

A Comparative Study of Fixed Effects Models and Random Intercept/Slope Models as a Special Case of Linear Mixed Models for Repeated Measurements

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Summary. Any dataset in which subjects are measured repeatedly over time or space can be described as *repeated measurements data*. A *linear mixed model (LMM)* is a powerful method for analyzing repeated measurements data. It is made up of two components. The first component consists of a regression model for the average response over time and the effects of covariates on this average response. The second component provides a model for the pattern of covariances or correlations between the repeated measurements. In this study, a comparative evaluation of fixed effects models with random intercept models and random intercept and slope models as a special case of random effects models from linear mixed models are taken into consideration and the superiority of random intercept and slope models allow to modeling possible heterogeneity in intercepts and in slopes of the individual's own regression line for repeated measurements data is emphasized

Key words: Repeated measurement, linear mixed model, fixed effects model, random effects model, random intercept model, random intercept and slope model, compound symmetry pattern, Mauchly's sphericity test.

1. Introduction

The defining feature of a longitudinal study design is that measurements of the same individuals are taken repeatedly through time allowing the direct study of change over time. The primary goal of a longitudinal study is to characterize the change in response over time and the factors that influence change (Fitzmaurice et al., 2004). Analyzing such a repeated measurements data on individuals requires recognizing and estimating variability both between and within individuals. The linear mixed model approach to repeated measurements allows

explicit modeling and analysis of this variation between-subjects and within-subjects factors (Davis, 2002). A factor consisting of repeated measurements on the same subjects or experimental units, under different conditions is commonly called a “*within-subjects*” factor. A “*between-subjects*” factor is one in which each level of the factor contains different experimental units (İyit et al., 2006). Linear mixed models (LMMs) provide a tool for analyzing repeated measurements data by taking into consideration these two types of variability as well as the linear relationship between the explanatory variables and the response variable (McCulloch and Searle, 2001).

The theoretical base of linear mixed models is well-established and the methodology has applications in many areas not only involving repeated measurements. McLean et al. (1991) provide a general introduction to linear mixed models and Ware (1985) gives an overview of their application to the analysis of repeated measurements. Raudenbush and Bryk (2002) investigate random intercept models and random intercept and slope models in details and Bryk and Driscoll (1988) use these two model types to examine how characteristics of school organization are related to teachers’ sense of efficacy in their work in the field of education.

In this study a comparative evaluation of fixed effects models with random intercept models and random intercept and slope models as a special case of random effects models from linear mixed models are taken into consideration and the superiority of random intercept and slope models allows to modeling possible heterogeneity in intercepts and in slopes of the individual’s own regression line for repeated measurements data is emphasized.

This paper is organized as follows; In Section 2 a general introduction to linear mixed models is given and then fixed effects models and random intercept/slope models as a special case of random effects models from linear mixed models are introduced also the information criteria in model selection of fixed and random effects models are given. In Section 3 an application of a statistical analysis of repeated measurements data with both fixed effects models and random effects models is considered. Results and discussion also conclusion parts obtained from the analysis with these procedures are given in Section 4 and Section 5 respectively.

2. Linear Mixed Models for Repeated Measurements

A model where we have both *fixed effects* and *random effects* is called *mixed model* because of the mixture of different types of these effects on response variable. *Fixed effects* are the effects attributable to a finite set of levels of a factor on response variable representing all possible levels of the variable in which inferences are to be made. On the other hand *random effects* are the ones attributable to an infinite set of levels of a factor on response variable of which only a random sample of potential levels of the factor is taken to draw inferences for the complete population of levels (Searle et al., 1992). In mixed models *fixed effects* are used to explain the expected value of the observations and *random effects* to explain the variance-covariance structure of the dependent variable

(Hamer and Simpson, 1989). If the relationship between the observations of the response variable and these effects is linear, then the model is called *linear mixed model* (Davis, 2002).

Laird and Ware (1982) consider the LMM in matrix notation;

$$(1) \quad \underline{Y} = \mathbf{X}\underline{\beta} + \mathbf{Z}\underline{u} + \underline{\varepsilon}$$

where $\underline{Y}_{n \times 1}$ is the vector of response variables, $\mathbf{X}_{n \times p}$ is the known design matrix for fixed effects parameter vector, $\underline{\beta}_{p \times 1}$ is the unknown fixed effects parameter vector, $\mathbf{Z}_{n \times q}$ is the known design matrix for random effects vector, $\underline{u}_{q \times 1}$ is the unobserved random effects vector for $q = \sum_{i=1}^r q_i$ having $(\underline{u}_i)_{q_i \times 1}$ sub-vectors for $i = 1, 2, \dots, r$ random effects included in the model such as;

$$(2) \quad \underline{u} = [\underline{u}_1 \quad \underline{u}_2 \quad \cdots \quad \underline{u}_r]'$$

and corresponding $(\mathbf{Z}_i)_{n \times q_i}$ sub-matrices for $i = 1, 2, \dots, r$ random effects given by Eq.(2) included in the model such as;

$$(3) \quad \mathbf{Z} = [\mathbf{Z}_1 \quad \mathbf{Z}_2 \quad \cdots \quad \mathbf{Z}_r]$$

and $\underline{\varepsilon}_{n \times 1}$ is the unobserved vector of the independent and identically distributed Gaussian random error terms (McLean et al.,1991). The mean vector and the variance-covariance matrix for the components of LMM are as follows;

