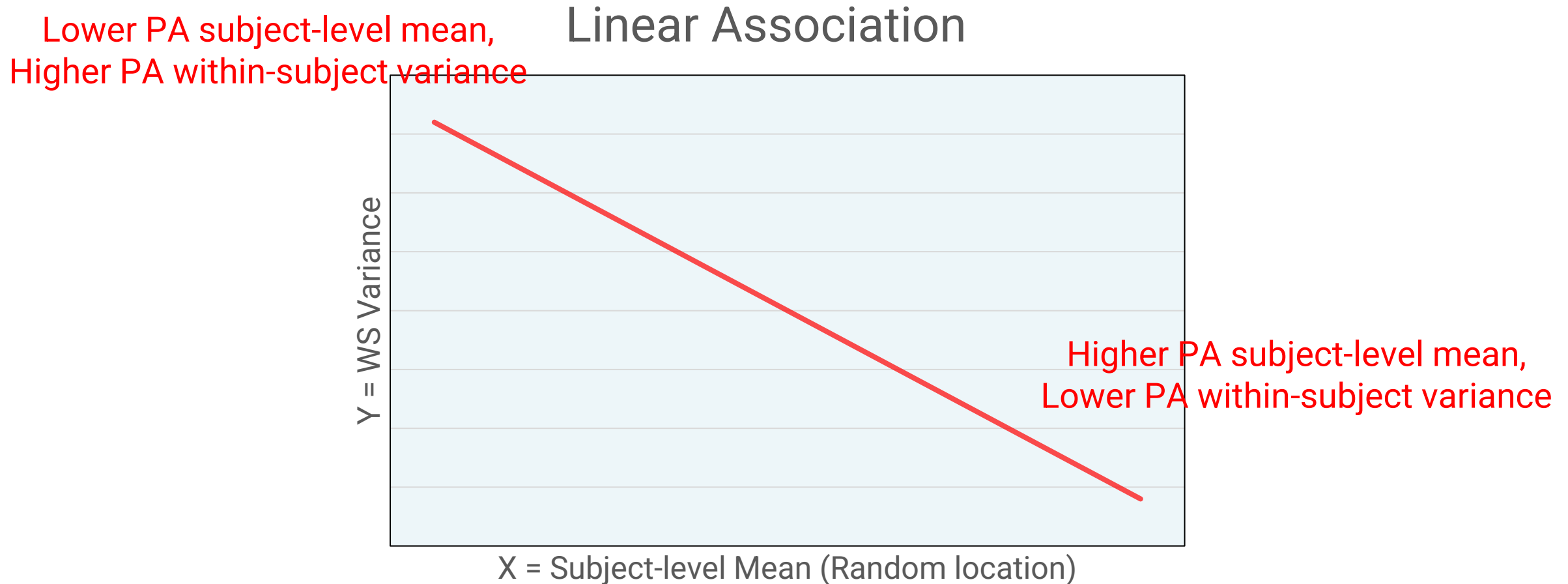
## Q1: Does the number of days in the study influence one's positive affect (PA)?

Variable	Estimate	AsymStdError	z-value	p-value
-----				
BETA (regression coefficients)				
intercept	4.13072	0.14192	29.10578	0.00000
Day_C	-0.10383	0.01829	-5.67717	0.00000
ALPHA (BS variance parameters: log-linear model)				
intercept	0.33320	0.17051	1.95409	0.05069
Day_C	0.11952	0.01981	6.03246	0.00000
TAU (WS variance parameters: log-linear model)				
intercept	-0.09010	0.10237	-0.88010	0.37881
Day_C	0.13547	0.02616	5.17843	0.00000
Random scale standard deviation				
Std Dev	0.73612	0.07047	10.44548	0.00000
Random location (mean) effect on WS variance				
Loc Eff	-0.36756	0.09955	-3.69233	0.00022

**The relationship between the random location and scale effects** is negative and significant indicating that subjects with higher average PA are also more consistent, and subjects with lower average PA are more erratic (also could be ceiling/cap effect).



Q1: Does the number of days in the study influence one's positive affect (PA)?



**The relationship between the random location and scale effects** is negative and significant indicating that subjects with higher average PA are also more consistent, and subjects with lower average PA are more erratic (also could be ceiling/cap effect).

## Research Question 2 (Q2)

Sleep  
Quality

Learning goal  
achievement

- Does day-to-day sleep quality influence one's day-to-day learning goal achievement (LGA)?

**{Stage 1 Model}**

- Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam?

**{Stage 2 Model}**



### **{Stage 1 Model}**

Does sleep quality (SQ) influence one's learning goal achievement (LGA)?

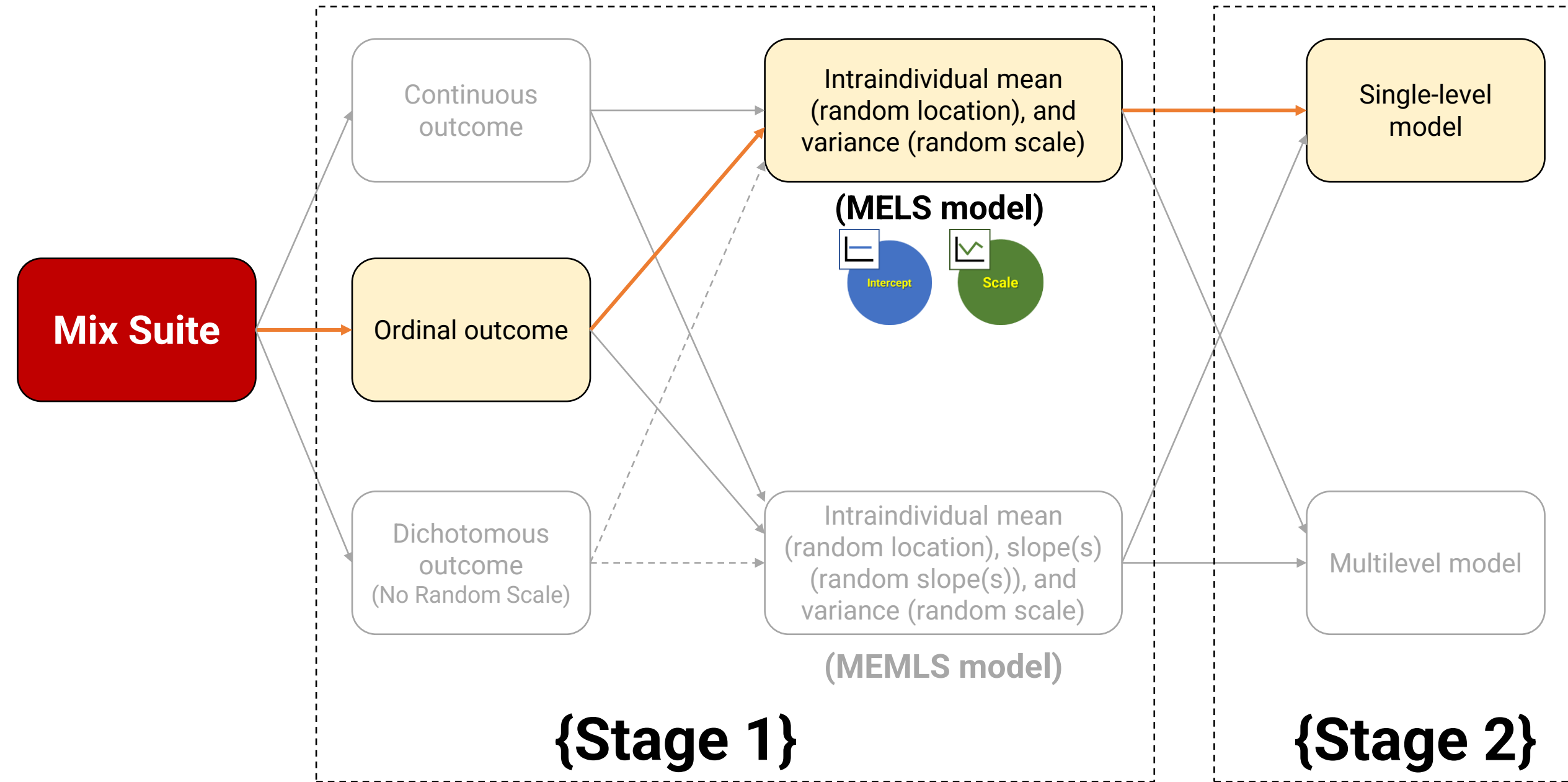
- (Mean model) Does a subject have a higher LGA on days with higher SQ?
- (BSV model) Does the sample become more homogeneous in LGA on days with higher SQ?
- (WSV model) Does a subject's LGA become more consistent on days with higher SQ?

### **{Stage 2 Model}**

Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam? (Subject-level Model)



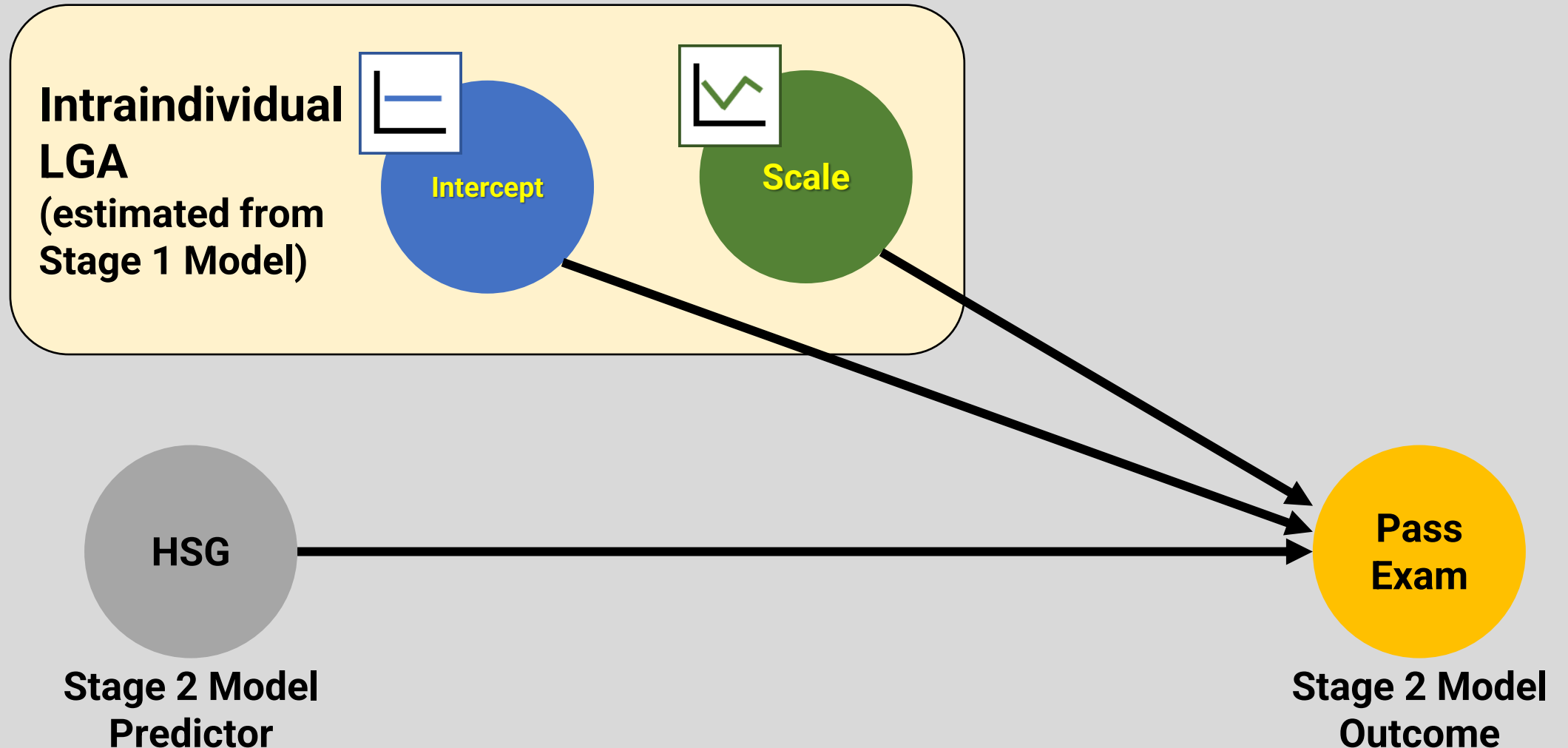
Q2: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam?



