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Linear Models for Multivariate Repeated Measures Data

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LINEAR MODELS FOR MULTIVARIATE REPEATED MEASURES DATA

by
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M.Sc. May 1988, Bangalore University
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May 1996

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ABSTRACT

LINEAR MODELS FOR MULTIVARIATE REPEATED MEASUREMENTS DATA

Shantha S. Rao
Old Dominion University, 1996
Director: Dr. Dayanand N. Naik

In this dissertation we focus mainly on the analysis of continuous multivariate repeated measurements data based on the assumption of multivariate normality. However certain aspects of the analysis of univariate repeated measures data are also considered. Typically, we have measurements on \( p \) variables (possibly correlated) in the form of \( p \times 1 \) vectors \( y_{ijk} \), observed at \( k = 1, 2, \ldots, t_{ij} \) occasions on \( j = 1, 2, \ldots, n_i \) individuals from \( i = 1, 2, \ldots, g \) groups. We assume a naturally occurring covariance structure \( V_{ij} \otimes \Sigma \) among the \( p \) variables on the \( j^{th} \) individual from \( i^{th} \) group made at \( t_{ij} \) occasions. Here \( V_{ij} \) and \( \Sigma \) are positive definite matrices of order \( t_{ij} \times t_{ij} \) and \( p \times p \) respectively. We develop a general linear model approach to accommodate both balanced and unbalanced repeated measures data.

Our main results are: (1) construction of Rao's score test for a simpler model with \( p = 1 \) (univariate case) and \( V_{ij} \) having a structure as in a mixed effects model, (2) comparison of all the methods for analyzing univariate repeated measures data.
with time varying covariates, (3) derivation of the maximum likelihood estimates of the covariance matrices \( V \) and \( \Sigma \) in the balanced case, (4) derivation of Satterthwaite type approximation to the distribution of multivariate quadratic forms, (5) estimation of degrees of freedom for these approximations, and (6) derivation of the maximum likelihood estimates of the covariance parameters under certain specific covariance structures for unbalanced case.

Rao's score test is derived in Chapter 2. Analysis of repeated measures in the presence of time varying covariates is a useful but difficult problem. In Chapter 3, we review the existing methods for analyzing repeated measured data with time varying covariates and discuss their computational aspects using SAS software. We also point out that a linear model approach yields an unified tool to analyze these data. In Chapter 4, various results about balanced multivariate repeated measures models are derived. We present the entire scheme of analysis of balanced multivariate data including the computational details. Finally, the analysis of unbalanced multivariate repeated measures is discussed in Chapter 5. In this case we assume two commonly used covariance structures namely equicorrelation and autoregressive structures for \( V_{ij} \) and derive the maximum likelihood estimates of the unknown parameters.
Dedicated

to

My parents
ACKNOWLEDGMENTS

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Many thanks are due to all my friends who helped me get through difficult times. Above all I am proud to acknowledge the immense contribution made by members of my family. This dream would not have been realized without the constant support and encouragement received from them. The love and affection showered by each one of them went a long way in guiding me through times when I had almost given up hope. As a mark of affection I dedicate this work to my family.
Contents

1 Introduction 1

2 Mixed Effects Models For Univariate Repeated Measures 12
   2.1 Introduction .......................................................... 12
   2.2 Score test for $H_0 : \xi = 0.$ ................................. 15
      2.2.1 A special case ................................................. 24
   2.3 Analysis under the reduced model ......................... 27
      2.3.1 Generalized least squares method ..................... 27
      2.3.2 Likelihood ratio method ................................. 28
      2.3.3 Score test ..................................................... 29
      2.3.4 Approximate $F$ test ...................................... 31
   2.4 Appendix ............................................................. 33
      2.4.1 Model and estimators ..................................... 33
      2.4.2 Estimation under the null hypothesis $K\beta = 0$ .. 35
List of Tables

3.1 Serum Anti-Rotavirus Antibody Concentration ...................................... 46
3.2 Plasma renin activity data .......................................................................... 54
4.1 Repeated measures data ............................................................................. 65
4.2 Manova table for mixed effects model ....................................................... 70
4.3 Dental data .................................................................................................... 86
4.4 Approximate MANOVA ............................................................................. 88
Chapter 1

Introduction

Repeated measurements, as indicated by the name, are successive measurements made over time on each experimental unit. If measurements are made on \( n \) individuals over \( t \) time points on a single characteristic, we have a set of univariate repeated measurements data. On the other hand, if measurements are made on a set of \( t \) repeated measurements on \( p \) variables (or characteristics) on each of the \( n \) individuals, we have a set of multivariate repeated measurements data. Thus in this case the data on each individual is a \( p \times t \) matrix.

The \( n \) individuals may be randomly divided into one of several groups to which the levels of an experimental factor are assigned. Repeated measurements occur very frequently in a variety of scientific fields where statistical models are used. The distinguishing characteristics of these measurements from those in the usual or more traditional regression setting is that one or more response variables are measured on
the same observational unit more than once, hence making the responses dependent.

The questions that one would generally like to ask and answer, in this context are

- Is there a group effect?
- Is there a time effect?
- Is there an interaction between time and group effects?

Let us first consider a set of univariate repeated measures. In an attempt to answer the above questions treat the $t$ measurements $Y'_{ij} = (y_{ij1}, \ldots, y_{ijt})$, on the $j^{th}$ individual belonging to the $i^{th}$ group as a single multivariate observation $Y_{ij}$ rather than as a set of separate univariate observations. Assuming a general covariance structure, we can use multivariate techniques to analyze these type of data. To be more specific, the above questions can be formulated as linear hypotheses of the form $H_0: C \mu_{ij} = 0$, for appropriate choices of $C$, where $\mu_{ij}$ is the mean or the expected value of $Y_{ij}$. Then a profile analysis (MANOVA) can be used to test $H_0$.