$$(4) \quad E(\underline{u}) = \underline{0} \text{ and } cov(\underline{u}) = E(\underline{u}\underline{u}') = \mathbf{D} \Rightarrow \underline{u} \sim N(\underline{0}, \mathbf{D})$$

$$(5) \quad E(\underline{\varepsilon}) = \underline{0} \text{ and } cov(\underline{\varepsilon}) = E(\underline{\varepsilon}\underline{\varepsilon}') = \mathbf{R} = \sigma_\varepsilon^2 \mathbf{I}_N \Rightarrow \underline{\varepsilon} \sim N(\underline{0}, \mathbf{R})$$

$$(6) \quad cov(\underline{u}, \underline{\varepsilon}') = cov(\underline{\varepsilon}, \underline{u}') = \mathbf{0}$$

Then

$$(7) \quad E(\underline{Y}) = \mathbf{X}\underline{\beta} \text{ and } cov(\underline{Y}) = \mathbf{V} = \mathbf{Z}\mathbf{D}\mathbf{Z}' + \mathbf{R} \Rightarrow \underline{Y} \sim N(\mathbf{X}\underline{\beta}, \mathbf{Z}\mathbf{D}\mathbf{Z}' + \mathbf{R})$$

In repeated measurements designs $\mathbf{Z}\mathbf{D}\mathbf{Z}'$ represents the between-subject portion of the covariance structure and \mathbf{R} represents the within-subject portion of the covariance structure.

Parameters in LMMs are estimated by *maximum likelihood (ML)* or by a technique known as *restricted maximum likelihood (REML)*. For $\underline{Y}_{n \times 1}$; the vector of

response variables having the assumption given by Eq.(7), the likelihood function of the parameters $\underline{\beta}$ and \mathbf{V} having elements known as variance components σ_i^2 ; $i = 0, 1, 2, \dots, r$ ($\sigma_0^2 = \sigma_\varepsilon^2$) is as follows;

$$(8) \quad \ln L(\underline{\beta}, \mathbf{V} / \underline{\mathbf{Y}}) = -\frac{N}{2} \ln(2\pi) - \frac{1}{2} \ln |\mathbf{V}| - \frac{1}{2} (\underline{\mathbf{Y}} - \mathbf{X}\underline{\beta})' \mathbf{V}^{-1} (\underline{\mathbf{Y}} - \mathbf{X}\underline{\beta})$$

The ML equations that maximize the likelihood function given by Eq.(8) for a $\mathbf{P}_{n \times n}$; symmetric square matrix satisfying

$$(9) \quad \mathbf{P} = \mathbf{V}^{-1} - \mathbf{V}^{-1} \mathbf{X} (\mathbf{X}' \mathbf{V}^{-1} \mathbf{X})^{-1} \mathbf{X}' \mathbf{V}^{-1} = \mathbf{K} (\mathbf{K}' \mathbf{V} \mathbf{K})^{-1} \mathbf{K}'$$

condition are as follows;

$$(10) \quad (\mathbf{X}' \hat{\mathbf{V}}^{-1} \mathbf{X}) \hat{\underline{\beta}} = \mathbf{X}' \hat{\mathbf{V}}^{-1} \underline{\mathbf{Y}}$$

$$(11) \quad \text{tr}(\hat{\mathbf{V}}^{-1} \mathbf{Z}_i \mathbf{Z}_i') = \underline{\mathbf{Y}}_i' \hat{\mathbf{P}} \mathbf{Z}_i \mathbf{Z}_i' \hat{\mathbf{P}} \underline{\mathbf{Y}} \quad i = 0, 1, 2, \dots, r$$

where $\underline{\mathbf{u}}_0 = \underline{\varepsilon}$, $\sigma_0^2 = \sigma_\varepsilon^2$, $q_0 = N$ and $\mathbf{Z}_0 = \mathbf{I}_N$. On the other hand the second ML equation given by Eq.(11) is nonlinear with respect to the variance components occurred in \mathbf{V}^{-1} . Thus closed form expressions for the solutions of Eq.(11) can be obtained iteratively by Newton-Raphson or Fisher's Scoring algorithms (Searle et al., 1992).

For $\underline{\mathbf{Y}}_{n \times 1}$; the vector of response variables having the assumption given by Eq.(7), REML method is based on $\mathbf{K}' \underline{\mathbf{Y}}$; linear combinations of elements of the responses chosen in such a way that those combinations do not contain any fixed effects satisfying $\mathbf{K}' \mathbf{X} = \mathbf{0}$ condition for a $\mathbf{K}' = [\underline{k}'_1, \underline{k}'_2, \dots, \underline{k}'_{N-r_X}]$ matrix having elements $N - r_X$ linearly independent \underline{k}' vectors satisfying $\underline{k}' \mathbf{X} = \mathbf{0}$ by making suitable replacements,

$$(12) \quad \begin{array}{l} \underline{\mathbf{Y}} \text{ by } \mathbf{K}' \underline{\mathbf{Y}} \\ \mathbf{X} \text{ by } \mathbf{K}' \mathbf{X} = \mathbf{0} \end{array} \quad \text{and} \quad \begin{array}{l} \mathbf{Z} \text{ by } \mathbf{K}' \mathbf{Z} \\ \mathbf{V} \text{ by } \mathbf{K}' \mathbf{V} \mathbf{K} \end{array}$$