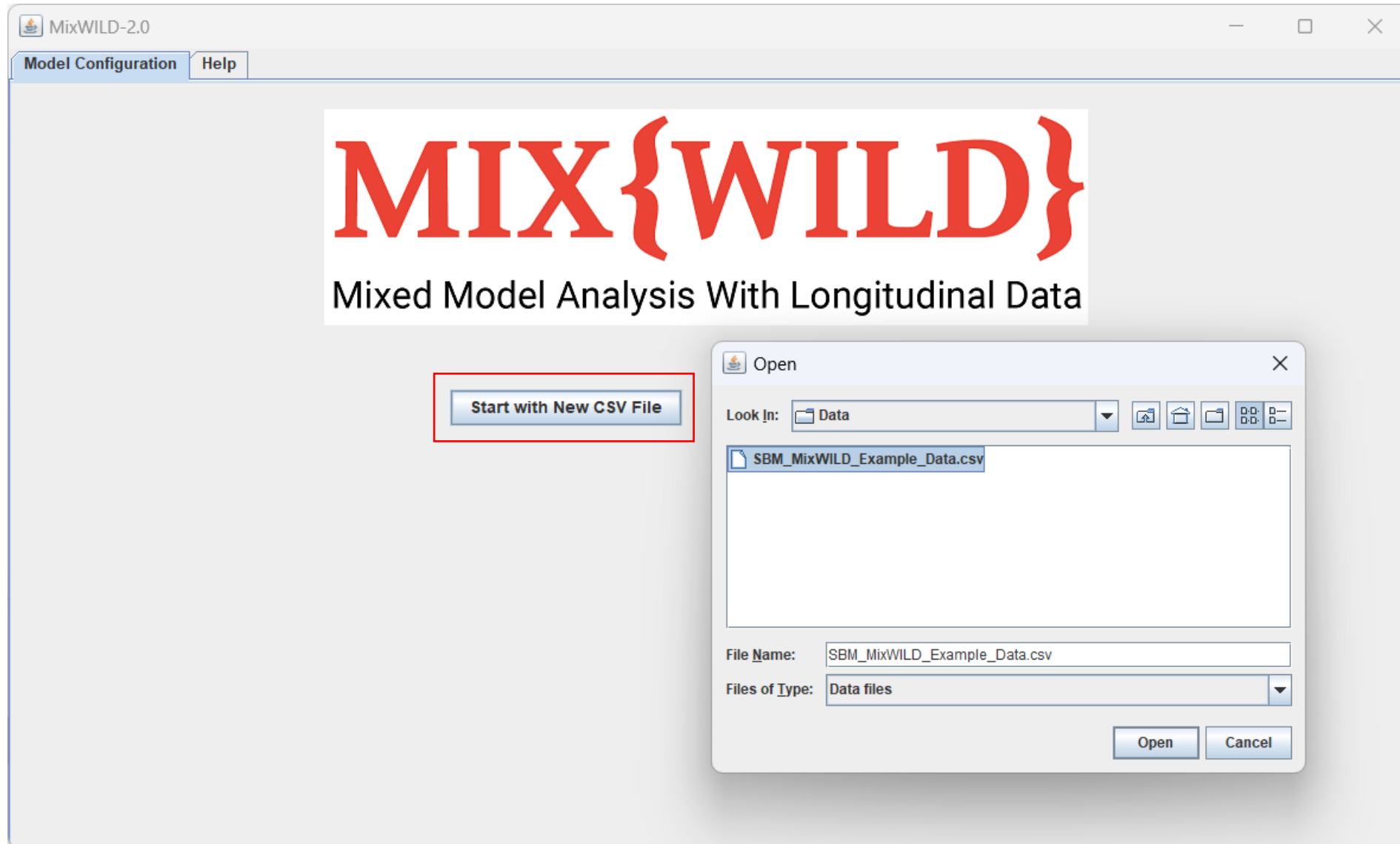


# MixWILD (Stage 2 Model)

- Use mean level of- and variance in LGA to predict exam results (Single-level model)



## Research Question 2 (Q2)



Start with “New CSV File” and locate the MixWILD example dataset



## Q2: (Stage 1) Does sleep quality (SQ) influence one's learning goal achievement (LGA)?

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path:  [Change Dataset](#)

Title (optional):

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as?

**Stage 1 Model**

Stage 1 outcome: ☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type: ☐ Probit ☒ Logistic

Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

Include Stage 2 model: ☒ Yes ☐ No

Include separate Stage 2 data file: ☐ Yes ☒ No

Stage 2 CSV file path:  [Import Dataset](#)

Stage 2 model type: ☒ Single level ☐ Multilevel

Stage 2 outcome: ☐ Continuous ☒ Dichotomous/Ordinal ☐ Count ☐ Multinomial

Set a seed for Stage 2 resampling (optional):

[Save Model](#) [Reset](#) [Continue](#)

Select Stage 1 outcome “Ordinal” (Outcome = “Learning goal achievement”)



## Q2: (Stage 1) Does sleep quality (SQ) influence one's learning goal achievement (LGA)?

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path:  [Change Dataset](#)

Title (optional):

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as?

**Stage 1 Model**

Stage 1 outcome: ☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type: ☐ Probit ☒ Logistic

Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

Include Stage 2 model: ☒ Yes ☐ No

Include separate Stage 2 data file: ☐ Yes ☒ No

Stage 2 CSV file path:  [Import Dataset](#)

Stage 2 model type: ☒ Single level ☐ Multilevel

Stage 2 outcome: ☐ Continuous ☒ Dichotomous/Ordinal ☐ Count ☐ Multinomial

Set a seed for Stage 2 resampling (optional):

[Save Model](#) [Reset](#) [Continue](#)

Specify random effects (Select “Intercept only” and include “Random scale”)



Q2: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam?

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path:  [Change Dataset](#)

Title (optional):

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as?

**Stage 1 Model**

Stage 1 outcome: ☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type: ☐ Probit ☒ Logistic

Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

Include Stage 2 model: ☒ Yes ☐ No

Include separate Stage 2 data file: ☐ Yes ☒ No

Stage 2 CSV file path:  [Import Dataset](#)

Stage 2 model type: ☒ Single level ☐ Multilevel

Stage 2 outcome: ☐ Continuous ☒ Dichotomous/Ordinal ☐ Count ☐ Multinomial

Set a seed for Stage 2 resampling (optional):

[Save Model](#) [Reset](#) [Continue](#)

Select Stage 2 Model and locate the Stage 2 data in your folder



Q2: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam?

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path:  **Change Dataset**

Title (optional):

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as?

**Stage 1 Model**

Stage 1 outcome: ☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type: ☐ Probit ☒ Logistic

Specify random location effects: ☒ Intercept only ☐ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

Include Stage 2 model: ☒ Yes ☐ No

Include separate Stage 2 data file: ☐ Yes ☒ No

Stage 2 CSV file path:  **Import Dataset**

Stage 2 model type: ☒ Single level ☐ Multilevel

Stage 2 outcome: ☐ Continuous ☒ Dichotomous/Ordinal ☐ Count ☐ Multinomial

Set a seed for Stage 2 resampling (optional):

**Save Model** **Reset** **Continue**

Specify Stage 2 model type ("Single level"); Outcome ("Exam" 1 = Pass; 0 = Fail)



## Q2: (Stage 1) Does sleep quality (SQ) influence one's learning goal achievement (LGA)?

MixWILD-2.0

Model Configuration Stage 1 Configuration Stage 2 Configuration View Data Help

**Selected Model Configuration**  
Stage 1 model: **Intercept Only**  
State 1 outcome: **Ordinal**

**ID Variable:**  
ID

**Stage 1 Outcome:**  
LGA

**Configure Stage 1 Regressors ...**

**Options ...**

**Specify the relationship between the mean and WS variance.**

☐ No Association  
☒ Linear Association  
☐ Quadratic Association

**Stage 1 Regressors**

	Mean	BS Variance	WS Variance
<b>Level-1</b>			
SQ	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Disaggregate?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
<b>Level-2</b>			

**Save Model** **Clear Stage 1** **Configure Stage 2**

Select “LGA” as the Stage 1 outcome and “SQ” as a time-varying predictor



## Q2: (Stage 1) Does sleep quality (SQ) influence one's learning goal achievement (LGA)?

MixWILD-2.0

Model Configuration Stage 1 Configuration Stage 2 Configuration View Data Help

**Selected Model Configuration**  
Stage 1 model: **Intercept Only**  
State 1 outcome: **Ordinal**

**ID Variable:**  
ID

**Stage 1 Outcome:**  
LGA

**Configure Stage 1 Regressors ...**  
**Options ...**

**Specify the relationship between the mean and WS variance.**

☐ No Association  
☒ Linear Association  
☐ Quadratic Association

**Stage 1 Regressors**

	Mean	BS Variance	WS Variance
<b>Level-1</b>			
SQ	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Disaggregate?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
<b>Level-2</b>			

**Save Model** **Clear Stage 1** **Configure Stage 2**

Specify the regressors in Stage 1 Models and click on “Configure Stage 2”





Q2: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam?