The main reason that prompted us to use the multivariate approach is that we had no particular structure assumed for $\text{cov}(Y_{ij}) = V$. However in many practical situations, $V$ is found to have some simpler structures. In that case, multivariate tests are less powerful. As an alternative to the multivariate approach, by considering each subject as a random block and time points as plots within blocks a split plot design model can be used to analyze repeated measures data.
The model in that case is

\[ y_{ijk} = \mu + \alpha_i + \beta_k + (\alpha\beta)_{ik} + \delta_{j(t)} + \epsilon_{ijk}, i = 1, \ldots, g, \]

\[ j = 1, \ldots, n_i, \quad k = 1, \ldots, t, \]

\[ \delta_{j(t)} \sim N(0, \sigma^2), \]

\[ \epsilon_{ijk} \sim N(0, \sigma^2), \]

where \( y_{ijk} \) is the \( k^{th} \) observation on the \( j^{th} \) individual from the \( i^{th} \) group and \( \mu, \alpha_i, \beta_k, (\alpha\beta)_{ik} \) and \( \delta_{j(t)} \) have the usual meaning as in split plot design. The problems of interest are to test the following null hypothesis:

(i) \( \alpha_1 = \ldots = \alpha_g \)

(ii) \( \beta_1 = \ldots = \beta_t \) and

(iii) \( (\alpha\beta)_{ik}s \) are all equal.

We know that the usual ANOVA technique can be applied here although the observations \( y_{ijkl} \), \( y_{ijkt} \) on an individual are correlated. As before define, \( Y_{ij} = (y_{ij1}, \ldots, y_{ijt}) \). Then the covariance matrix of \( Y_{ij} = \text{cov}(Y_{ij}) = V = \sigma^2 \mathbf{I} + \sigma^2 \mathbf{J}, \) where \( \mathbf{I} \) is an identity matrix of order \( t \) and \( \mathbf{J} \) is a \( t \times t \) matrix of ones. Further, if we define \( Y' = (Y'_{11}, \ldots, Y'_{1n_1}, Y'_{21}, \ldots, Y'_{2n_2}, \ldots, Y'_{g1}, \ldots, Y'_{gn_g}) \) then \( \text{cov}(Y) = \Omega = \text{diag}(\mathbf{I}_{n_1} \otimes \mathbf{V}, \ldots, \mathbf{I}_{n_g} \otimes \mathbf{V}). \)

The ANOVA table for the above model is
<table>
<thead>
<tr>
<th>Source</th>
<th>D.O.F.</th>
<th>SS</th>
<th>F statistic</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Between individuals</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Groups</td>
<td>$g - 1$</td>
<td>$Q_1 = Y'AY$</td>
<td>$F_1 = (n - g)Q_1/(g - 1)Q_2$</td>
</tr>
<tr>
<td>Individuals</td>
<td>$n - g$</td>
<td>$Q_2 = Y'BY$</td>
<td></td>
</tr>
<tr>
<td><strong>Within individuals</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Time</td>
<td>$t - 1$</td>
<td>$Q_3 = Y'CY$</td>
<td>$F_2 = (n - g)Q_3/Q_5$</td>
</tr>
<tr>
<td>Time*Group</td>
<td>$(g - 1)(t - 1)$</td>
<td>$Q_4 = Y'DY$</td>
<td>$F_3 = (n - g)Q_4/(g - 1)Q_5$</td>
</tr>
<tr>
<td>Error</td>
<td>$(t - 1)(n - g)$</td>
<td>$Q_5 = Y'EY$</td>
<td></td>
</tr>
<tr>
<td>Total</td>
<td>$n - 1$</td>
<td>$Y'TY$</td>
<td></td>
</tr>
</tbody>
</table>

Here $n = \sum_{i=1}^{g} n_i$. The quadratic forms $Q_1$ to $Q_5$ are:

\[
Q_1 = k \sum_{i=1}^{g} n_i (\bar{y}_{..i} - \bar{y}_{..})^2 = Y'AY
\]

\[
Q_2 = k \sum_{i=1}^{g} \sum_{j=1}^{t} (\bar{y}_{ij} - \bar{y}_{..})^2 = Y'BY
\]

\[
Q_3 = n \sum_{k=1}^{t} (\bar{y}_{..k} - \bar{y}_{..})^2 = Y'CY
\]

\[
Q_4 = \sum_{i=1}^{g} n_i \sum_{k=1}^{t} (\bar{y}_{ik} - \bar{y}_{..i} - \bar{y}_{..k} + \bar{y}_{..})^2 = Y'DY
\]

\[
Q_5 = \sum_{i=1}^{g} \sum_{j=1}^{t} \sum_{k=1}^{t} (y_{ijk} - \bar{y}_{ij} - \bar{y}_{i.k} + \bar{y}_{..})^2 = Y'EY
\]

and $Q_1 + \ldots + Q_5 = \sum_{i=1}^{g} \sum_{j=1}^{t} \sum_{k=1}^{t} (y_{ijk} - \bar{y}_{..})^2 = Y'TY$

with appropriate choice of matrices $A$ through $E$, and $T$. These $n \times n$ matrices are all symmetric and can be easily derived.
The tests of hypotheses are carried out using the usual ANOVA F-tests. The split-plot design that we considered accommodates a simple correlation among repeated measures. However, there are certain other covariance structures which also admit the same F tests.

Huynh and Feldt (1970) and Rouanet and Le'pine (1970) have derived a set of necessary and sufficient conditions on the covariance structures such that the distributions of the F statistics remains invariant. Consequently, there are certain dependence among repeated measures which still admit the usual tests. The class of all covariance structures which admit the usual F tests is said to have type H structure. A typical member of this class is $V = \sigma^2(I + a1' + la')$, where $a' = (a_1, \ldots, a_t)$ is such that $V$ is positive definite and $I$ is a vector of all ones.

Recently, Chaganty and Vaish (1995) studied this characterization more closely and produced the following form for $H(a)$:

$$H^*(a) = I + \frac{1}{t}(a1' + la') - \frac{1}{t}(1 + \bar{a})11',$$

where $a' = (a_1, \ldots, a_t)$ is such that

$$\frac{1}{t} \sum_{i=1}^{t} (a_i - \bar{a})^2 \leq \bar{a} \quad (1.2)$$

and $\bar{a}$ is the mean of the components of the vector $a$. This characterization by Chaganty and Vaish (1995) is most explicit in the sense that it gives easy condition (1.2) on the elements of $a$ such that $V$ is positive (semi) definite. Since the form of covariance structure determines the choice of the analysis, we have a need for...
testing for the type H structure for $V$. We have the following result due to Huynh and Feldt (1970) to test for type H structure.