Then the restricted likelihood function of the parameter $\mathbf{K}' \mathbf{V} \mathbf{K}$ having only variance components for

$$(13) \quad \mathbf{K}' \underline{\mathbf{Y}} \sim N(\underline{\mathbf{0}}, \mathbf{K}' \mathbf{V} \mathbf{K})$$

is as follows;

$$(14) \quad \ln L_R(\mathbf{K}' \mathbf{V} \mathbf{K} / \mathbf{K}' \underline{\mathbf{Y}}) = -\frac{(N-r)}{2} \ln(2\pi) - \frac{1}{2} \ln |\mathbf{K}' \mathbf{V} \mathbf{K}| - \frac{1}{2} \underline{\mathbf{Y}}' \mathbf{K} (\mathbf{K}' \mathbf{V} \mathbf{K})^{-1} \mathbf{K}' \underline{\mathbf{Y}}$$

Then the REML equations that maximize the restricted likelihood function given by Eq.(14) for a $\mathbf{P}_{n \times n}$ matrix satisfying Eq.(9) are as follows;

$$(15) \quad tr \left(\widehat{\mathbf{P}} \mathbf{Z}_i \mathbf{Z}'_i \right) = \underline{Y}' \widehat{\mathbf{P}} \mathbf{Z}_i \mathbf{Z}'_i \widehat{\mathbf{P}} \underline{Y} \quad i = 0, 1, 2, \dots, r$$

Again the REML equations given by Eq.(15) are nonlinear with respect to the variance components occurred in \mathbf{P} matrix including \mathbf{V}^{-1} can be obtained iteratively by Newton-Raphson or Fisher's Scoring algorithms (McCulloch and Searle, 2001).

From Eq.(1) it can be easily seen that *fixed effects model* is a special case of LMM when $\mathbf{Z} = \mathbf{0}$ and $\mathbf{R} = \sigma_\epsilon^2 \mathbf{I}_n$. *Random effects models* are special case of *linear mixed models*. The main importance of *random effects models* for longitudinal data is the necessity of modeling natural heterogeneity across individuals in their response profile over time. In the following subsections *fixed effects models*, *random intercept model* and *random intercept and slope model* from *random effects models* as a special case of *linear mixed models* are taken into consideration.

2.1. Fixed Effects Models

A *fixed effects model* or commonly called *general linear model (GLM)* for the statistical analysis of such a repeated measurements design includes only modeling the expected value of the observations as a linear function of explanatory variables based on the assumption that the observations from different subjects are statistically independent and uncorrelated and that the variance-covariance structure is the same for each subject (Landau and Everitt, 2004). The *fixed effects model* for the response given by individual i at time t_j , y_{ij} is modeled as;

$$(16) \quad y_{ij} = \beta_0 + \beta_1 (group_i) + \beta_2 t_j + \varepsilon_{ij}$$

where $group_i$ is the dummy variable indicating the treatment group to which individual i belongs, β_0 , β_1 and β_2 are the usual regression coefficients for the model; β_0 is the intercept, β_1 represents the treatment group effect and β_2 the slope of the linear regression of outcome on time. ε_{ij} is the usual residual or error term, assumed to be normally distributed with zero mean and variance σ^2 .

Fixed effects model analysis in repeated measurements data is valid under three assumptions about the data:

1. **Normality:** the data arise from populations with normal distribution.
2. **Homogeneity of variance:** the variances of the assumed normal distributions are equal. Levene (1960) developed a test statistic for testing the *homogeneity of variance* assumption using the F distribution with $g - 1$ and $n - g$ degrees of freedom at a significance level of α that is $F_{g-1, n-g, \alpha}$ called *Levene test statistic*. The formula for *Levene's test statistic L* is as follows;

$$(17) \quad L = \frac{(n-g) \sum_{i=1}^g n_i (\bar{Z}_{i.} - \bar{Z}_{..})^2}{(g-1) \sum_{i=1}^g \sum_{j=1}^{n_i} (Z_{ij} - \bar{Z}_{i.})^2}$$

where n is the total number of subjects, g is the number of groups, n_i is the number of subjects in the I^{th} group, $\bar{Z}_{i.}$ are the group means of the Z_{ij} , $\bar{Z}_{..}$ is the overall mean of the Z_{ij} and Z_{ij} is defined as $Z_{ij} = |y_{ij} - \bar{y}_{i.}|$ and $\bar{y}_{i.}$ as $\bar{y}_{i.} = \sum_{j=1}^{n_i} y_{ij}/n_i$ the mean of the I^{th} group.