MixWILD-2.0

Model Configuration Stage 1 Configuration **Stage 2 Configuration** View Data Help

**Selected Model Configuration**  
Stage 1 model: **Intercept Only**  
Stage 1 outcome: **Ordinal**  
Stage 2 model type: **Single-level**  
Stage 2 outcome: **Dichot/Ord**  
Number of resamples (stage 2): **500**

**Stage 2 Outcome:**  
Exam

**Configure Stage 2 Regressors...**

**Check outcome categories**  
2 Categories:  
1) 0.0  
2) 1.0

**Stage 2 Interactions**

**Level-1**

Main Effects	Random Location	Random Scale	Location X Scale

**Level-2**

Main Effects	Random Location	Random Scale	Location X Scale
HSG_Rank	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>

☒ Suppress 2-way Location X Scale Interaction

**Save Model** **Clear Stage 2** **Run Stage 1 and 2**

Select “Exam” as Stage 2 Outcome; Check the outcome



Q2: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam?

MixWILD-2.0

Model Configuration Stage 1 Configuration Stage 2 Configuration View Data Help

**Selected Model Configuration**  
Stage 1 model: **Intercept Only**  
Stage 1 outcome: **Ordinal**  
Stage 2 model type: **Single-level**  
Stage 2 outcome: **Dichot/Ord**  
Number of resamples (stage 2): **500**

**Stage 2 Outcome:**  
Exam

**Configure Stage 2 Regressors...**

**Check outcome categories**  
2 Categories:  
1) 0.0  
2) 1.0

**Stage 2 Interactions**

**Add Stage 2 regressors**

**Variables**  
Day\_C  
Sex\_F  
Age\_C  
Sem  
HSG  
BDI  
SQ  
PhysAct  
PhysAct\_LN  
PA  
PA\_D  
PA\_Ord  
NA  
NA\_D  
NA\_Mean  
LGA

**Level 1 (Time Variant)**

**Level 2 (Time Invariant)**  
HSG\_Rank

Cancel Reset Submit

Click on “Configure Stage 2 Regressors to select “HSG\_Rank” as covariate”



Q2: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam?

MixWILD-2.0

Model Configuration Stage 1 Configuration **Stage 2 Configuration** View Data Help

**Selected Model Configuration**  
Stage 1 model: **Intercept Only**  
Stage 1 outcome: **Ordinal**  
Stage 2 model type: **Single-level**  
Stage 2 outcome: **Dichot/Ord**  
Number of resamples (stage 2): **500**

**Stage 2 Outcome:**  
Exam

**Configure Stage 2 Regressors...**

**Check outcome categories**  
2 Categories:  
1) 0.0  
2) 1.0

**Stage 2 Interactions**

**Main Effects Random Location Random Scale Location X Scale**

Level-1

Level-2

	Main Effects	Random Location	Random Scale	Location X Scale
HSG_Rank	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>

☒ Suppress 2-way Location X Scale Interaction

**Save Model Clear Stage 2 Run Stage 1 and 2**

Specify the regressor in the single-level model (Check “Main Effects” )



Q2: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam?

MixWILD-2.0

Model Configuration Stage 1 Configuration Stage 2 Configuration View Data Help

**Selected Model Configuration**  
Stage 1 model: Intercept Only  
Stage 1 outcome: Ordinal  
Stage 2 model type: Single-level  
Stage 2 outcome: Dichot/Ord  
Number of resamples (stage 2): 500

Stage 2 Outcome:  
Exam

Configure Stage 2 Regressors...

Check outcome categories

2 Categories:  
1) 0.0  
2) 1.0

**Stage 2 Interactions**

Main Effects Random Location Random Scale Location X Scale

Level-1

Level-2

	Main Effects	Random Location	Random Scale	Location X Scale
HSG_Rank	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>

☒ Suppress 2-way Location X Scale Interaction

Save Model Clear Stage 2 Run Stage 1 and 2

Please note random location and scale estimates are **default** regressors



Q2: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam?

MixWILD-2.0

Model Configuration Stage 1 Configuration **Stage 2 Configuration** View Data Help

**Selected Model Configuration**  
Stage 1 model: **Intercept Only**  
Stage 1 outcome: **Ordinal**  
Stage 2 model type: **Single-level**  
Stage 2 outcome: **Dichot/Ord**  
Number of resamples (stage 2): **500**

**Stage 2 Outcome:**  
Exam

**Configure Stage 2 Regressors...**

**Check outcome categories**  
2 Categories:  
1) 0.0  
2) 1.0

**Stage 2 Interactions**

**Main Effects Random Location Random Scale Location X Scale**

**Level-1**

**Level-2**

	Main Effects	Random Location	Random Scale	Location X Scale
HSG_Rank	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>

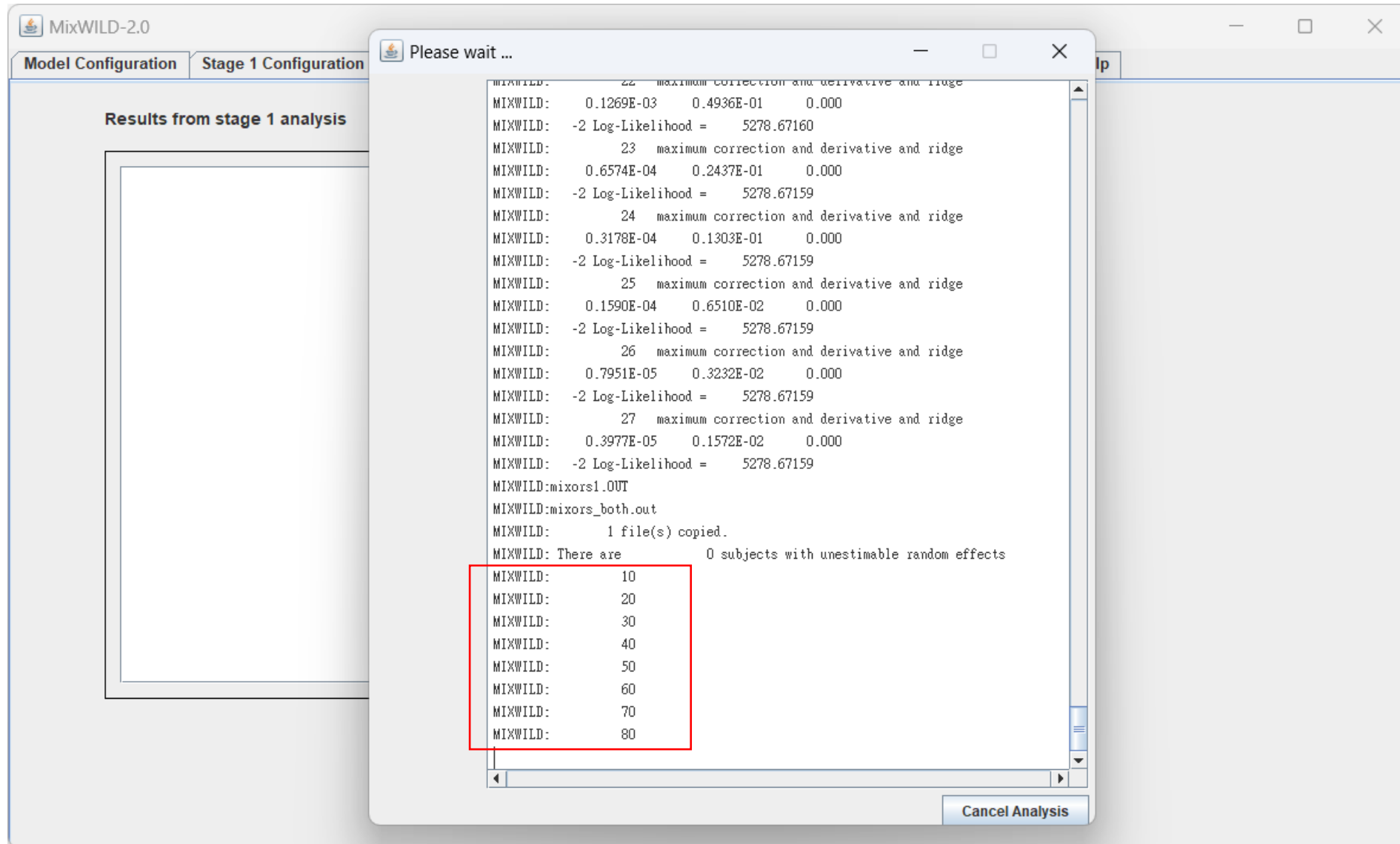
☒ Suppress 2-way Location X Scale Interaction

**Save Model Clear Stage 2 Run Stage 1 and 2**

Run Stage 1 and 2



Q2: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam?



After completing Stage 1, it performs 500 logistic regressions



Q2: (Stage 1) Does sleep quality (SQ) influence one’s learning goal achievement (LGA)?

Variable	Estimate	AsymStdError	z-value	p-value
-----				
BETA (regression coefficients)				
SQ	0.22590	0.05959	3.79082	0.00015
ALPHA (BS variance parameters: log-linear model)				
Intercept	0.25980	0.42487	0.61147	0.54089
SQ	0.00193	0.10377	0.01858	0.98518
TAU (WS variance parameters: log-linear model)				
SQ	-0.03627	0.03271	-1.10881	0.26751
Thresholds (for identification)				
1	-2.11209	0.30704	-6.87893	0.00000
2	-0.60166	0.22607	-2.66143	0.00778
3	1.11348	0.25694	4.33365	0.00001
4	3.41990	0.44740	7.64389	0.00000
Random location effects on WS variance (log-linear model)				
Linear	0.10151	0.05563	1.82469	0.06805
Random scale standard deviation				
Std Dev	0.31285	0.03770	8.29837	0.00000

**Mean model (BETA):** the slope of sleep quality (SQ) is positive and significant (beta = 0.22590;  $p < 0.001$ ). On average, a subject has a higher learning goal achievement (LGA) on days with higher SQ.



Q2: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam?

```
Number of replications =          500

-----
Final Results
-----

Average Log Likelihood      =      -42.094 (sd=      0.762)
Akaike's Information Criterion =      -46.094
Schwarz's Bayesian Criterion =      -50.647

==> multiplied by -2
Log Likelihood              =          84.188
Akaike's Information Criterion =          92.188
Schwarz's Bayesian Criterion =         101.295

Variable          Estimate      AsymStdError      z-value      p-value
-----
Intercept         -2.33488          0.91741      -2.54509      0.01092
HSG_Rank          0.17185          0.06764       2.54081      0.01106
Locat_1           0.65342          0.31175       2.09596      0.03609
Scale_1           0.01781          0.31544       0.05646      0.95498
```

Subject's high school grades (HSG\_Rank) is positive and significant. For every one unit increase in student's HSG\_Rank, the odds of being more likely to pass exam is multiplied 1.19 times ( $\exp(0.17185) = 1.19$ ).