**Result 1**: Let $C_{t-1 \times t}$ be such that $CC' = I_{t-1}$ and $C1 = 0$, and $U_{t-1 \times 1} = C_{t-1 \times t}Y_{1 \times 1}$. Then $\text{cov}(Y) = V$ is of type H structure iff $\text{cov}(U) = CVC' = \lambda I_{t-1}$.

A likelihood ratio test for testing $\text{cov}(U) = \lambda I_{t-1}$ can be constructed using a test of sphericity (Mauchly (1940)) on the transformed variables. The likelihood ratio is given by

$$
\lambda = \frac{|CSC'|}{[1/(t - 1)tr(CSC')]^{t-1}},
$$

where $S$ is the sample variance covariance matrix. Under the null hypothesis of sphericity, that is $\text{cov}(U) = \lambda I_{t-1}$,

$$
-\{(n - 1) - (2(t - 1)^2 + (t - 1) + 2)/(6(t - 1))\} \ln \lambda
$$

has an approximate $\chi^2$ distribution with $\frac{1}{2}t(t - 1) - 1$ degrees of freedom. Standard statistical software like SAS provides a test for sphericity. For details see Khatree and Naik (1995).

So far we have explored the possibility of using the usual F statistics under certain dependent covariance structures. That is, we noted that the ANOVA remains invariant as long as $V$ is of type H structure. Now we need to address the situation where we reject the hypothesis of sphericity. Rejecting the hypothesis of sphericity implies the departure of $V$ from type H structure. Under no structure
on $V$, Geisser and Greenhouse (1958) have shown that $A\Omega B = A\Omega E = B\Omega E = C\Omega E = D\Omega E = 0$. Hence the quadratic forms $Q_1$ and $Q_2$, $Q_1$ and $Q_5$, $Q_2$ and $Q_5$, $Q_3$ and $Q_5$, and $Q_4$ and $Q_5$, are all pairwise independent. However, these quadratic forms no longer have exact $\chi^2$ distributions. The results that follow enable us to find the distributions of these quadratic forms and their ratios.

**Result 2:** Consider $X_{t\times 1} \sim N_t(0, V)$. Let $A$ be any symmetric matrix of order $t \times t$ with $\text{rank}(A) = r$. Then the distribution of any quadratic form $X'AX$ is the same as the distribution of certain linear combinations of independent $\chi^2$ variables. That is, $X'AX \overset{d}{=} \sum_{j=1}^{r} \lambda_j \chi_j^2(1)$, where $\lambda_1, ..., \lambda_r$ are the eigenvalues of a certain matrix to be determined in the proof. Further $\chi_j^2(1)'s$ are all independent.

**Proof:**

Since $V$ is a positive definite matrix we can express the quadratic form $X'AX$ as

$$
X'AX = X'V^{-1/2}V^{1/2}AV^{1/2}V^{-1/2}X
$$

$$
= Z'BZ (Z = V^{-1/2}X, Z \sim N(0, I)),
$$

$$
= Z'\Lambda \Gamma'Z \quad (B = \Gamma\Lambda\Gamma', \, \Gamma' = I \text{ and } \Lambda = \text{diag}(\lambda_1, ..., \lambda_r, 0, ..., 0))
$$

$$
= U'\Lambda U, \quad (U = \Gamma'Z, \, U \sim N(0, I))
$$

$$
= \sum_{j=1}^{r} \lambda_j u_i^2, \quad u_i^2 \sim \chi_j^2(1), \text{ and } \chi_1^2(1), ..., \chi_r^2(1) \text{ are independent.}
$$

We observe that $\lambda_1, ..., \lambda_r$ are the nonzero eigenvalues of $B$. (note that $\text{rank} \ (A) = \text{rank} \ (B) = r$). Hence the result.

We see that the quadratic form $X'AX$ is the weighted sum of independent $\chi^2$ vari-
Satterthwaite (1941) has developed the following approximate \( \chi^2 \) distribution for \( X'AX \).

Suppose the distribution of \( X'AX \) is approximated by \( a\chi^2_r \), where \( a \) and \( b \) are determined satisfying the following conditions:

\[
(i) \ E(X'AX) = E(a\chi^2_r), \quad (1.3)
\]
\[
(ii) \ var(X'AX) = var(a\chi^2_r). \quad (1.4)
\]

Using (1.3) and (1.4) we can show that \( a = \frac{\sum_{j=1}^{r} \lambda_j^2}{\sum_{j=1}^{r} \lambda_j} \) and \( b = \frac{(\sum_{j=1}^{r} \lambda_j)^2}{\sum_{j=1}^{r} \lambda_j^2} \).

From Result 2 we have the following:

\[
Q_1 \sim \chi^2(g - 1)
\]
\[
Q_2 \sim \chi^2(n - g)
\]
\[
Q_3 \sim \sum_{j=1}^{i-1} \lambda_j \chi^2_1(1)
\]
\[
Q_4 \sim \sum_{j=1}^{i-1} \lambda_j \chi^2_1(g - 1)
\]
\[
Q_5 \sim \sum_{j=1}^{i-1} \lambda_j \chi^2_1(n - g),
\]

where \( \lambda_1, \ldots, \lambda_{i-1} \) are the eigenvalues of \( (I - 1/tJ)V \).