3. Sphericity condition: the covariance matrix of the observations must have *compound symmetry pattern* given by Eq.(22). Mauchly (1940) developed a test statistic for testing the *sphericity* assumption using the chi-square distribution called *Mauchly's sphericity test*. The formulas for *Mauchly's test statistic* W are as follows;

$$(18) \quad W = |\mathbf{CSC}'| / (\text{trace}\mathbf{CSC}'/p)^p$$

$$(19) \quad \chi_{p(p+1)/2-1}^2 = - \left(\sum_{i=1}^g n_i - g \right) \left(1 - \frac{2p^2 + p + 2}{6p \left(\sum_{i=1}^g n_i - g \right)} \right) \ln(W)$$

where p is the number of parameters in the model, g is the number of groups, n_i is the number of subjects in the I^{th} group, \mathbf{C} is a contrast matrix with p rows suitable for testing a main effect or interaction, \mathbf{S} is a $g \times g$ matrix of the pooled group covariances;

$$(20) \quad \mathbf{S} = \frac{\sum_{i=1}^g \sum_{j=1}^{n_i} (y_{ij} - \bar{y}_{i.})(y_{ij} - \bar{y}_{i.})'}{\sum_{i=1}^g n_i - g}$$

where $\bar{y}_{i.} = \sum_{j=1}^{n_i} y_{ij}/n_i$ is the mean of the I^{th} group (Rencher, 1995).

When the last assumption is not thought to be valid, then LMMs having flexible covariance structures are used to analyze data in repeated measurements design (Landau and Everitt, 2004).

On the other hand making such a restrictive assumption like repeated measurements of a response variable are independent and uncorrelated with the same

variance-covariance structure for each subject is an unrealistic assumption for modeling longitudinal data. Because of this reason *random effects models* allowing within-group correlations between repeated measurements and modeling the covariance structure of each subject given in the following subsection are better than fixed effects models (Wolfinger and Chang, 1999).

2.2. Random Effects Models

Suppose that we have observations made in a repeated measurements study at time points t_1, t_2, \dots, t_T . Assume that we have a single covariate called *treatment group* coded as a zero or one dummy variable. We want to model the response of an individual at time point t_j in terms of *treatment group* and *time*. We assume that there is no *treatment* \times *time* interaction. For this aim two possible *random effects model* types called *random intercept model* and *random intercept and slope model* can be used to model individuals' responses over time.

2.2.1. Random Intercept Model

The simplest case of a linear mixed model is the *random intercept model* based on the *fixed effects model* with a randomly varying subject effect. The *random intercept model* for the response given by individual i at time t_j, y_{ij} is modeled as;

$$(21) \quad y_{ij} = \beta_0 + \beta_1 (\text{group}_i) + \beta_2 t_j + u_i + \varepsilon_{ij}$$

where group_i is the dummy variable indicating the *treatment group* to which individual i belongs, β_0 , β_1 and β_2 are the usual regression coefficients for the model; β_0 is the intercept, β_1 represents the *treatment group* effect and β_2 the slope of the linear regression of response variable on *time*. ε_{ij} is the usual residual or random error term, assumed to be normally distributed with mean zero and variance σ^2 that is $\varepsilon_{ij} \sim N(0, \sigma^2)$. u_i terms are the random effects that model possible heterogeneity in the intercepts of the individuals and are assumed normally distributed with zero mean and variance σ_u^2 that is $u_i \sim N(0, \sigma_u^2)$. The ε_{ij} and u_i terms are assumed independent of one another that is $\text{cov}(\varepsilon_{ij}, u_i) = 0$.

In the *random intercept model* each individual's trend over time is assumed parallel to the treatment group's average trend only the intercepts of each individual's regression line differ by including the random subject effect term u_i into the model is graphically illustrated in Figure1.

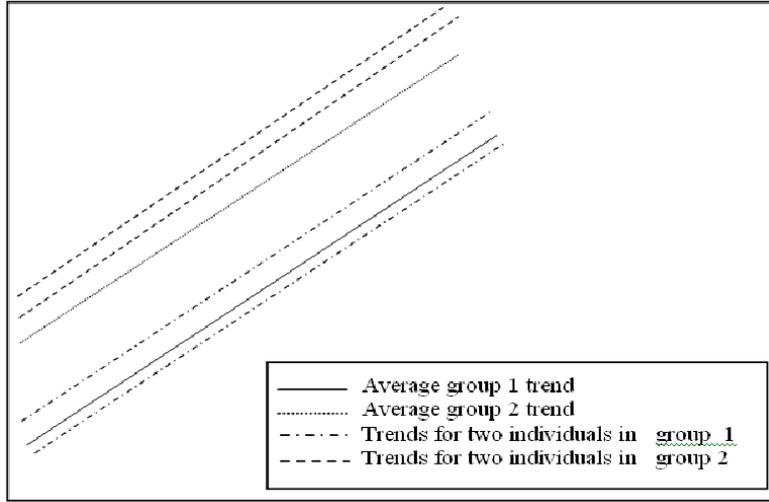


Figure1. Graphical illustration of *random intercept model* by time versus response

The presence of the u_i terms specifies modeling covariance pattern for the repeated measurements of the response variable by an appropriate covariance matrix called *compound symmetry (CS) pattern* as follows;

$$(22) \quad R_i = \begin{bmatrix} \sigma_\epsilon^2 + \sigma_u^2 & \sigma_u^2 & \sigma_u^2 & \sigma_u^2 \\ \sigma_u^2 & \sigma_\epsilon^2 + \sigma_u^2 & \sigma_u^2 & \sigma_u^2 \\ \sigma_u^2 & \sigma_u^2 & \sigma_\epsilon^2 + \sigma_u^2 & \sigma_u^2 \\ \sigma_u^2 & \sigma_u^2 & \sigma_u^2 & \sigma_\epsilon^2 + \sigma_u^2 \end{bmatrix}$$

where σ^2 represents variance within subjects that is $var(\varepsilon_{ij}) = \sigma^2$, and σ_u^2 ; covariance between measurements within subjects or variance between subjects for repeated measurements on the i^{th} subject that is $var(u_i) = \sigma_u^2$ (Iyit et al., 2006). As seen from Eq.(22) *compound symmetry (CS) covariance structure* defines equal variances and covariances between measurements with the same diagonal elements given by $\sigma^2 + \sigma_u^2$ and the off-diagonal elements given by σ_u^2 . So this structure is very important to satisfy the *sphericity* assumption as mentioned in Section 2.1.