Q2: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam?

```
Number of replications =          500

-----
Final Results
-----

Average Log Likelihood      =      -42.094 (sd=      0.762)
Akaike's Information Criterion =      -46.094
Schwarz's Bayesian Criterion =      -50.647

==> multiplied by -2
Log Likelihood              =          84.188
Akaike's Information Criterion =          92.188
Schwarz's Bayesian Criterion =         101.295

Variable          Estimate      AsymStdError      z-value      p-value
-----
Intercept         -2.33488          0.91741      -2.54509      0.01092
HSG Rank          0.17185          0.06764       2.54081      0.01106
Locat_1           0.65342          0.31175       2.09596      0.03609
Scale_1           0.01781          0.31544       0.05646      0.95498
```

**Subject-level random location effect:** the random location (subject's mean level) of learning goal achievement (LGA) is positive and significant. For every one unit increase in student's LGA, the odds of being more likely to pass exam is multiplied 1.92 times ( $\exp(0.65342) = 1.92$ ).



Q2: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA influence one's chance of passing exam?

```
Number of replications =          500

-----
Final Results
-----

Average Log Likelihood      =      -42.094 (sd=      0.762)
Akaike's Information Criterion =      -46.094
Schwarz's Bayesian Criterion =      -50.647

==> multiplied by -2
Log Likelihood              =      84.188
Akaike's Information Criterion =      92.188
Schwarz's Bayesian Criterion =     101.295

Variable          Estimate      AsymStdError      z-value      p-value
-----
Intercept         -2.33488          0.91741         -2.54509      0.01092
HSG_Rank           0.17185          0.06764          2.54081      0.01106
Locat_1           0.65342          0.31175          2.09596      0.03609
Scale_1           0.01781          0.31544          0.05646      0.95498
```

**Subject-level random scale effect:** the random scale (intraindividual variance) of learning goal achievement (LGA) is not significantly associated with the odds of being more likely to pass exam.



## Research Question 2b (Q2b) *Interaction Effects*

### {Stage 1 Model}

Does sleep quality (SQ) influence one's learning goal achievement (LGA)?

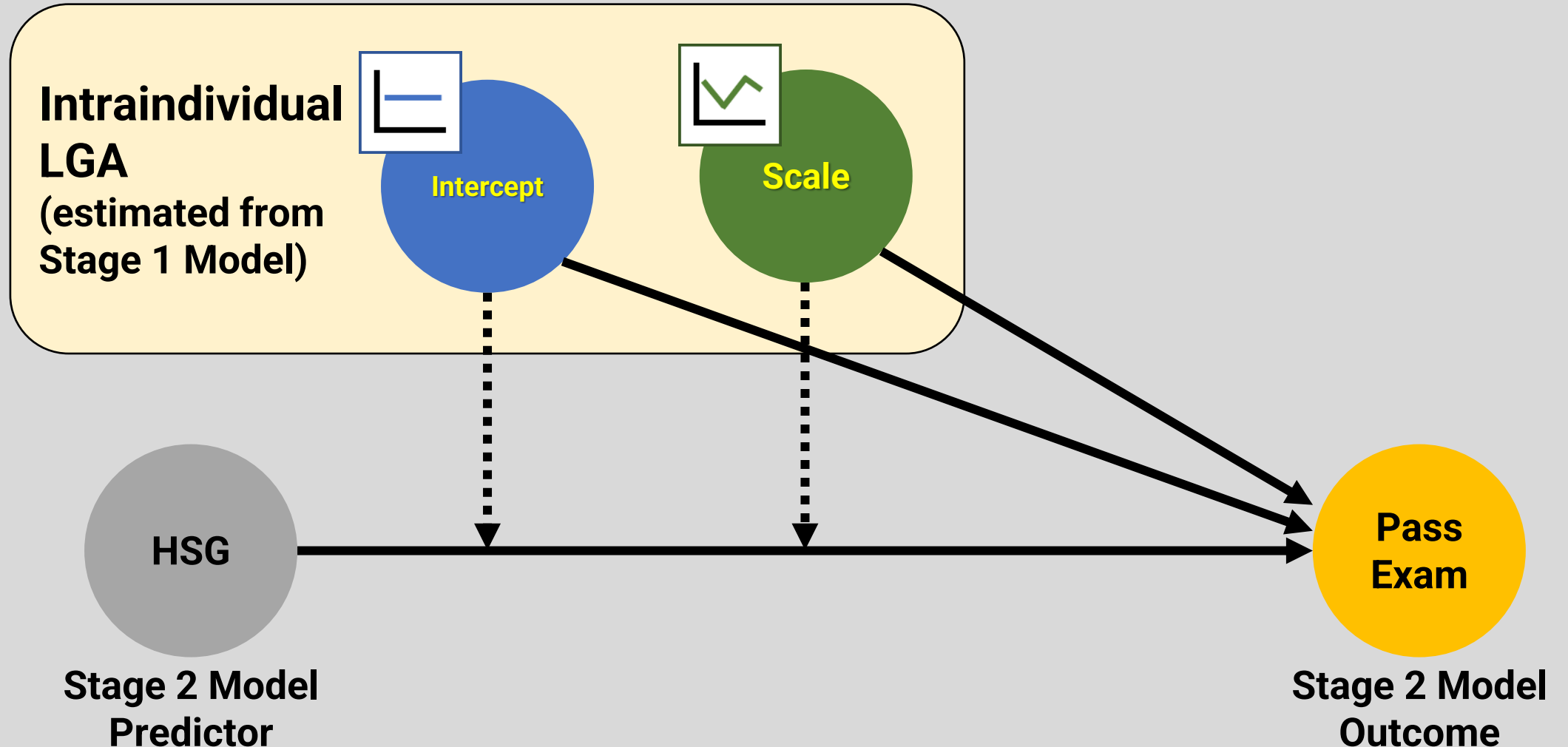
- (Mean model) Does a subject have a higher LGA on days with higher SQ?
- (BSV model) Does the sample become more homogeneous in LGA on days with higher SQ?
- (WSV model) Does a subject's LGA become more consistent on days with higher SQ?

### {Stage 2 Model}

Does subject's mean level of-, or intraindividual variance in LGA, or **interaction effects** influence one's chance of passing exam? (Subject-level Model)

# MixWILD (Stage 2 Model)

- Use mean level of-, variance in LGA, or the interaction effects to predict exam results





Q2b: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA, or interaction effects influence one's chance of passing exam?

MixWILD-2.0

Model Configuration Stage 1 Configuration Stage 2 Configuration View Data Help

**Selected Model Configuration**

Stage 1 model: **Intercept Only**  
Stage 1 outcome: **Ordinal**  
Stage 2 model type: **Single-level**  
Stage 2 outcome: **Dichot/Ord**  
Number of resamples (stage 2): **500**

**Stage 2 Outcome:**  
Exam

**Configure Stage 2 Regressors...**

**Check outcome categories**

2 Categories:  
1) 0.0  
2) 1.0

**Stage 2 Interactions**

**Main Effects Random Location Random Scale Location X Scale**

**Level-1**

**Level-2**

	Main Effects	Random Location	Random Scale	Location X Scale
HSG_Rank	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>

☐ Suppress 2-way Location X Scale Interaction

**Save Model Clear Stage 2 Run Stage 1 and 2**

Try interaction effects and uncheck “Suppress 2-way Location x Scale”



Q2b: (Stage 2) Does subject's mean level of- or intraindividual variance in LGA, or interaction effects influence one's chance of passing exam?

Number of replications = 500

-----  
Final Results  
-----

Average Log Likelihood = -41.213 (sd= 1.042)  
Akaike's Information Criterion = -48.213  
Schwarz's Bayesian Criterion = -56.181

==> multiplied by -2

Log Likelihood = 82.425  
Akaike's Information Criterion = 96.425  
Schwarz's Bayesian Criterion = 112.362

Variable	Estimate	AsymStdError	z-value	p-value
-----	-----	-----	-----	-----
Intercept	-2.49042	1.00474	-2.47867	0.01319
HSG_Rank	0.18321	0.07466	2.45385	0.01413
Locat_1	0.17765	1.11884	0.15878	0.87384
Locat_1*HSG_Rank	0.04033	0.08421	0.47885	0.63204
Scale_1	-0.16927	1.36956	-0.12359	0.90164
Scale_1*HSG_Rank	0.01372	0.10099	0.13582	0.89197
Locat_1*Scale	0.21538	0.35947	0.59917	0.54906



## Research Question 2c (Q2c) *Random Slope*

### {Stage 1 Model}

Does sleep quality (SQ) influence one's learning goal achievement (LGA)?

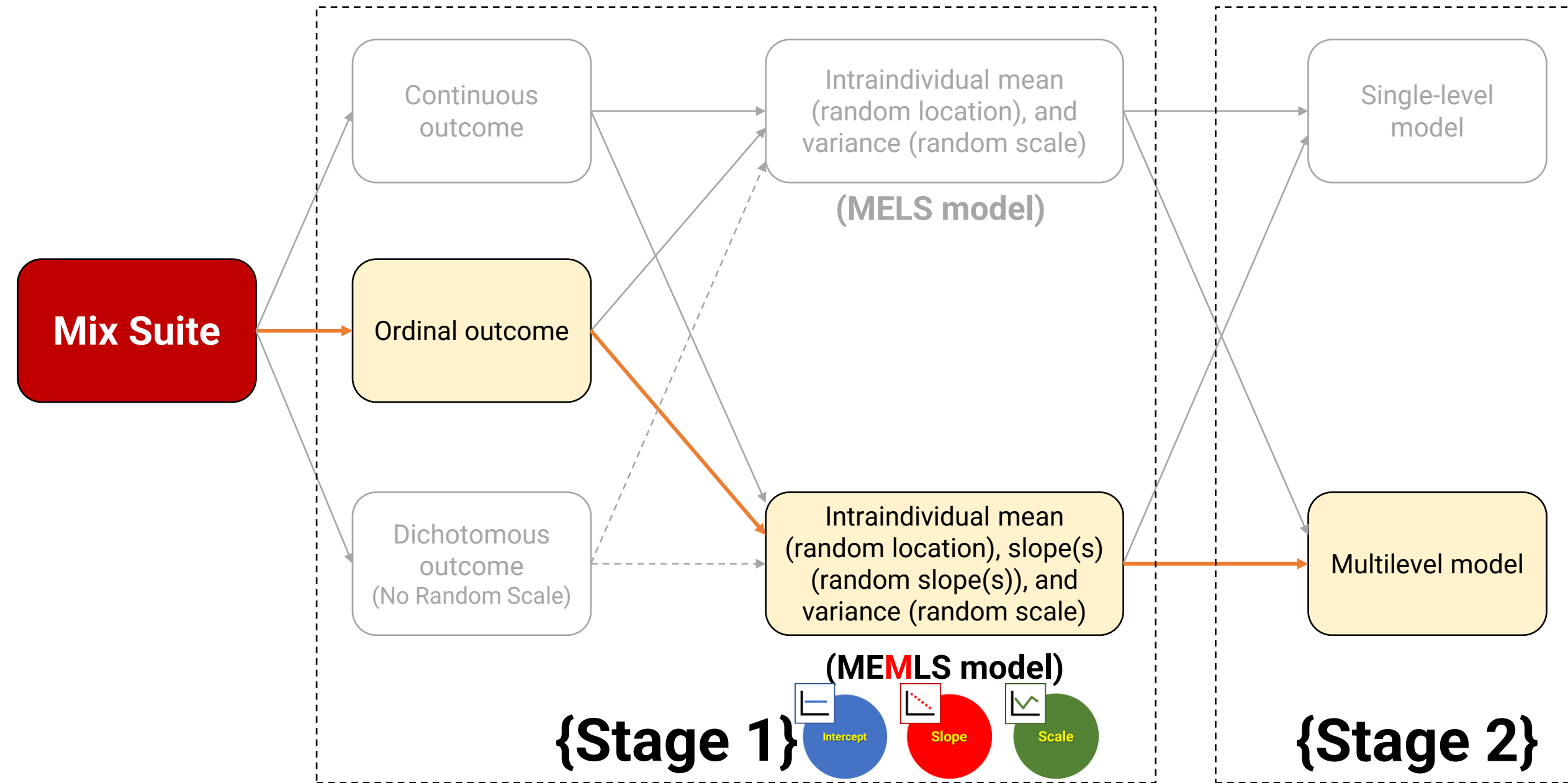
- (Mean model) Does a subject have a higher LGA on days with higher SQ?  
**Is the association between LGA and SQ different across subjects?**
- (WSV model) Does a subject's LGA become more consistent on days with higher SQ?