Box (1954) applied Satterthwaite approximation to obtain the distribution of the ratio of the quadratic forms. Writing \( \epsilon = \frac{(\sum_{j=1}^{r} \lambda_j)^2}{(t - 1) \sum_{j=1}^{r} \lambda_j^2} \), we can verify the
following

\[ F_1 \sim F_{(\nu-1),(n-\nu)} \text{ (exact)} \]  \hfill (1.5)

\[ F_2 \sim F_{\nu(t-1),\nu(t-1)(n-\nu)} \text{ (approx.)} \]  \hfill (1.6)

\[ F_3 \sim F_{\nu(t-1)(\nu-1),\nu(t-1)(n-\nu)} \text{ (approx.)} \]  \hfill (1.7)

We use the above \( F \) statistics given by (1.5), (1.6) and (1.7) and test the hypotheses of interest. Notice that in applications we need to estimate \( \epsilon \). Greenhouse and Geisser (1959) suggested the estimate of \( \epsilon \) given by

\[ \hat{\epsilon}_{GG} = \frac{[\text{tr}(C'SC)]^2}{(t-1)\text{tr}(C'SC)^2}, \]

where \( S \) is the sample variance covariance matrix. Huynh and Feldt (1976) have suggested an alternative estimate of \( \epsilon \) given by

\[ \hat{\epsilon}_{HF} = \frac{n(t-1)\hat{\epsilon}_{GG} - 2}{(t-1)[n-1-(t-1)]\hat{\epsilon}_{GG}}. \]

We summarize the scheme for analysis of univariate repeated measures using the ANOVA approach as follows:

- If \( \epsilon \) is very small, which is an indication of lack of sphericity, use of the multivariate approach may be suggested.

- If \( \epsilon \) is very close to 1, usual ANOVA for a split plot design may be carried out.
• Otherwise use of ANOVA with an approximate $F$ (use of $\epsilon$ correction) may be appropriate.

Next, we give an outline of the subsequent chapters.

In Chapter 2 we study a generalization of the basic ANOVA model to analyze univariate repeated measures data. Laird and Ware (1982), utilizing the ideas introduced by Harville (1977), introduced mixed effects models to analyze these data. This modeling approach gives us a choice of various covariance structures for both individual random effects component and within individual errors. We consider a mixed effects model assuming within individual errors that follow an autoregressive (AR(1)) structure. We propose a score test for the parameter that accounts for the between individual random effects component. This test is very useful because it tests for the possibility of a simpler model.

In Chapter 3 we study methods of analyzing repeated measures data in the presence of covariates. We review methods of analyzing univariate repeated measures data in presence of two types of covariates: (a) covariates fixed over time and (b) time varying covariates. We provide computer programs, using SAS software to perform each of these analyses.

In Chapter 4 we consider analysis of balanced multivariate repeated measures data. Analysis of these data using a MANOVA model is considered. The well known Satterthwaite type approximation to the distribution of a quadratic form in normal variables is extended to the distribution of a multivariate quadratic form in
multivariate normal variates. The multivariate tests using this approximation are developed for testing the usual hypotheses.

In Chapter 5 we consider analysis of unbalanced multivariate repeated measures data. Use of multivariate linear models for analysis of these data is illustrated. Analysis of these models are discussed under some special covariance structures.
Chapter 2

Mixed Effects Models For

Univariate Repeated Measures

2.1 Introduction

In this chapter we study a generalization of the basic ANOVA model, to analyze repeated measures or longitudinal data. The defining characteristic of a repeated measures or longitudinal study is that the individuals are measured repeatedly over time. Longitudinal studies are designed to investigate changes in a characteristic which is measured over time. In medical experiments, the characteristic of interest might be blood pressure, cholesterol level, lung volume or serum glucose. Suppose we have \( t \) measurements taken repeatedly over time on \( n \) experimental units on a certain characteristic. Our main problem of interest is to make inference about the
differences between the levels of the characteristic at various time points. Stated in other words, our interest is to test a hypothesis about the time effect. To model longitudinal data collected over time for each member of a group of experimental units, one must recognize the correlation between serial observations on the same experimental unit.

In practice, longitudinal data are often unbalanced or incomplete, that is, all individuals are not observed the same number of times or with the same design matrix $X$. In these situations it is imperative to consider models that account for both unbalancedness and correlation among the observations on an individual.

Laird and Ware (1982), utilizing the ideas introduced by Harville (1977), introduced a generalization of the basic ANOVA model to analyze repeated measures data. Let $\beta$ denote a $m \times 1$ vector of unknown population parameters and $X_k$ be a known $t_k \times m$ design matrix. Let $\tau_k$ denote a $q \times 1$ vector of unknown individual random effects and $C_k$ be a known $t_k \times q$ design matrix for individual effects. For each individual $k$, Laird and Ware (1982) considered the following model:

$$Y_k = X_k \beta + C_k \tau_k + \epsilon_k,$$  \hspace{1cm} (2.1)

where $\epsilon_k$ is distributed as $N(0, V_k)$. Here $V_k$ is a $t_k \times t_k$ positive definite covariance matrix, depending on $k$ through its dimension $t_k$. However, the set of unknown parameters in $V_k$ do not depend on $k$. Next, the random vectors $\tau_k$ are distributed as $N(0, \Gamma)$, independently of each other and of $\epsilon_k$. Here $\Gamma$ is a $q \times q$ positive definite
matrix. To summarize we have model (2.1) with the following specifications.

\[ E(\epsilon_k) = 0, \quad V(\epsilon_k) = V_k, \quad (2.2) \]
\[ E(\tau_k) = 0, \quad V(\tau_k) = \Gamma \quad \text{and} \quad Cov(\tau_k, \epsilon_k) = 0. \quad (2.3) \]

Thus for the model (2.1) we have,

\[ E(Y_k) = X_k \beta \quad \text{and} \]
\[ V(Y_k) = C_k \Gamma C_k^t + V_k. \]

For this model with \( V_k \) equal to an identity matrix, Laird and Ware (1982) utilized the EM algorithm to obtain the maximum likelihood (ML) and restricted maximum likelihood (REML) estimates of the variance components. However, the tests of hypothesis about the parameters of the covariance matrix were not considered by them.