On the other hand this covariance structure satisfying *sphericity* assumption is not realistic for most repeated measurements data because it specifies that each pair of the repeated measurements on the same subject is equally correlated. It seems reasonable that the observations closer in time will be more highly correlated than those taken further apart from the same subject.

2.2.2. Random Intercept and Slope Model

A model that allows a more realistic structure than the *random intercept model* for specifying covariance structures of the observations is called *random intercept*

and slope model that allows heterogeneity in both intercepts and slopes (İyit and Genç, 2005a,b). The *random intercept and slope model* for the response given by individual i at time t_j, y_{ij} is modeled as;

$$(23) \quad y_{ij} = \beta_0 + \beta_1 (\text{group}_i) + \beta_2 t_j + u_{i1} + u_{i2} t_j + \varepsilon_{ij}$$

where u_{i1} and u_{i2} terms are the random effects that model possible heterogeneity in intercepts and in slopes of the individual's own regression line respectively and are assumed bivariate normally distributed with zero means, variances $\sigma_{u_1}^2$, $\sigma_{u_2}^2$ and covariance $\sigma_{u_1 u_2}$.

In the *random intercept and slope model* each individual's trend over time is not assumed parallel to the treatment group's average trend as in *random intercept model* so the intercepts and slopes of each individual's regression line differ in this case by including the random subject effect terms u_{i1} and u_{i2} into the model is graphically illustrated in Figure 2.

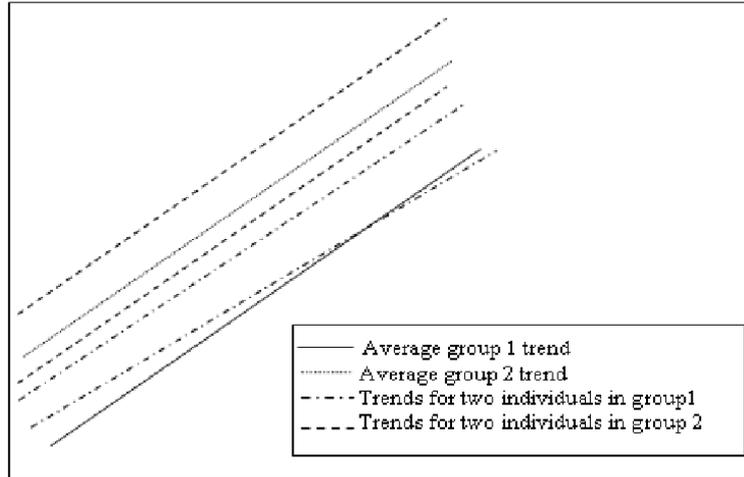


Figure 2. Graphical illustration of *random intercept and slope model* by time versus response

This flexibility ensures the *random intercept and slope model* to model the covariance matrix of the repeated measurements with a more complex pattern allowing variances and covariances to change over time. These two advantages make the *random intercept and slope model* superior in the competition with the *random intercept model* in *random effects models*.

2.3. Model Selection in Fixed and Random Effects Models

Likelihood ratio (LR) test obtained by taking twice the difference in the respective maximized log-likelihoods;

$$(24) \quad G^2 = 2(\log L_{full} - \log L_{reduced})$$

compared with a chi-square distribution with degrees of freedom equal to the difference between the number of parameters in the full and reduced models or the information criteria shown below can be used as a goodness-of-fit statistics to test whether one model is significantly better than the other (Landau and Everitt, 2004, Fitzmaurice et al., 2004);

$$(25) \quad \begin{array}{ll} \text{Akaike's Information} & \\ \text{Criterion;} & AIC = -2 \log L + 2p \\ \text{Hurvich ve Tsai's} & \\ \text{Criterion;} & AICC = -2 \log L + 2p(n/(n-p-1)) \\ \text{Bozdogan's} & \\ \text{Criterion;} & CAIC = -2 \log L + p(\log(n) + 1) \\ \text{Schwarz's Bayesian} & \\ \text{Information Criterion;} & BIC = 2 \log L + p \log(n) \end{array}$$

where $\log L$ is the maximized log-likelihood (ML) or maximized restricted log-likelihood (REML), p is the total number of parameters in the model (ML) or number of parameters in the covariance structure (REML) and n is the number of subjects included in the data set. The smaller the information criteria, the better the model happens to be (Fitzmaurice, 2004).