### {Stage 2 Model}

Does subject's mean level of-, or intraindividual variance in LGA, or **random-subject slope of SQ predicting LGA** influence one's daily positive affect? (Multilevel model, Level 1: Day; Level 2: Subject)



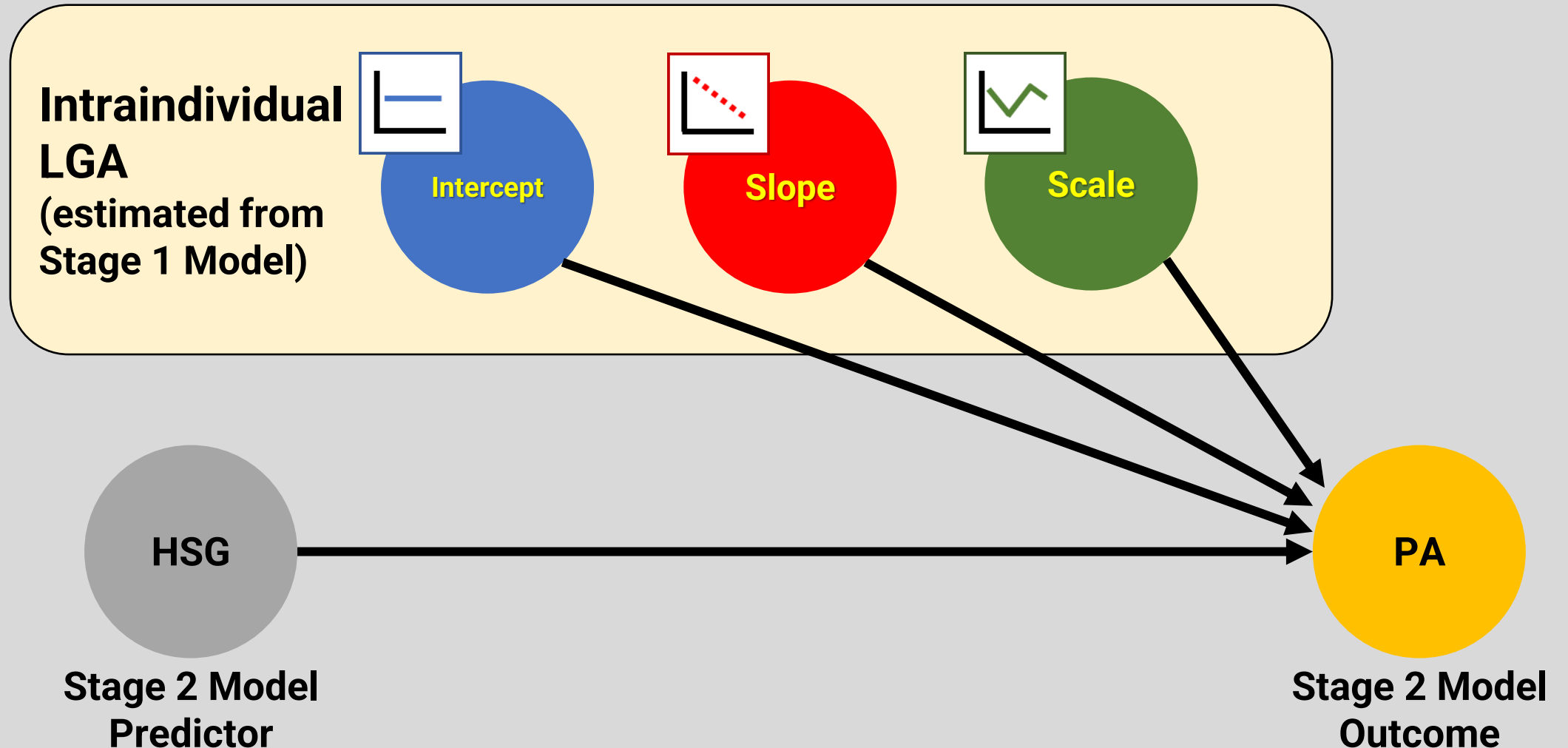
Q2c: (Stage 2) Does subject's mean level of-, or intraindividual variance in LGA, or random-subject slope of SQ predicting LGA influence one's daily positive affect?





# MixWILD (Stage 2 Model)

- Use mean level of-, slope of and variance in LGA to predict positive affect





Q2c: (Stage 1) Does sleep quality (SQ) influence one's learning goal achievement (LGA)?  
Is the association between LGA and SQ different across subjects?

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path:  [Change Dataset](#)

Title (optional):

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as?

**Stage 1 Model**

Stage 1 outcome: ☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type: ☐ Probit ☒ Logistic

Specify random location effects: ☐ Intercept only ☒ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

Include Stage 2 model: ☒ Yes ☐ No

Include separate Stage 2 data file: ☐ Yes ☒ No

Stage 2 CSV file path:  [Import Dataset](#)

Stage 2 model type: ☐ Single level ☒ Multilevel

Stage 2 outcome: ☒ Continuous ☐ Dichotomous/Ordinal ☐ Count ☐ Multinomial

Set a seed for Stage 2 resampling (optional):

[Save Model](#) [Reset](#) [Continue](#)

Specify random effects (Select “Intercept only and slope(s)”)



Q2c: (Stage 2) Does subject's mean level of-, or intraindividual variance in LGA, or random-subject slope of SQ predicting LGA influence one's daily positive affect?

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path:  **Change Dataset**

Title (optional):

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as?

**Stage 1 Model**

Stage 1 outcome: ☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type: ☐ Probit ☒ Logistic

Specify random location effects: ☐ Intercept only ☒ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

Include Stage 2 model: ☒ Yes ☐ No

Include separate Stage 2 data file: ☐ Yes ☒ No

Stage 2 CSV file path:  **Import Dataset**

Stage 2 model type: ☐ Single level ☒ Multilevel

Stage 2 outcome: ☒ Continuous ☐ Dichotomous/Ordinal ☐ Count ☐ Multinomial

Set a seed for Stage 2 resampling (optional):

**Save Model** **Reset** **Continue**

Select “Multilevel” since the new outcome, “positive affect”, is at level 1



Q2c: (Stage 2) Does subject's mean level of-, or intraindividual variance in LGA, or random-subject slope of SQ predicting LGA influence one's daily positive affect?

MixWILD-2.0

Model Configuration View Data Help

**Dataset**

CSV file path:  **Change Dataset**

Title (optional):

Does your data contain missing values? ☒ Yes ☐ No

What is your missing data coded as?

**Stage 1 Model**

Stage 1 outcome: ☐ Continuous ☐ Dichotomous ☒ Ordinal

Stage 1 regression type: ☐ Probit ☒ Logistic

Specify random location effects: ☐ Intercept only ☒ Intercept and slope(s)

Include estimates of random scale: ☒ Yes ☐ No

**Stage 2 Model**

Include Stage 2 model: ☒ Yes ☐ No

Include separate Stage 2 data file: ☐ Yes ☒ No

Stage 2 CSV file path:  **Import Dataset**

Stage 2 model type: ☐ Single level ☒ Multilevel

**Stage 2 outcome:** ☒ Continuous ☐ Dichotomous/Ordinal ☐ Count ☐ Multinomial

Set a seed for Stage 2 resampling (optional):

**Save Model** **Reset** **Continue**

Select “Continuous” for the new Stage 2 outcome, “positive affect (PA)”



Q2c: (Stage 1) Does sleep quality (SQ) influence one's learning goal achievement (LGA)?  
Is the association between LGA and SQ different across subjects?

MixWILD-2.0

Model Configuration Stage 1 Configuration Stage 2 Configuration View Data Help

**Selected Model Configuration**  
Stage 1 model: **Intercept + Slope(s)**  
State 1 outcome: **Ordinal**

**ID Variable:**  
ID

**Stage 1 Outcome:**  
LGA

**Configure Stage 1 Regressors ...**

**Options ...**

**Association of random location & scale?**  
☒ Yes  
☐ No

**Stage 1 Regressors**

	Mean	Random Slope	WS Variance
<b>Level-1</b>			
SQ	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Disaggregate?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
<b>Level-2</b>			
	<b>Mean</b>	<b>WS Variance</b>	

**Save Model** **Clear Stage 1** **Configure Stage 2**

Select “LGA” as the Stage 1 outcome and “SQ” as a time-varying predictor



Q2c: (Stage 1) Does sleep quality (SQ) influence one's learning goal achievement (LGA)?  
Is the association between LGA and SQ different across subjects?

MixWILD-2.0

Model Configuration Stage 1 Configuration Stage 2 Configuration View Data Help

**Selected Model Configuration**  
Stage 1 model: **Intercept + Slope(s)**  
State 1 outcome: **Ordinal**

**ID Variable:**  
ID

**Stage 1 Outcome:**  
LGA

**Configure Stage 1 Regressors ...**  
**Options ...**

**Association of random location & scale?**  
☒ Yes  
☐ No

**Stage 1 Regressors**

	Mean	Random Slope	WS Variance
<b>Level-1</b>			
SQ	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Disaggregate?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
<b>Level-2</b>			

**Mean** **WS Variance**

**Save Model** **Clear Stage 1** **Configure Stage 2**

Add a random slope of “SQ” in the Mean model



Q2c: (Stage 2) Does subject's mean level of-, or intraindividual variance in LGA, or random-subject slope of SQ predicting LGA influence one's daily positive affect?