Chi and Reinsel (1989) considered model (2.1) that contain both individual random effect component and within individual errors that follow an autoregressive (AR(1)) time series process. They developed a score test for the autocorrelation in the within individual errors for the 'conditional independence' random effects model. The AR(1) structure is particularly appealing because of the fact that it incorporates the natural decay in the correlation between observations taken on the same experimental unit with increasing time lags. In other words, observations on the same experimental unit which are closer together should be more highly correlated than those that are far apart in time.
Chi and Reinsel (1989) illustrated their methods on a data set with $C_k = 1_k$, where $1_k$ is a $t_k \times 1$ vector of ones. This choice of $C_k$ seems to be most useful in practical situations. With this choice of $C_k$, in the following section we derive a score test for the parameter that accounts for the between individual random effects component. Our model for individual $k$ is

$$Y_k = X_k \beta + 1_k \tau_k + \epsilon_k, \quad k = 1, \ldots, n, \quad (2.4)$$

with

$$E(Y_k) = X_k \beta$$

$$\Sigma_k = cov(Y_k) = \sigma^2 11' + \sigma^2 V_k \quad (2.5)$$

where the $(i,j)^{th}$ element of $V_k$ is $\frac{1}{1 - \rho^2 |i-j|}$. Notice that (2.5) can be written as

$$\Sigma_k = cov(Y_k) = \sigma^2 (\xi J_k + V_k), \quad (2.6)$$

where $\xi = \sigma^2 / \sigma^2$ and $J_k$ is a $t_k \times t_k$ matrix of one's.

2.2 Score test for $H_0 : \xi = 0$.

We propose a score test for testing $H_0 : \sigma^2 = 0 \Leftrightarrow H_0 : \xi = 0$. The score test is based on the score criterion suggested by Rao (1948), and it has the advantage over other large sample tests, such as Wald's test and the likelihood ratio test, of requiring
estimation only under the null hypothesis. Computation of maximum likelihood estimates is quite tedious in most cases. However the score test enables considerable savings in terms of computation. It requires estimation only under the null model, and also yields a test asymptotically equivalent to the corresponding likelihood ratio test. Thus the null distribution of the score statistic is asymptotically $\chi^2$ with degrees of freedom equal to the number of parametric restrictions imposed by the hypothesis.

As was mentioned before, Chi and Reinsel (1989) proposed a score test for testing the random effects model (2.1) with $\text{cov}(\epsilon_k) = \sigma^2 I$ against the same model with autocorrelated (AR(1)) errors for the $\epsilon_k$'s. Thus a score test for testing $H_0 : \rho = 0$ (the autoregressive parameter=0) provides a simple check for the presence of possible autocorrelation in the errors. The test for $H_0 : \xi = 0$, that we develop would provide a check for the possibility of a simpler model without the random effects.

Let $\alpha = (\beta', \sigma^2, \xi, \rho)'$ be the vector of parameters of the model (2.4) and let $\hat{\alpha}_0 = (\hat{\beta}'_0, \hat{\sigma}^2_0, \hat{\xi}_0, \hat{\rho}_0)'$ be the parameters of $\alpha$ replaced their corresponding ML estimates under the null model. The log-likelihood function of $\alpha$, given $Y_1, Y_2, \ldots, Y_n$, for the model (2.4) can be written as

$$l(\beta, \sigma^2, \xi, \rho) = \text{constant} - \frac{1}{2} \sum_{k=1}^{n} \log | \Sigma_k | - \frac{1}{2} \sum_{k=1}^{n} (Y_k - X_k \beta)' \Sigma_k^{-1} (Y_k - X_k \beta),$$

where $\Sigma_k = \text{cov}(Y_k) = \sigma^2 (\xi J_k + V_k)$. It is true that when evaluated at $\hat{\alpha}_0$, the score
vector $\frac{\partial \ell}{\partial \alpha}$ has all elements equal to zero except the derivative with respect to $\xi$ denoted by $\frac{\partial \ell}{\partial \xi}$ evaluated at $\hat{\alpha}_0$. The $(i,j)^{th}$ element of information matrix is $I_{ij} = E \left( -\frac{\partial^2 \ell}{\partial \alpha_i \partial \alpha'_j} \right)$, which has a form partitioned in accordance with $\alpha = (\beta, \sigma^2, \xi, \rho)$. Then the information matrix denoted by $I$ is given by

$$I = \begin{bmatrix}
I_{\beta\beta} & 0 & 0 & 0 \\
0 & I_{\sigma^2\sigma^2} & I_{\sigma^2\xi} & I_{\sigma^2\rho} \\
0 & I_{\xi\sigma^2} & I_{\xi\xi} & I_{\xi\rho} \\
0 & I_{\rho\sigma^2} & I_{\rho\xi} & I_{\rho\rho}
\end{bmatrix}$$

The score statistic denoted by $\lambda$, then takes the form

$$\lambda = \left( \frac{\partial \ell}{\partial \alpha} \right)_{\hat{\alpha}_0} \left( I^{-1} \right)_{\hat{\alpha}_0} \left( \frac{\partial \ell}{\partial \alpha} \right)_{\hat{\alpha}_0}$$  \hspace{1cm} (2.7)

which in the present context reduces to

$$\lambda = \left( \frac{\partial \ell}{\partial \xi} \right)_{\hat{\alpha}_0} \left( I_{\xi\xi}^{-1} \right)_{\hat{\alpha}_0} \left( \frac{\partial \ell}{\partial \xi} \right)_{\hat{\alpha}_0}$$  \hspace{1cm} (2.8)

where $I_{\xi\xi}^{-1}$ is the $(m+2,m+2)^{nd}$ element of the matrix $I^{-1}$. The score test statistic $\lambda$ is asymptotically distributed as chi-square with 1 degree of freedom under the null hypothesis. The second derivatives of $l(\theta^*)$ with respect to $\theta^* = (\sigma^2, \xi, \rho)$ have the form

$$\frac{\partial^2 \ell}{\partial \theta^*_i \partial \theta^*_j} = \frac{1}{2} \sum_{k=1}^{n} tr \left( \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \theta^*_j} \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \theta^*_i} \right) - \sum_{k=1}^{n} (Y_k - X_k\beta)^T \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \theta^*_j} \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \theta^*_i} (Y_k - X_k\beta).$$  \hspace{1cm} (2.9)
Therefore, it is easy to see that

\[ E(-\frac{\partial^2 \ell}{\partial \theta_i \partial \theta_j}) = \frac{1}{2} \sum_{k=1}^{n} \text{tr}\left( \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \theta_i} \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \theta_j} \right) \]  

(2.10)

The elements of the information matrix can be calculated using (2.10).