3. Application

The data used in this study are taken from a placebo controlled clinical trial of a treatment for post-natal (PND) depression in psychiatry given by Landau and Everitt (2004). Around 1 in every 10 women has PND after having a baby starting within a month of the birth up to six months later. A mother who suffers from PND shows some symptoms such as feeling depressed, guilty and anxious, getting irritable with her baby, being tired and sleepless, unable to enjoy anything and unable to cope with her baby (Wheatley, 2005). On the other hand the hormones play an important role on women with PND. It is known that levels of estrogen hormone drop suddenly after the baby is born (Lawrie et al., 2004, Gregoire et al., 1996). This trial is designed to assess the effectiveness of an estrogen treatment compared with a placebo treatment for patients suffer from PND. The main outcome measure used in this trial is depression scores recorded on eight monthly visits.

The main interest about these data is to determine whether the estrogen treatment helps reduce PND by using the appropriate random effects models. The data handed are for 63 patients, 27 in the placebo treatment group and 36 in the estrogen treatment group. These kind of data are repeated measurements data with *time* as the single *within-subject factor* which explains the within-subject variability, i.e., they are longitudinal data and *treatment group* as the

between-subject factor which explains the between-subject variability coded as follows: placebo treatment=0, estrogen treatment=1.

In this study *fixed effects models* and *random effects models* such as *random intercept model*, *random intercept and slope model* as a special case of *linear mixed models* are fitted to the repeated measurements data by using SPSS 13.0 for Windows package programme.

The variables that we wish to include in the linear mixed model are defined as follows; measure of depression scores recorded on eight monthly visits is specified as dependent variable labeled **depscores** (at 1 month after start of the treatment: **depscores1m** up to 8 months after start of the treatment: **depscores8m**). The effect of *time* factor having levels with values 1 up to 8 months modeled by a linear effect of generating a continuous variable labeled **c_month** by recoding the time factor levels and standardizing them and treatment effect labeled **treatment** as a categorical explanatory variable are taken as fixed effects and a subject identifier (**subject**) is included into the model as a random effect.

In order to get a first impression of the data a clustered error bar graph corresponding to the treatment groups for repeated measurements is given in Figure 3.

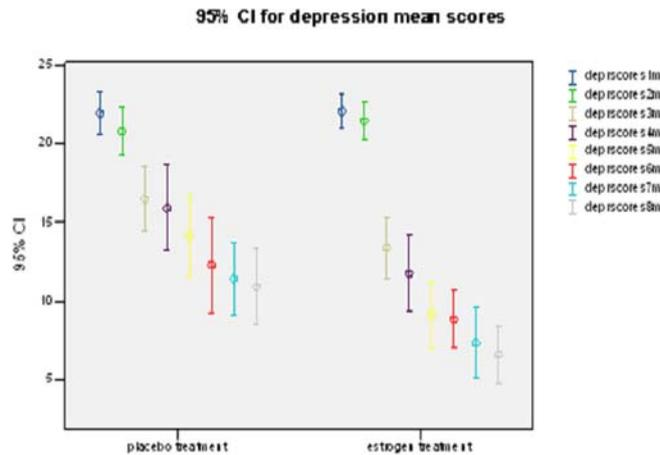


Figure 3. Clustered error bargraph for PND repeated measurements data

From Figure 3 it can be concluded that the depression scores are tending to decrease over time in both treatment groups with an apparent larger decrease in the estrogen treatment group at 3 months after starting the treatment.

Before fitting either of the fixed and random effects models to the repeated measurements data it is important to examine the correlational structure of the depression scores within each treatment group correlation matrices. In the placebo treatment group generally there is an apparent first increase and then decrease in the strength of the correlations with increasing length of time between the

repeated measurements and in the estrogen treatment group the strength of the correlations is tending to increase also including irregular patterns with respect to the time. So the statistical analysis of such a repeated measurements design including modeling within-subject variance-covariance structure depends on such a restrictive assumption that the repeated measurements of the outcome variable are independent, uncorrelated with the same variance-covariance structure for each subject is an invalid assumption for modeling the PND scores data. Because of this reason random effects models allowing within-treatment group correlations between repeated measurements and modeling the covariance structure of each subject are better than fixed effects models.

Modeling the PND scores data with the fixed effects model includes an intercept parameter, two regression coefficients for each of the explanatory variables and the variance parameter of the residual terms in the fitted fixed effects model.

The “Estimates of Fixed Effects” table given by Table 1 gives the t-tests and associated p-values for assessing the regression coefficients. For a model including only fixed effects the results suggest that the intercept term and the fixed effects of time (c_month) and treatment explanatory variables are statistically significant at the 5% significance level and are predictive of modeling the PND scores repeated measurements data.

Table 1. Estimates of Fixed Effects for Fixed Effects Model

						95% Confidence Interval	
Parameter	Estimate	Std. Error	df	t	Sig.	Lower Bound	Upper Bound
Intercept	15.286	0.403	416	37.966	0.000	14.495	16.077
treatment	-2.641	0.519	416	-5.088	0.000	-3.661	-1.621
c_months	-4.806	0.255	416	-18.826	0.000	-5.307	-4.304

The fixed effects model for the PND scores repeated measurements data is a multiple regression model in which there is no random effect as follows;

$$(26) \quad y_{ij} = 15.286 - 2.641 (\text{group}_i) - 4.806t_j + \varepsilon_{ij}$$

The “Estimates of Covariance Parameters” table given by Table 2 gives details of the single covariance parameter in the fixed effects model, namely, the variance of the residual term. This table provides an estimate, confidence interval and test for zero variance. The variance of the residual term is clearly not zero as seen from the confidence interval for the parameter [23.708, 31.112] given in Table 2 which gives an alternate test, the Wald test (Wald z=14.422, p<0.05), for testing the null hypothesis that the variance parameter is zero.