MixWILD-2.0

Model Configuration Stage 1 Configuration Stage 2 Configuration View Data Help

**Selected Model Configuration**

Stage 1 model: **Intercept + Slope(s)**  
Stage 1 outcome: **Ordinal**  
Stage 2 model type: **Multi-level**  
Stage 2 outcome: **Continuous**  
Number of resamples (stage 2): **500**

Stage 2 Outcome:  
PA

Configure Stage 2 Regressors...

**Stage 2 Interactions**

Level-1

Main Effects	Random Location	Random Scale	Location X Scale

Level-2

Main Effects	Random Location	Random Scale	Location X Scale
HSG_Rank	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>

☒ Suppress 2-way Location X Scale Interaction

Save Model Clear Stage 2 Run Stage 1 and 2

Select "PA" as Stage 2 Outcome and add "HSG\_Rank" as Level-2 covariate



## Q2c: (Stage 1) Does sleep quality (SQ) influence one's learning goal achievement (LGA)? Is the association between LGA and SQ different across subject?

Variable	Estimate	AsymStdError	z-value	p-value
BETA (regression coefficients)				
SQ	0.19774	0.07007	2.82197	0.00477
Random (location) Effect Variances and Covariances				
Intercept	1.58607	0.27494	5.76887	0.00000
Covariance12	-0.06054	0.07820	-0.77419	0.43882
SQ	0.03154	0.05814	0.54247	0.58749
TAU (WS variance parameters: log-linear model)				
SQ	-0.03770	0.03321	-1.13524	0.25628
Thresholds (for identification)				
1	-2.19205	0.78148	-2.80500	0.00503
2	-0.68741	0.28521	-2.41015	0.01595
3	1.02594	0.10671	9.61429	0.00000
4	3.33576	0.47548	7.01558	0.00000
Random location effects on WS variance (log-linear model)				
Intercept	0.16154	0.07123	2.26769	0.02335
SQ	-0.10438	0.03679	-2.83755	0.00455
Random scale standard deviation				
Std Dev	0.26672	0.02501	10.66580	0.00000

**Mean model (BETA):** the slope of SQ is positive (beta = 0.19774; p = -.00477). For every one unit increase in student's SQ, the odds of being more likely to make higher learning goal achievement (versus “not at all”) is multiplied 1.22 times (i.e., increases 22%), holding constant all other variables.





Q2c: (Stage 1) Does sleep quality (SQ) influence one's learning goal achievement (LGA)?  
Is the association between LGA and SQ different across subject?

Variable	Estimate	AsymStdError	z-value	p-value
-----	-----	-----	-----	-----
BETA (regression coefficients)				
SQ	0.19774	0.07007	2.82197	0.00477
Random (location) Effect Variances and Covariances				
Intercept	1.58607	0.27494	5.76887	0.00000
Covariance12	-0.06054	0.07820	-0.77419	0.43882
SQ	0.03154	0.05814	0.54247	0.58749
TAU (WS variance parameters: log-linear model)				
SQ	-0.03770	0.03321	-1.13524	0.25628
Thresholds (for identification)				
1	-2.19205	0.78148	-2.80500	0.00503
2	-0.68741	0.28521	-2.41015	0.01595
3	1.02594	0.10671	9.61429	0.00000
4	3.33576	0.47548	7.01558	0.00000
Random location effects on WS variance (log-linear model)				
Intercept	0.16154	0.07123	2.26769	0.02335
SQ	-0.10438	0.03679	-2.83755	0.00455
Random scale standard deviation				
Std Dev	0.26672	0.02501	10.66580	0.00000

**Random (Location) Effect:** the subjects differ significantly between each other based on mean levels (random intercept) of LGA (estimate = 1.58607;  $p < 0.001$ ).



## Q2c: (Stage 1) Does sleep quality (SQ) influence one's learning goal achievement (LGA)? Is the association between LGA and SQ different across subject?

Variable	Estimate	AsymStdError	z-value	p-value
-----	-----	-----	-----	-----
BETA (regression coefficients)				
SQ	0.19774	0.07007	2.82197	0.00477
Random (location) Effect Variances and Covariances				
Intercept	1.58607	0.27494	5.76887	0.00000
Covariance12	-0.06054	0.07820	-0.77419	0.43882
SQ	0.03154	0.05814	0.54247	0.58749
TAU (WS variance parameters: log-linear model)				
SQ	-0.03770	0.03321	-1.13524	0.25628
Thresholds (for identification)				
1	-2.19205	0.78148	-2.80500	0.00503
2	-0.68741	0.28521	-2.41015	0.01595
3	1.02594	0.10671	9.61429	0.00000
4	3.33576	0.47548	7.01558	0.00000
Random location effects on WS variance (log-linear model)				
Intercept	0.16154	0.07123	2.26769	0.02335
SQ	-0.10438	0.03679	-2.83755	0.00455
Random scale standard deviation				
Std Dev	0.26672	0.02501	10.66580	0.00000

**Random (Location) Effect:** the random intercept and random slope were not statistically associated with each other (Covariance), indicating that there is no relationship between the mean levels of LGA and the coupling association of SQ and LGA (estimate = -0.06054; p = 0.43882).



Q2c: (Stage 1) Does sleep quality (SQ) influence one's learning goal achievement (LGA)?  
Is the association between LGA and SQ different across subject?

Variable	Estimate	AsymStdError	z-value	p-value
-----				
BETA (regression coefficients)				
SQ	0.19774	0.07007	2.82197	0.00477
Random (location) Effect Variances and Covariances				
Intercept	1.58607	0.27494	5.76887	0.00000
Covariance12	-0.06054	0.07820	-0.77419	0.43882
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2	-0.68741	0.28521	-2.41015	0.01595
3	1.02594	0.10671	9.61429	0.00000
4	3.33576	0.47548	7.01558	0.00000
Random location effects on WS variance (log-linear model)				
Intercept	0.16154	0.07123	2.26769	0.02335
SQ	-0.10438	0.03679	-2.83755	0.00455
Random scale standard deviation				
Std Dev	0.26672	0.02501	10.66580	0.00000

**Random (Location) Effect:** there is no statistical difference in their association (random slope) between SQ and LGA (estimate = 0.03154; p = 0.58749) across subjects.



## Q2c: (Stage 1) Does sleep quality (SQ) influence one's learning goal achievement (LGA)? Is the association between LGA and SQ different across subject?

Variable	Estimate	AsymStdError	z-value	p-value
-----				
BETA (regression coefficients)				
SQ	0.19774	0.07007	2.82197	0.00477
Random (location) Effect Variances and Covariances				
Intercept	1.58607	0.27494	5.76887	0.00000
Covariance12	-0.06054	0.07820	-0.77419	0.43882
SQ	0.03154	0.05814	0.54247	0.58749
TAU (WS variance parameters: log-linear model)				
SQ	-0.03770	0.03321	-1.13524	0.25628
Thresholds (for identification)				
1	-2.19205	0.78148	-2.80500	0.00503
2	-0.68741	0.28521	-2.41015	0.01595
3	1.02594	0.10671	9.61429	0.00000
4	3.33576	0.47548	7.01558	0.00000
Random location effects on WS variance (log-linear model)				
Intercept	0.16154	0.07123	2.26769	0.02335
SQ	-0.10438	0.03679	-2.83755	0.00455
Random scale standard deviation				
Std Dev	0.26672	0.02501	10.66580	0.00000

The relationship between **the random intercept** and **scale effects of LGA** is positive and significant, indicating that subjects with higher average are also more erratic.

The relationship between **the random slope of SQ predicting LGA** and **scale effect of LGA** is negative and significant, indicating that subjects with higher slope are less erratic.



Q2c: (Stage 1) Does sleep quality (SQ) influence one's learning goal achievement (LGA)?  
Is the association between LGA and SQ different across subject?

Variable	Estimate	AsymStdError	z-value	p-value
-----				
BETA (regression coefficients)				
SQ	0.19774	0.07007	2.82197	0.00477
Random (location) Effect Variances and Covariances				
Intercept	1.58607	0.27494	5.76887	0.00000
Covariance12	-0.06054	0.07820	-0.77419	0.43882
SQ	0.03154	0.05814	0.54247	0.58749
TAU (WS variance parameters: log-linear model)				
SQ	-0.03770	0.03321	-1.13524	0.25628
Thresholds (for identification)				
1	-2.19205	0.78148	-2.80500	0.00503
2	-0.68741	0.28521	-2.41015	0.01595
3	1.02594	0.10671	9.61429	0.00000
4	3.33576	0.47548	7.01558	0.00000
Random location effects on WS variance (log-linear model)				
Intercept	0.16154	0.07123	2.26769	0.02335
SQ	-0.10438	0.03679	-2.83755	0.00455
Random scale standard deviation				
Std Dev	0.26672	0.02501	10.66580	0.00000

**The standard deviation of the random scale effect** is estimated to be 0.26672, and this is a highly significant effect. Thus, subjects vary considerably in terms of how consistent/erratic they are in their LGA.



Q2c: (Stage 2) Does subject's mean level of-, or intraindividual variance in LGA, or random-subject slope of SQ predicting LGA influence one's daily positive affect?

----- Final Results -----				
Average Log Likelihood	=	-3311.270	(sd=	2.057)
Akaike's Information Criterion	=	-3317.270		
Schwarz's Bayesian Criterion	=	-3324.100		
==> multiplied by -2				
Log Likelihood	=	6622.540		
Akaike's Information Criterion	=	6634.540		
Schwarz's Bayesian Criterion	=	6648.200		
-----				
Variable	Estimate	AsymStdError	z-value	p-value
-----				
Intercept	4.33165	0.37209	11.64135	0.00000
HSG_Rank	-0.01435	0.02756	-0.52080	0.60250
Locat_1	0.47664	0.11872	4.01484	0.00006
Locat_2	0.57032	0.17890	3.18791	0.00143
Scale_1	0.13953	0.15861	0.87974	0.37900
-----				
Random.Int.Var	0.89552	0.16570	5.40441	0.00000
Residual.Varianc	1.21729	0.03818	31.88212	0.00000

**Subject-level random intercept effect:** the subject-level random intercept of learning goal achievement (Locat\_1) is positively associated with their reported positive affect (beta = 0.47664;  $p < 0.001$ ). It suggests that students with a higher subject-level mean of LGA, on average, have higher PA.