**Result 1:** With the usual notations, the elements of the information matrix are given by

(a) \[ E(-\frac{\partial^2 \ell}{\partial \sigma^4}) = \frac{1}{2} \sum_{k=1}^{n} \text{tr}\left( \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \sigma^2} \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \sigma^2} \right) \]

(b) \[ E(-\frac{\partial^2 \ell}{\partial \sigma^2 \partial \xi}) = \frac{1}{2} \sum_{k=1}^{n} \text{tr}\left( \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \sigma^2} \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \xi} \right) \]

(c) \[ E(-\frac{\partial^2 \ell}{\partial \sigma^2 \partial \rho}) = \frac{1}{2} \sum_{k=1}^{n} \text{tr}\left( \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \sigma^2} \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \rho} \right) \]

(d) \[ E(-\frac{\partial^2 \ell}{\partial \xi^2}) = \frac{1}{2} \sum_{k=1}^{n} \text{tr}\left( \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \xi} \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \xi} \right) \]

(e) \[ E(-\frac{\partial^2 \ell}{\partial \xi \partial \rho}) = \frac{1}{2} \sum_{k=1}^{n} \text{tr}\left( \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \xi} \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \rho} \right) \]

(f) \[ E(-\frac{\partial^2 \ell}{\partial \rho^2}) = \frac{1}{2} \sum_{k=1}^{n} \text{tr}\left( \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \rho} \Sigma_k^{-1} \frac{\partial \Sigma_k}{\partial \rho} \right) \]

**Proof:** Easy to see from (2.10)

**Result 2:** Under the null hypothesis, the expressions (a)-(f) reduce to the fol-
following: Define $V^* = \frac{\partial}{\partial \rho}(V_k)$

(a) $E\left(-\frac{\partial^2 \ell}{\partial \sigma^4}\right) = \frac{1}{2\sigma^4} \sum_{k=1}^{n} tr(I_k) = \frac{1}{2\sigma^4} \sum_{k=1}^{n} t_k$

(b) $E\left(-\frac{\partial^2 \ell}{\partial \sigma^2 \partial \xi}\right) = \frac{1}{2\sigma^2} \sum_{k=1}^{n} tr(V_k^{-1}J_k) = \frac{1}{2\sigma^2} \sum_{k=1}^{n} \frac{\rho^2(t_k - 2) - 2\rho(t_k - 1) + t_k}{2\sigma^2}$

(c) $E\left(-\frac{\partial^2 \ell}{\partial \sigma^2 \partial \rho}\right) = \frac{1}{2\sigma^2} \sum_{k=1}^{n} tr(V_k^{-1}V_k^*V_k^{-1}V_k^*) = \frac{1}{\sigma^2(1 - \rho^2)}$

(d) $E\left(-\frac{\partial^2 \ell}{\partial \xi^2}\right) = \frac{1}{2} \sum_{k=1}^{n} \sum_{k=1}^{n} tr(V_k^{-1}J_kV_k^{-1}J_k) = \frac{1}{2} \sum_{k=1}^{n} \frac{[\rho^2(t_k - 2) - 2\rho(t_k - 1) + t_k]^2}{2}$

(e) $E\left(-\frac{\partial^2 \ell}{\partial \xi \partial \rho}\right) = \frac{1}{2} \sum_{k=1}^{n} \sum_{k=1}^{n} tr(V_k^{-1}J_kV_k^{-1}V_k^*) = \frac{1}{2} \sum_{k=1}^{n} \frac{(t_k - 1) - \rho(t_k - 2)}{(1 - \rho^2)^2}$

Proof: We now give proof of only (e), as the rest similarly follow. Under the null hypothesis

$E\left(-\frac{\partial^2 \ell}{\partial \xi \partial \rho}\right) = \frac{1}{2} \sum_{k=1}^{n} tr(V_k^{-1}J_kV_k^{-1}V_k^*)$

Consider $tr(V_k^{-1}J_kV_k^{-1}V_k^*) = tr(V_k^{-1}1_kV_k^{-1}V_k^*)$

$V_k^{-1} = I + \rho^2 C_1 - \rho C_2$

$1_kV_k^{-1} = 1_k + \rho^2 1_k C_1 - \rho 1_k C_2$

$= 1_k + \rho^2 (0, 1, 1, ..., 1, 1, 0) - \rho (1, 2, 2, ..., 2, 2, 1)$

$= (1 - \rho, 1 - 2\rho + \rho^2, ..., 1 - 2\rho + \rho^2, 1 - \rho)$

$= (1 - \rho)(1, 1 - \rho, ..., 1 - \rho, 1)$ (2.11)

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Recall that $\mathbf{V}_k^* = \frac{\partial}{\partial \rho} (\mathbf{V}_k)$ and $\mathbf{V}_k$ is of order $t_k \times t_k$. For notational convenience, we carry out the further calculations by dropping the subscript $k$ of $t_k$. Let us now examine the elements of the matrix $\mathbf{V}^* = (v_{ij}^*), \text{ for } i, j = 1, ..., t$.