Table 2. Estimates of Covariance Parameters for Fixed Effects Model

					95% Confidence Interval	
Parameter	Estimate	Std. Error	Wald Z	Sig.	Lower Bound	Upper Bound
Residual	27.159	1.883	14.422	0.000	23.708	31.112

On the other hand the *Mauchly's test statistic* for testing the sphericity assumption that is the covariance matrix of the observations have *compound symmetry pattern* is rejected with Mauchly's W-value; 0.062 and related χ^2 -value; 112.562, p-value less than 0.05.

Now we shall fit some random effects models proper to the PND scores data, beginning with the random intercept model. Modeling the PND scores data with the random intercept model indicates that subject random effects for the intercept have been added to the model. This results in fitting one extra model parameter, the variance of the subject random effects.

In the random intercept model case for the PND scores data, it can be easily seen from Table 3 and Table 4 that the model includes an intercept parameter, two regression coefficients; one for each of the explanatory variables and the variance parameters of the residual and the intercept terms in the model.

Table 3. Estimates of Fixed Effects for Random Intercept Model

						95% Confidence Interval	
Parameter	Estimate	Std. Error	df	t	Sig.	Lower Bound	Upper Bound
Intercept	15.221	0.719	63.076	21.155	0.000	13.784	16.659
treatment	-2.308	0.945	61.317	-2.443	0.017	-4.197	-0.419
c_months	-4.738	0.206	376.242	-23.033	0.000	-5.143	-4.334

The random intercept model allowing to model possible heterogeneity in the intercepts of the individuals for the PND scores repeated measures data is as follows;

$$(27) \quad y_{ij} = 15.221 - 2.308(\text{group}_i) - 4.738t_j + u_i + \varepsilon_{ij}$$

From Table 4, for testing the null hypothesis that the variance of the residual term and the subject intercept effect are zero, the Wald test (Wald z=13.350 and 4.422, p<0.05) results and the confidence intervals for the parameters [14.019, 18.804] and [6.997, 16.979] respectively show that the covariance parameters are statistically significantly different from zero.

Table 4. Estimates of Covariance Parameters for Random Intercept Model

					95% Confidence Interval	
Parameter	Estimate	Std. Error	Wald Z	Sig.	Lower Bound	Upper Bound
Residual	16.237	1.216	13.350	0.000	14.019	18.804
Intercept [subject ID diagonal SC]	10.899	2.465	4.422	0.000	6.997	16.979

Finally modeling the PND scores data with the random intercept and slope model indicates that both the intercept term and the regression coefficient of time effect (c_month) are allowed to vary between subjects. This case results in fitting two extra model parameters to the model, the variance of the subject intercept random effects and the variance of the random slope effects.

From Table 5 the finally fitted random intercept and slope model describes the average profiles of the PND scores data in the two treatment groups over the eight time point visits including random intercepts for subjects and fixed effects of treatment and time.

Table 5. Estimates of Fixed Effects for Random Intercept and Slope Model

						95% Confidence Interval	
Parameter	Estimate	Std. Error	df	t	Sig.	Lower Bound	Upper Bound
Intercept	15.307	0.618	110.534	24.783	0.000	14.083	16.531
treatment	-2.521	0.801	108.131	-3.148	0.002	-4.108	-0.934
c_months	-4.773	0.394	108.341	-12.125	0.000	-5.554	-3.993

The random intercept and slope model allowing to model possible heterogeneity in intercepts and in slopes of the individual's own regression line for the PND repeated measurements data is as follows:

$$(28) \quad y_{ij} = 15.307 - 2.521(\text{group}_i) - 4.773t_j + u_{i1} + u_{i2}t_j + \varepsilon_{ij}$$

From Table 6, for testing the null hypothesis that the variance of the residual term and the variance of the subject intercept random effect and the random slope effect are zero, the Wald test (Wald $z=12.314$ and 5.569 , $p<0.05$) results and the confidence intervals for these parameters $[12.034, 16.544]$ and $[4.693, 9.487]$ respectively show that the covariance parameters are statistically significantly different from zero. Then it can be concluded that the random intercept and slope model provides an adequate description of the PND scores data.

Table 6. Estimates of Covariance Parameters for Random Intercept and Slope Model

Parameter	Estimate	Std. Error	Wald Z	Sig.	95% Confidence Interval	
					Lower Bound	Upper Bound
Residual	14.110	1.146	12.314	0.000	12.034	16.544
Intercept+ c_months [subject ID diagonal SC]	6.672	1.198	5.569	0.000	4.693	9.487

The “Information Criteria” table given by Table 7 will be helpful for comparing the fixed effects model and the random intercept model with the finally fitted random intercept and slope model. The information criteria are displayed in Table 7 in smaller-is-better forms.

