

# The patient-as-fixed-effect fallacy: Consequences for power and Type I errors

Examination for the course: ‘Open science and reproducible research’

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## 1 Introduction

In clinical psychology, and in many other fields, linear mixed-effects models (LMMs) have quickly risen in popularity during the 21st century (Gueorguieva and Krystal 2004). Their usage in clinical psychology is often motivated by LMMs ability to include participants with missing data (e.g., Kahn 2011). However, LMMs are highly sensitive to some types of model misspecification (Kwok, West, and Green 2007), and investigators are faced with many modelling choices, or researchers degrees of freedom (Wicherts et al. 2016). When analyzing patients that have been repeatedly measured during a treatment, investigators must decide whether to model subjects’ change over time as fixed or varying (random). The issues of treating and effect as constant for all individuals (fixed), or as varying between individuals (random), goes back a long time. For instance, in linguistics Clark (1973) coined the term “language-as-fixed-effect fallacy”, and Martindale (1978) soon followed by pointing out the unreasonable assumption that therapists have exactly the same success with their patients (“therapist-as-a-fixed-effect fallacy”). In an influential simulation study Barr et al. (2013) recommended to “keep it maximal”, i.e. include as many random effects as possible. Others (e.g., Matuschek et al. 2017) have noted that keeping it maximal might be too conservative, and that investigators need to balance the risk of Type I or II errors.

In this paper we will focus on one of the most basic decisions an analyst must make, when analyzing longitudinal treatment data—whether subjects’ trajectories over time should be seen as a fixed or random effect. Specifically, this paper focuses on the consequence of ignoring subject-specific varying slopes, on both Type I and II errors.

## 2 Methods

In typical multilevel notation, the simplest case of the two-level model is,

Level 1

$$Y_{ij} = \beta_{0j} + \beta_{1j}t_{ij} + R_{ij} \quad (1)$$

Level 2

$$\beta_{0j} = \gamma_{00} + \gamma_{01}TX_j + U_{0j} \quad (2)$$

$$\beta_{1j} = \gamma_{10} + \gamma_{11}TX_j + U_{1j} \quad (3)$$

$$(4)$$

$$\text{with } \begin{pmatrix} U_{0j} \\ U_{1j} \end{pmatrix} \sim \mathcal{N} \left( \begin{pmatrix} 0 \\ 0 \end{pmatrix}, \begin{pmatrix} \sigma_{u_0}^2 & \sigma_{u_{01}} \\ \sigma_{u_{01}} & \sigma_{u_1}^2 \end{pmatrix} \right), \text{ and } R_{ij} \sim \mathcal{N}(0, \sigma^2) \quad (5)$$

The parameter of interest is  $\gamma_{11}$ , which is the mean difference in change between the two groups. The aim of this paper is to investigate if accounting for subject-specific slopes ( $\sigma_{u_1}^2 > 0$ ) is important, both when planning the study, or when analyzing the outcome.

## 2.1 Simulation

To investigate the impact of wrongly omitting a random slope on the risk of committing a Type I error, a Monte Carlo simulation was performed. A parallel group RCT with 11 weekly time points was assumed, with 50 participants in each treatment group. We also assumed that at baseline there was an equal amount of variance between and within subjects, which would translate to an intraclass correlation of 0.5, if there was no variation between subjects in change over time,  $\sigma_{u_1}^2 = 0$ . The simulation consisted of comparing 5 different amounts of random slope variance,  $\sigma_{u_1}^2/\sigma^2 = \{0, 0.01, 0.02, 0.03, 0.04\}$ . We write the slope variance as a fraction of the error variance, since it's the ratio that matters, not the absolute value of,  $\sigma_{u_1}^2$ .<sup>1</sup>

Power was calculated for the same model, assuming a Cohen's  $d$  of 0.5 (standardized using the pretest standard deviation). Simulations and power calculations were done in R (version 3.4.3; R Core Team, 2017), using *powerlmm* (version 0.2.0; Magnusson 2018). LMMs were fit with *lme4* (version 1.1-15; Bates et al. 2015), using restricted maximum likelihood estimation. For each model 5000 data sets were generated, resulting in a 95 % Monte Carlo CI of 0.044–0.056, for a nominal  $\alpha$  of 0.05. Simulation code and additional material is available at: <https://github.com/rpsychologist/open-science-exam>.

## 3 Results

Figure 1 shows that.