Case (i)

For $|i - j| = 0$

$$v_{ij}^* = \frac{2\rho}{(1 - \rho^2)^2}$$

Case (ii)

For $|i - j| > 0$

$$v_{ij}^* = \frac{(1 - \rho^2) |i - j| \rho^{|i-j|-1} + 2\rho^{|i-j|+1}}{(1 - \rho^2)^2}$$

Note that $\mathbf{V}^*$ is of the form

$$\mathbf{V}^* = \begin{bmatrix}
v_1 & v_2 & v_3 & \ldots & v_{t-1} & v_t \\
v_2 & v_1 & v_2 & \ldots & v_{t-2} & v_{t-1} \\
\vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\
v_{t-1} & v_{t-2} & v_{t-3} & \ldots & v_1 & v_2 \\
v_t & v_{t-1} & v_{t-2} & \ldots & v_2 & v_1 \\
\end{bmatrix} \tag{2.12}$$
Using (2.11) and (2.12) we can see that

\[
V^{-1} V' V^{-1} \mathbf{1} = \begin{bmatrix}
v_1 + (1 - \rho) \sum_{i=2}^{t-1} v_i + v_t \\
v_2 + (1 - \rho) \sum_{i=1}^{t-2} v_i + v_{t-1} \\
\vdots \\
v_{t-1} + (1 - \rho) \sum_{i=1}^{t-2} v_i + v_2 \\
v_t + (1 - \rho) \sum_{i=2}^{t-1} v_i + v_1
\end{bmatrix}
\]

Then

\[
1' V^{-1} V' V^{-1} \mathbf{1} = 2v_1 + 2v_t + 4(1 - \rho) \sum_{i=2}^{t-1} v_i + (1 - \rho)^2 (t - 2)v_1 \\
+ (1 - \rho)^2 \sum_{j=3}^{t-1} \sum_{i=2}^{j-1} v_i + (1 - \rho)^2 \sum_{j=2}^{t-1} \sum_{i=2}^{t-2} v_i
\]

(2.13)

It can be verified that

\[
\sum_{j=3}^{t-1} \sum_{i=2}^{j-1} v_i = \sum_{j=2}^{t-1} \sum_{i=2}^{t-j} v_i.
\]

Therefore, the above equation (2.13) becomes

\[
1' V^{-1} V' V^{-1} \mathbf{1} = 2v_1 + 2v_t + 4(1 - \rho) \sum_{i=2}^{t-1} v_i \\
+ (1 - \rho)^2 (t - 2)v_1 + 2 \sum_{j=2}^{t-1} \sum_{i=2}^{t-j} v_i.
\]

(2.14)
We now evaluate
\[ \sum_{i=2}^{i-1} v_i \quad \text{and} \quad \sum_{j=2}^{j-1} \sum_{i=2}^{i-1} v_i \]
in terms of the elements of \( V^* \).

The elements of \( V^* \) for \( i = 1, \ldots, t \) can be written in the form
\[ v_i = \rho^{i-2}[(i - 1) - (i - 3)\rho] \]

Now,
\[
\sum_{i=2}^{i-1} v_i = \sum_{i=2}^{i-1} [\rho^{i-2}(i - 1) - (i - 3)\rho] \\
= \sum_{i=2}^{i-1} i\rho^{i-2} - \sum_{i=2}^{i-1} \rho^{i-2} - \sum_{i=2}^{i-1} i\rho^i + 3\sum_{i=2}^{i-1} \rho^i \\
= \frac{2\rho^2 + \rho + 1}{1 - \rho} - \frac{\rho^{t-2}}{1 - \rho} (4\rho^2 + \rho - 1) - t(1 + \rho)\rho^{t-2} \tag{2.15}
\]

\[
\sum_{j=2}^{j-1} \sum_{i=2}^{i-1} v_i = \frac{(t - 2)(2\rho^2 + \rho + 1)}{(1 - \rho)} - \frac{(4\rho^2 + \rho - 1)(1 - \rho^{t-2})}{(1 - \rho^2)} \tag{2.16}
\]

\[ v_1 = 2\rho \tag{2.17} \]

\[ v_t = \rho^{t-2}[(t - 1) - (t - 3)\rho^2] \tag{2.18} \]

From equations (2.15), (2.16), (2.17) and (2.18) we have
\[ 1'V^{-1}V^{-1}V^{-1} = 2[(t - 1) - \rho(t - 2)]. \tag{2.19} \]

Now substituting \( t = t_k \) back in (2.19), we have
\[
E(-\frac{\partial^2 \ell}{\partial \xi \partial \rho}) = \frac{1}{2} \sum_{k=1}^{n} tr(V_k^{-1}J_kV_k^{-1}V_k^*) \quad = \sum_{k=1}^{n} (t_k - 1) - \rho(t_k - 2).
\]

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Similarly, we calculate the other elements of the information matrix.

Partition the information matrix

\[
I = \begin{bmatrix}
D & E \\
E' & J
\end{bmatrix}
\]

where

\[
D = \frac{\partial^2 \ell}{\partial \beta \partial \beta'}
\]

\[
E = 0_{m \times 3}
\]

\[
J = \begin{bmatrix}
\sum_{k=1}^{n} \frac{1}{2\sigma^4} A_1 \\
\sum_{k=1}^{n} \frac{\rho^2(t_k-2)-2\rho(t_k-1)+t_k}{2\sigma^2} \\
\sum_{k=1}^{n} \frac{n\rho}{\sigma^2(1-\rho^2)} \\
\end{bmatrix}
\]

Writing \( \sum_{k=1}^{n} t_i = A \) and \( \sum_{k=1}^{n} t_i^2 = B \), we have

\[
J = \begin{bmatrix}
\frac{1}{2\sigma^4} A \\
\frac{A(1-\rho)^2+2\rho n(1-\rho)}{2\sigma^2} \\
\frac{n\rho}{\sigma^2(1-\rho^2)} \\
\end{bmatrix}
\]

Recall that the score statistic for testing \( H_0 : \xi = 0 \) is given by

\[
\lambda = \left( \frac{\partial \ell}{\partial \xi} \right)_{\hat{\alpha}} \left( I_{\xi} \right)_{\hat{\alpha}} \left( \frac{\partial \ell}{\partial \xi} \right)_{\hat{\alpha}} ,
\]

(2.20)
where
\[
\left( \frac{\partial \ell}{\partial \xi} \right)_{\xi_0} = -\frac{1}{2} \sum_{k=1}^{n} 1_k' V^{-1} 1_k + \frac{1}{2\sigma^2} \sum_{k=1}^{n} (Y_k - X_k\beta)' V^{-1} J_k V^{-1} (Y_k - X_k\beta),
\]
\[
(I_{\xi_0}^*) = J_{22} - \begin{pmatrix} J_{21} & J_{23} \\ J_{31} & J_{33} \end{pmatrix}^{-1} \begin{pmatrix} J_{11} & J_{13} \\ J_{31} & J_{33} \end{pmatrix}
\]