Q2c: (Stage 2) Does subject's mean level of-, or intraindividual variance in LGA, or random-subject slope of SQ predicting LGA influence one's daily positive affect?

----- Final Results -----				
Average Log Likelihood	=	-3311.270	(sd=	2.057)
Akaike's Information Criterion	=	-3317.270		
Schwarz's Bayesian Criterion	=	-3324.100		
==> multiplied by -2				
Log Likelihood	=	6622.540		
Akaike's Information Criterion	=	6634.540		
Schwarz's Bayesian Criterion	=	6648.200		
Variable	Estimate	AsymStdError	z-value	p-value
-----				
Intercept	4.33165	0.37209	11.64135	0.00000
HSG_Rank	-0.01435	0.02756	-0.52080	0.60250
Locat_1	0.47664	0.11872	4.01484	0.00006
Locat_2	0.57032	0.17890	3.18791	0.00143
Scale_1	0.13953	0.15861	0.87974	0.37900
Random.Int.Var	0.89552	0.16570	5.40441	0.00000
Residual.Varianc	1.21729	0.03818	31.88212	0.00000

**Subject-level random slope effect:** the subject-level random slope of sleep quality predicting learning goal achievement (Locat\_2) is positively associated with their reported positive affect (beta = 0.57032; p = 0.00143). It indicates that students with a higher subject-level slope of SQ predicting LGA (association between SQ and LGA) have higher PA.

# Troubleshooting





# Troubleshooting

## Limited variable number (Maximum ≈ 256)

Although there seems to be no limited of the sample size, the capacity of the maximum variable number could potentially be capped. Please keep the dataset as lite as possible and only include the variables that you will use in analysis.

	A	B	C	D	E	F	G	H	I	J	K	L	M	N
1	ID	Day	Sex	Age	Sem	SQ	PhysAct	PA	NA	LGA	Exam	HSG	BDI	
2	1	1	1	22	2	3	30	2.666667	3.666667	2	1	4.6	2	
3	1	2	1	22	2	3	60	3.333333	3	1	1	4.6	2	
4	1	3	1	22	2	3	360	2.666667	4	0	1	4.6	2	
5	1	4	1	22	2	3	780	1.333333	5	1	1	4.6	2	
6	1	6	1	22	2	3	210	3	4.666667	2	1	4.6	2	
7	1	8	1	22	2	3	310	4	3.666667	2	1	4.6	2	
8	1	9	1	22	2	3	90	3.333333	3.333333	0	1	4.6	2	
9	1	10	1	22	2	3	405	3.333333	2.666667	2	1	4.6	2	
10	1	11	1	22	2	3	405	2.333333	2.666667	2	1	4.6	2	
11	1	12	1	22	2	3	360	3	4	2	1	4.6	2	
12	1	13	1	22	2	3	270	1.666667	5.666667	1	1	4.6	2	
13	1	14	1	22	2	3	30	3.333333	2.333333	2	1	4.6	2	
14	1	15	1	22	2	3	540	2.666667	2	1	1	4.6	2	
15	1	17	1	22	2	2	405	4.333333	3.333333	1	1	4.6	2	
16	1	18	1	22	2	3	405	2.666667	3	1	1	4.6	2	
17	1	19	1	22	2	3	540	2.666667	3.333333	1	1	4.6	2	
18	1	21	1	22	2	3	60	3.666667	2.666667	2	1	4.6	2	
19	1	22	1	22	2	3	405	3.333333	3	1	1	4.6	2	
20	1	23	1	22	2	3	405	3.333333	3.333333	2	1	4.6	2	
21	1	24	1	22	2	2	60	1.666667	4.666667	2	1	4.6	2	
22	1	25	1	22	2	3	405	5	2.666667	2	1	4.6	2	
23	1	26	1	22	2	3	330	3.666667	2.666667	1	1	4.6	2	
24	1	27	1	22	2	3	435	2	3.666667	2	1	4.6	2	
25	1	29	1	22	2	3	525	3.666667	2.666667	2	1	4.6	2	

## Starting your model with a simplified variables/specification

Adding the random slope/scale effect may make your model overly complicated and generate some estimate difficulties. If you experience the model crash issue with no issue above, please try to start your model with simple settings. Also using Probit/Logistic model for dichotomous/ordinal outcome may increase estimate difficulties. Please check your outcomes before doing nonlinear models.

$$\begin{bmatrix} v_{0i} \\ v_{1i} \\ \omega_i \end{bmatrix} \sim N \left\{ \begin{bmatrix} 0 \\ 0 \\ 0 \end{bmatrix} \begin{bmatrix} \sigma_{v_0}^2 & \sigma_{v_0 v_1} & \sigma_{v_0 \omega} \\ \sigma_{v_0 v_1} & \sigma_{v_1}^2 & \sigma_{v_1 \omega} \\ \sigma_{v_0 \omega} & \sigma_{v_1 \omega} & \sigma_{\omega}^2 \end{bmatrix} \right\}$$

**Website:** <https://reach-lab.github.io/MixWildGUI/>

**User Guide:** [https://reach-lab.github.io/MixWildGUI/MixWild\\_User\\_Guide.pdf](https://reach-lab.github.io/MixWildGUI/MixWild_User_Guide.pdf)



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CHICAGO



Northeastern  
University



USC University of  
Southern California



# Troubleshooting



## Correct



## Error

### 1 No blank SPACE in dataset name / folder name

The dataset should be saved in a folder, and the folder name **CANNOT** have any **blank SPACES** ( ).

**Error: The data be loaded correctly, but the analysis will be shut down immediately when running the data.**

### 2 Variable names in the first row

The dataset should be saved as a **.CSV file** with variable names in the **first row**.

**Error: The software could not access to correct variable names/labels, and the first row data will be cut.**

### 3 No blank ( ) / Periods (.) / String ('miss') in data

Missing values should **NOT** be **blank ( )** or **periods (.)** or **string** in the dataset. All data should be coded as **numeric values** only, except for the first row (variable names).

**Error: The data cannot be read correctly, and it will end up to wrong estimates or model crash.**

### 4 Sorted by ID

Data should be sorted ascending or descending by ID number.

**Error: The data cannot be read correctly, and it will end up to get wrong estimates or model crash.**



C:\MyData\data\_1.csv

	A	B	C	D	E	F
1	ID	Sex_F	Age_C	Exam	HSG	PA_Mean
2	1	1	-1	1	4.6	3.11
3	2	1	-2	0	4.9	5.76
4	3	1	13	1	4.7	3.94
5	4	1	-2	0	4.3	2.9
6	5	1	-1	1	4.7	4.61

	A	B	C	D	E	F
1	ID	Sex_F	Age_C	Exam	HSG	PA_Mean
6	5	1	-1	1	4.7	4.61
7	6	1	-3	0	4.5	3.22
8	7	0	8	1	4.4	6.04
9	8	-99	-99	0	4.5	6.62
10	9	1	-3	1	5.3	5.03

	A	B	C	D	E	F
1	ID	Sex_F	Age_C	Exam	HSG	PA_Mean
2	1	1	-1	1	4.6	3.11
3	2	1	-2	0	4.9	5.76
4	3	1	13	1	4.7	3.94
5	4	1	-2	0	4.3	2.9
6	5	1	-1	1	4.7	4.61



C:\My Data\data 1.csv

	A	B	C	D	E	F
1	1	1	-1	1	4.6	3.11
2	2	1	-2	0	4.9	5.76
3	3	1	13	1	4.7	3.94
4	4	1	-2	0	4.3	2.9
5	5	1	-1	1	4.7	4.61
6	6	1	-3	0	4.5	3.22

	A	B	C	D	E	F
1	ID	Sex_F	Age_C	Exam	HSG	PA_Mean
6	5	1	-1	1	4.7	4.61
7	6	1	-3	0	4.5	3.22
8	7	0	8	1	4.4	6.04
9	8	.	.	0	4.5	6.62
10	9	1	-3	1	5.3	5.03

	A	B	C	D	E	F
1	ID	Sex_F	Age_C	Exam	HSG	PA_Mean
2	1	1	-1	1	4.6	3.11
3	2	1	-2	0	4.9	5.76
4	5	1	13	1	4.7	3.94
5	4	1	-2	0	4.3	2.9
6	3	1	-1	1	4.7	4.61





# Troubleshooting

## Quick Summary of the Optimal Options

(Please try to change these key parameters one by one in the model)

	Default	Optimization
<b>Quadrature Point</b>	11	15 - 25
<b>Maximum Iterations</b>	200	300 - 500
<b>Ridge</b>	0.1	0.15 – 0.25
<b>Discard Subjects with no Variance</b>	Uncheck	Check

### Quadrature Point

Usually, 11 points is sufficient, but if model convergence is not achieved, then increasing the points can sometimes help. So, for example, one might try 15, 21, or 25 points rather than the default of 11.

### Maximum Iterations

For example, beyond some number of iterations there are no practical gains. You can increase the number of iterations allowed to see if they will converge if the estimation doesn't converge within the default number. By default, the number of maximum iterations is 200.

### Ridge

The ridge increases the values of the diagonal elements of the 2nd derivative matrix by a factor of 1 multiplied by the ridge value. The reason that this is helpful is that this matrix must be inverted at each iteration of the solution, and inversion of this matrix becomes computationally difficult to the extent that the off-diagonal elements of this matrix get large, relative to the diagonal elements. Thus, in cases of non-convergence, one might try increasing the ridge value to 0.15, 0.20, or even 0.25. This will slow down the estimation, but in some cases can aid in model convergence.