Table 7. Information Criteria for Fixed Effects Models and Random Intercept/Slope Models

	Fixed Effects Model	Random Intercept Model	Random Intercept and Slope Model
-2 Restricted Log Likelihood	2570.748	2457.092	2370.748
Akaike's Information Criterion (AIC)	2572.748	2461.092	2374.748
Hurvich and Tsai's Criterion (AICC)	2572.757	2461.121	2374.777
Bozdogan's Criterion (CAIC)	2577.778	2471.154	2384.809
Schwarz's Bayesian Criterion (BIC)	2576.778	2469.154	2382.809

From Table 7 the difference in $-2 \times$ restricted log likelihood for the three models can be tested as a chi-square distribution with degrees of freedom given by the difference in the number of parameters in each of the three models. This is known as *likelihood ratio test*. Here the difference between the fixed effects model likelihood and the random intercept model likelihood is $2570.748 - 2457.092 = 113.656$, which is tested as a chi-square with one degree of freedom, $\chi^2_{1, 0.95} = 3.84$ has an associated p-value 0.01. So the random intercept model is found better than the fixed effects model. When we compare the random intercept model with the random intercept and slope model, the difference between these two model likelihoods is $2457.092 - 2370.748 = 86.344$ with the same chi-square and associated p-value. Then the random intercept and slope model clearly provides a better fit for the PND scores data than the random intercept model. On the other hand in the light of the other information criteria, the lower the information criteria, the better fit the random intercept and slope model has.

There are a number of residual diagnostics that can be used to check the assumptions of a linear mixed model. Firstly the assumption of normality of the random effect terms will be checked by constructing histograms of the variables; residuals and random effects for subjects given by Figure 4.

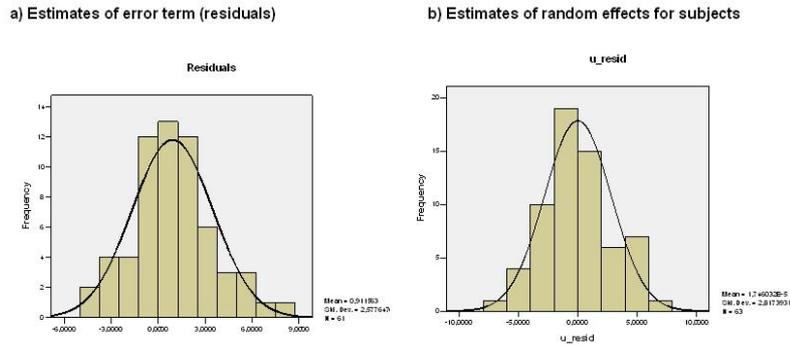


Figure 4. Histograms for estimates of random effects

The resulting histogram given by Figure 4 does not indicate any departure from the normality assumption. The next one that will be checked is the assumption of homogeneity of variance that is the constant error variance across repeated measurements. For checking this assumption Levene test is used to test the null hypothesis that the error variance of the dependent variable is equal across groups and not rejected with F-value; 0.067 and related p-value; 0.797.

4. Results and Discussion

In this study a comparative evaluation of fixed effects models with random intercept models and random intercept and slope models as a special case of random effects models from linear mixed models are taken into consideration following a repeated measurements design having one between-subject factor (referred to as treatment) with two treatment levels and one within-subject continuous covariate (referred to as time).

Before fitting either of the fixed and random effects models to the repeated measurements data, the correlational structure of the PND scores within each treatment group correlation matrices is examined and it is observed that generally covariances between observations made closer together in time are likely to be higher than those made at greater time points. So the fixed effects model including only modeling the expected value of the observations as a linear function of explanatory variables based on the assumption that the observations from different subjects are statistically independent and uncorrelated and that the variance-covariance structure is the same for each subject is not found appropriate for the statistical analysis of such a repeated measurements design. Also as a result of *Mauchly's test* for testing the null hypothesis that the covariance matrix of the observations have *compound symmetry pattern* is rejected then the sphericity assumption is not satisfied for constructing fixed effects models fitted

to PND scores repeated measurements data. On the other hand the random intercept model implying a particular structure for the covariance matrix of the repeated measurements that variances at different time points are equal and covariances between each pair of time points are equal is in fact a very restrictive and unrealistic assumption. The random intercept and slope model that allows a less restrictive covariance structure for the repeated measurements provides an improvement in fit over the random intercept model.

In the random intercept model, the groups differ with respect to the value of the intercept graphically illustrated by Figure 1. This results in fitting one extra model parameter, the variance of the subject random effects. On the other hand in the random intercept and slope model, there are random slopes as well as random intercepts, one could try to explain the variability of slopes as well as intercepts and both the intercept term and the regression coefficient of time effect (c_month) are allowed to vary between subjects graphically illustrated by Figure 2. This results in fitting two extra model parameters, the variance of the subject random effects and the variance of the random slope effects. Also from tests of fit of these two competing models from Table 7, it can be easily seen that random intercept and slope model better fit to the data than random intercept model. These advantages make the random intercept and slope model allowing a more complex pattern for the covariance matrix of the repeated measurements superior in the competition with random intercept model in random effects models.

5. Conclusion

The statistical analysis of such a repeated measurements design includes modeling not only the expected value of the observations as in the fixed effects models but also their within-subject variance-covariance structure. Because making such a restrictive assumption like repeated measurements of a response variable are independent, uncorrelated with the same variance-covariance structure for each subject is an unrealistic assumption for modeling longitudinal data, it can be concluded that the random effects models allowing within-group correlations between repeated measurements and modeling the covariance structure of each subject are better than the fixed effects models. In the light of this study it would be interesting to model the covariance structure of random intercept and slope model in various forms other than compound symmetry pattern in a further study.

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