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<sup>1</sup>Here I use the famous proof: "proof is left as an exercise to the reader", or "It can easily be shown that..." ;)

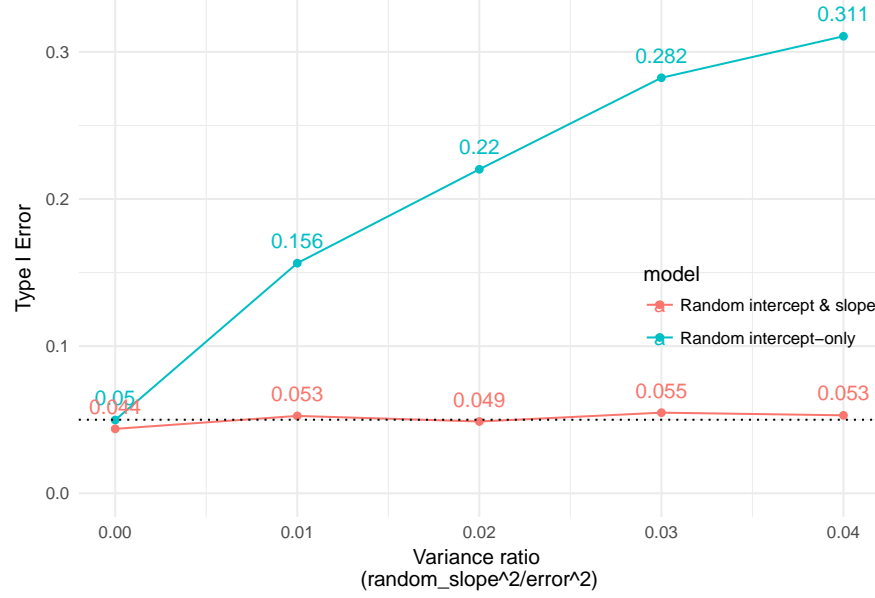


Figure 1: Type I errors for models with different amounts of true slope variance. Nominal levels is 0.05.

### 3.1 Power curves

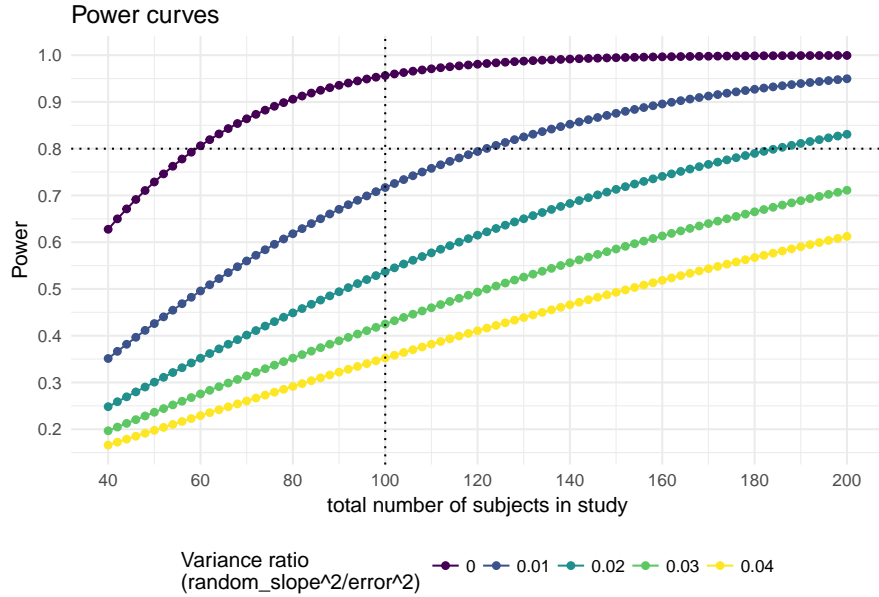


Figure 2: Power curves. Cohen's  $d = 0.5$ , 11 time points, and ICC at pretest = 0.5.

## 4 Discussion

Even if some model selection could be used, to select the most parsimonious model, investigators must still decide whether the test should be sensitive to detect relevant differences even if random slopes are included. Our recommendation is that researchers assume participants will change differently during a treatment trial, and include enough participants so that the statistical tests will not miss clinically relevant effects.

even under moderate to large amounts of heterogeneity in change. To facilitate this we encourage authors to actually report these variance components, so that some kind of reference point is available.

Lastly, our simulation showed that ignoring even small amounts of slopes variance can substantially inflate the Type I errors. Hence, we think it is reasonable that investigators who consider a random intercept-only model a reasonable choice, provide sufficient evidence for this assumption.

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