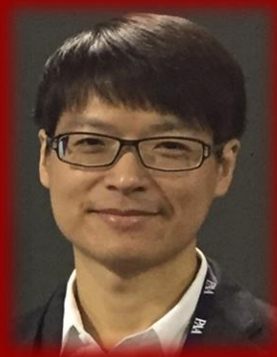
### Discard Subjects with no Variance

For such subjects with no variation on the outcome, the estimate of their random scale goes to negative infinity and can cause the program to fail to converge. In this case, the selection of the option can facilitate model convergence. **Please note selecting this option will remove these subjects from the stage 1 analysis.**



MixWILD website: <https://reach-lab.github.io/MixWildGUI/>

MixWILD GitHub: <https://github.com/reach-lab/MixWildGUI/discussions>



# Thank You!!!

Wei-Lin Wang

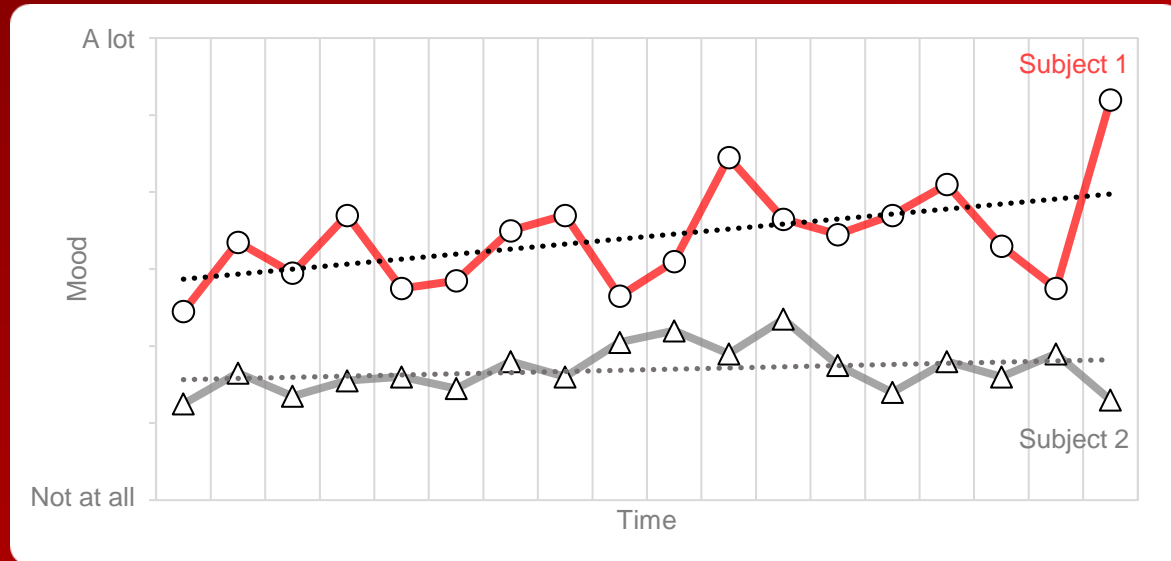
[weilinwa@usc.edu](mailto:weilinwa@usc.edu)

Department of Department of Population and Public Health Sciences

University of Southern California



# Harnessing the power of fluctuation: New horizons in modeling intraindividual variability with intensive longitudinal data



**Date:**

**Wednesday April 26, 2023**

**11:00 AM – 1:00 PM**



**Presenters:**

**Genevieve F. Dunton**

**Donald Hedeker**


**Wei-Lin Wang**

# Mini MixWILD Installation Handbook

**MIX{WILD}**  
Mixed Model Analysis With Longitudinal Data



# Download Software

1. Visit our website: <https://reach-lab.github.io/MixWildGUI/>
2. Click on **MacOS** or **Windows** to download the program.
  - **MacOS:** <https://github.com/reach-lab/MixWildGUI/releases/download/v2.0-stable/MixWILD-2.0.dmg>
  - **Windows:** <https://github.com/reach-lab/MixWildGUI/releases/download/v2.0-stable/MixWILD-2.0.exe>
3. Select your directory to save the program.
4. When finished downloading, double-click on the MixWILD icon  and follow the instructions to complete installation.

**MixWILD - Mixed models With Intensive Longitudinal Data**

Mix-WILD is a statistical software designed to perform multilevel modeling on intensive longitudinal experience sampling data.

[View the Project on GitHub](#)  
reach-lab/MixWildGUI

**MIX{WILD}**

Mixed Model Analysis With Intensive Longitudinal Data

**MixWild**

**What is this project about?**

MixWILD (Also Mixed model analysis with Intensive Longitudinal Data) is a desktop GUI-based application for examining the effects of variance and slope of time-varying variables in intensive longitudinal data, especially the ones collected using ecological momentary assessments.

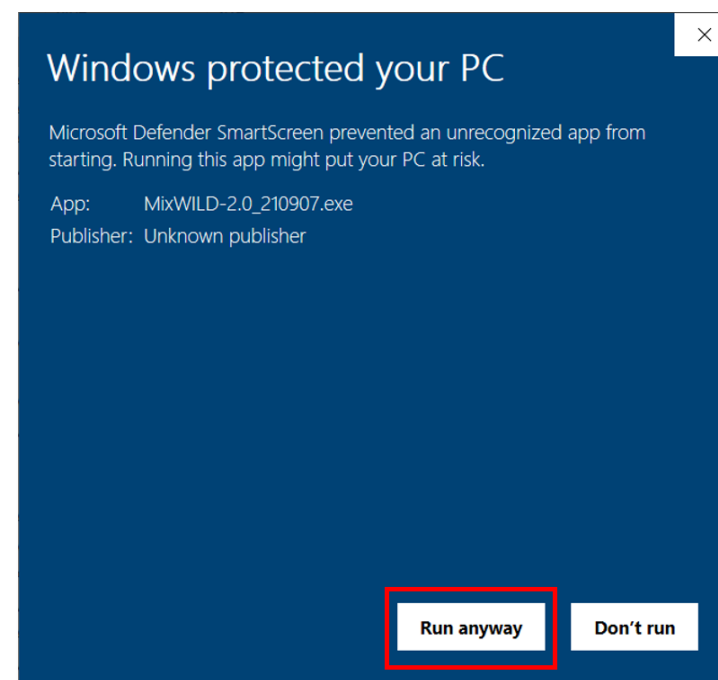
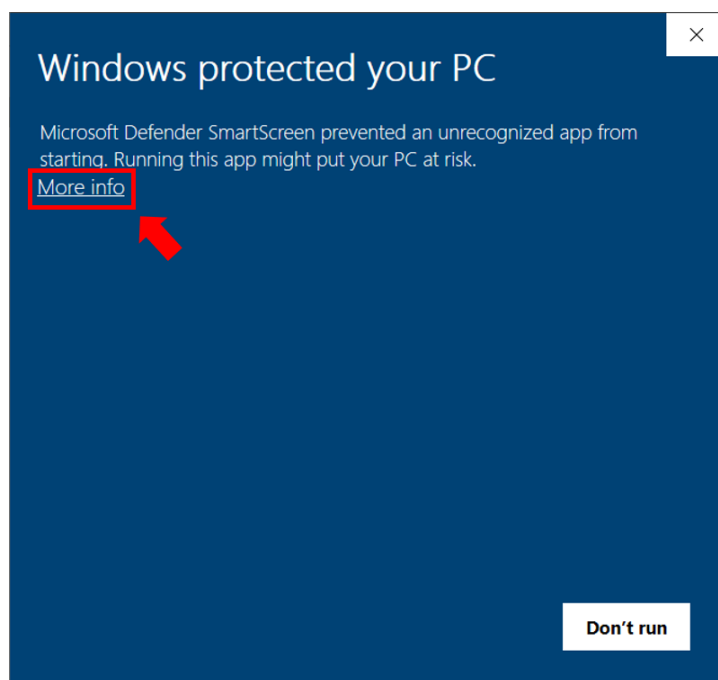




# Install Software

If this is your first time to install MixWILD, the Windows system may ask you to do some extra steps to successfully install the software.

1. Click on the MixWILD-2.0.exe, and click on **[More info]** to continue the process.

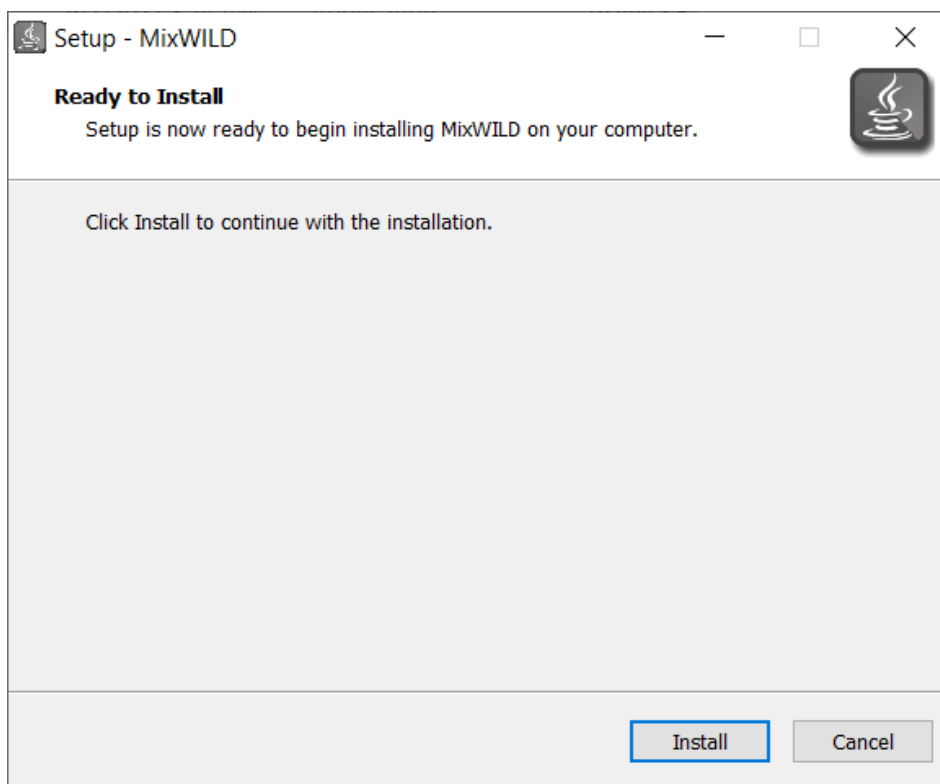



2. Click on **[Run anyway]**.



# Install Software

3. Click on **[Install]** to complete the installation.



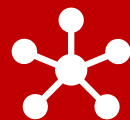
4. Well done!  You are all set! Please try MixWILD!





# Data for MixWILD

- The dataset should be saved in a folder, and the folder name **CANNOT** have any blank SPACES. (i.e., Please don't name your folder as "My Data" which will lead to an error. Please use **underscore** to replace space, the correct name should be "My\_Data").
- The dataset should be saved as a **.csv** file with variable names (no blank SPACES in variable names as well) in the **first row**.
- Data should be in the long format and sorted ascending or descending by **ID** number.
- Missing values should not be blank or periods (.) in the dataset and should be coded as **numeric values** only (i.e., "-999").



# Resources

MixWILD User's Guide:

<https://reach-lab.github.io/MixWildGUI/>

MixWILD Cheat Sheets:

[https://reach-lab.github.io/MixWildGUI/resources/cheat\\_sheets/MixWILD\\_UG\\_CS\\_220124.html](https://reach-lab.github.io/MixWildGUI/resources/cheat_sheets/MixWILD_UG_CS_220124.html)

MixWILD GitHub Discussion:

<https://github.com/reach-lab/MixWildGUI/discussions>

Introduction of Mixed-effects Location Scale Model Video:

<https://www.youtube.com/watch?v=wCEHuv9t1xw>

**MIX{WILD}**

Mixed Model Analysis With Longitudinal Data



Thank You Very Much! Look Forward to Meeting You  
on Wednesday April 26, 2023, in the SBM **MixWILD**  
Workshop!



**Wei-Lin Wang**

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