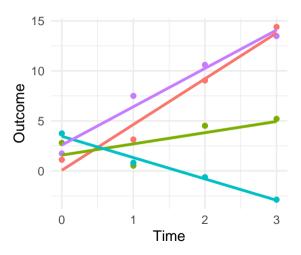
Linear Mixed Effects Models

What if we fit a different model for each individual?

We could find a unique β_{0i} and β_{1i} for each individual . . .



Problems with this...

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2. This procedure would ignore within-subject correlations.

Alternative idea...

Share parameters.

Use **individual-level** terms in addition to **population-level** terms which are shared across the population.

Parameter Sharing in Practice

Instead of β_{0i} and β_{1i} , we could break these down into $\beta_{0i} = \beta_0 + b_{0i}$ and $\beta_{1i} = \beta_1 + b_{1i}$.

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This way,

$$\begin{aligned} Y_{ij} &= \beta_{0i} + \beta_{1i}t_{ij} + \epsilon_{ij} \\ &= (\beta_0 + b_{0i}) + (\beta_1 + b_{1i})t_{ij} + \epsilon_{ij} \\ &= \underbrace{(\beta_0 + \beta_1t_{ij})}_{\text{Population Level}} + \underbrace{(b_{0i} + b_{1i}t_{ij})}_{\text{Individual Level}} + \epsilon_{ij}. \end{aligned}$$



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 - ▶ The measurement **variation** at time j: ϵ_{ij} .

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Can we do the same thing **here**?

Specification of a Linear Mixed Effects Model

For a continuous variate Y_{ij} , we take

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$$b_i \perp \epsilon_i$$
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 where $b_i \sim N(0, D)$, $\epsilon_i = (\epsilon_{i1}, \cdots, \epsilon_{ik_i})' \sim N(0, G_i)$, and with $b_i \perp \epsilon_i$.

Typically, we will set $G_i = \sigma^2 I$ to:

- 1. maintain the interpretation as sampling error; and
- 2. ensure **identifiability**.

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This is a specific form of a linear marginal model!

Specific Examples: Random Intercept and Random Slope Models

The Random Intercept Model

The most basic version of a mixed effects model takes $Z_{ij} = 1$, and as such, b_i is a scalar for each individual.

This is called the **random intercept model**.

The Random Intercept Model

We have that

$$Y_i = \beta_0 + b_{0i} + \widetilde{X}_i \beta + \epsilon_i,$$

with $b_{0i} \sim N(0, \sigma_b^2)$ and $\epsilon_i \sim N(0, G_i = \sigma^2 I)$.

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This gives

$$cov(Y_{ij}, Y_{il}) = \frac{\sigma_b^2}{\sigma_L^2 + \sigma^2}.$$

As a result, a random intercept model imposes the compound symmetry assumption!

The Random Intercept and Slope Model

If instead of just a random intercept, we also include a random time slope we get

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Here, D will be given by the variance of each b_{0i} and b_{1i} , as well as by the covariance between these terms.

The within-subject correlation will be time dependent in this model automatically!

Parameter Estimation and Hypothesis Testing

This is a Parametric Model

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This will give the (familiar) asymptotic results where

$$\widehat{\beta} \quad \stackrel{.}{\sim} \quad N\left(\beta, \left[\sum_{i=1}^n X_i' V_i^{-1}(\theta) X_i\right]^{-1}\right),$$

with
$$V_i(\theta) = \text{var}(Y_i) = Z_i D Z_i' + G_i$$
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- ▶ The parameters for the covariances, denoted θ , will have some **regularity concerns**.
- Can use the standard information criteria as well!



Estimation versus Prediction

We saw **estimation** of the parameters β , but the b_i are random! As a result, we must **predict** them.

The best[†] predictor for b_i will be $E[b_i|Y_i]$, a quantity that we call the **best linear unbiased prediction** (or BLUP).

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- Once estimated, we can estimate outcomes as

$$\widehat{Y}_i = X_i \widehat{\beta} + Z_i \widehat{b}_i = \dots = \widehat{G}_i \widehat{V}_i^{-1} X_i \beta + [I - \widehat{G}_i V_i^{-1}] Y_i.$$

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- This is a **weighted average** between the estimated population mean $(X_i\beta)$ and the individual observation Y_i .
- When G_i is large (more within-subject variation than between) there is more weight to the population average, and vice-versa.

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- ▶ We make normality assumptions, allowing for **standard asymptotic theory** when these are valid.
- ► Two basic, common models (random intercept and random intercept and slope) capture correlation structures that we have previously seen.
- ▶ We can use the **BLUP** to estimate individual effects, as-is necessary.

Drawbacks to Marginal Effects Models

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- ▶ They make distributional assumptions (GEEs did not).
- ▶ They may imply overly complex structures at the marginal level.