### 2.2.1 A special case

We now consider a special case of the model (2.4) with $t_k = t$ for all $k$ and give the score test for testing $H_0: \xi = 0$. In this case it can be shown that

\[
J = \begin{bmatrix}
\frac{nt}{\sigma^2} & \frac{n[t^2-2t(t-1)+t]}{2\sigma^2} & \frac{n\rho}{\sigma^2(1-\rho^2)} \\
\frac{n[t^2-2t(t-1)+t]}{2\sigma^2} & n[t-1 - \rho(t-2)] & \frac{(t-1) - \rho^2(t-3)}{(1-\rho^2)^2} \\
\frac{n\rho}{\sigma^2(1-\rho^2)} & n[t-1 - \rho(t-2)] & \frac{(t-1) - \rho^2(t-3)}{(1-\rho^2)^2}
\end{bmatrix}
\]

Now the components of the score test statistic are given by

\[
\left( \frac{\partial \ell}{\partial \xi} \right)_{\xi_0} = -\frac{n}{2} 1' V^{-1} 1 + \frac{1}{2\sigma^2} \sum_{k=1}^{n} U_k' V^{-1} J V^{-1} U_k,
\]

\[
U_k = (Y_k - X_k\beta)
\]

and

\[
\left( I_{\xi_0}^* \right) = \frac{2}{n(1-\rho)^4(t-2)(t-1)}.
\]

Therefore the test statistic $\lambda$ is given by

\[
\lambda = \frac{n^2 \left( (1-\hat{\rho}_0)(t(1-\hat{\rho}_0)+2\hat{\rho}_0)) + \frac{1}{2\hat{\rho}_0} n(1-\hat{\rho}_0)^2 \sum_{k=1}^{n} (\sum_{i=1}^{t} u_{ik} - \hat{\rho}_0 \sum_{i=2}^{t} u_{ik})^2 \right)}{4(1-\hat{\rho}_0)^4(t-2)(t-1)}
\]

(2.21)

Here $u_{ik}$ is the $i^{th}$ component of the vector $u_k$, and

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\[ u_k = \left( \begin{array}{c} y_{1k} - X_k \hat{\beta}_0 \\ y_{2k} - X_k \hat{\beta}_0 \\ \vdots \\ y_{tk} - X_k \hat{\beta}_0 \end{array} \right) \]

vector of measurements on individual \( k \). In order to compute the score test statistic given by (2.21), we need \( \hat{\beta}_0, \hat{\sigma}^2 \) and \( \hat{\rho} \), estimates of \( \beta, \sigma^2 \) and \( \rho \) respectively under the null hypothesis \( H_0 : \xi = 0 \). We use the estimates obtained in the Appendix to evaluate the test statistic. Once the statistic is evaluated it can be used to test \( H_0 \) by comparing its value with the \( \chi^2 \) cutoff point.

Before proceeding with the analysis with the other components of the model, we provide a brief discussion of the Rao's score test with the other two asymptotic tests, the likelihood ratio test and the Wald's test.

A discussion of the three large sample tests

We give a brief discussion of different tools for hypothesis testing in large samples. Testing of hypotheses in large samples has a long history. Wilks (1938) introduced the likelihood ratio test, Wald (1943) introduced what is now called Wald's test, and Rao (1948) proposed the score test (now known as Rao's score test) for testing in large samples.
These three tests are known to have a common asymptotic distribution, which is chi-square with degrees of freedom equal to the dimension of the vector of parameters under the null hypothesis. In a textbook on econometrics Amemiya (1985) has discussed these three tests. Hall and Mathiason (1990) showed that these tests are not just asymptotically equivalent in law (i.e., have the same asymptotic distribution) but are asymptotically pointwise equivalent. That is, the test statistics are indeed approximately equal. This equivalence is proved by these authors considering Pitman-type local alternatives.

By simulation and theoretical investigations, it has been shown by several authors that there are no strict dominance relationships among the Wald’s, likelihood ratio and score test statistics. For example, see Amemiya (1985). However, Chandra and Mukherjee (1991) observed that Wald’s and Rao’s tests are not necessarily unbiased while the likelihood ratio test is. In this sense the three tests are not therefore strictly comparable. They suggested a Bartlett type modification for Wald’s and score tests to correct for bias. They found that the modified score test has better local properties than the likelihood ratio or modified Wald test.

The conclusions from the above discussion and the fact that the score test is easier to compute than either the likelihood ratio or the Wald’s test, prompted us to restrict our attention to the score test for testing $H_0 : \xi = 0$. 

26
2.3 Analysis under the reduced model

Suppose we accept the null hypothesis $H_0 : \xi = 0$. Then the model (2.4) is reduced to

$$Y_k = X_k\beta + \epsilon_k, \quad k = 1, \ldots, n$$  \hspace{1cm} (2.22)

That is, we now have a linear model with fixed effects and correlated errors. In the following sections, we discuss various methods of testing hypothesis of the form $H_0 : K\beta = 0$. In order to test the above linear hypothesis about the mean parameter vector $\beta$ we can use one of the following alternatives.

1. Generalized least squares method
2. Likelihood ratio method
3. Score test
4. Approximate $F$ test

While the methods (1) and (2) are commonly used, methods (3) and (4) are developed in this thesis.

2.3.1 Generalized least squares method

The generalized least squares estimator $\hat{\beta}$ is found by minimizing the quadratic form $\sum_{k=1}^{r} Q_k(\beta, V)$, where $Q_k(\beta, V) = (Y_k - X_k\beta)'V_k^{-1}(Y_k - X_k\beta)$. In the hypothetical case of known $V_k$, the solution of $\beta$ is given by

$$\hat{\beta} = (\sum_{k=1}^{r} X_k'V_k^{-1}X_k)^{-1}(\sum_{k=1}^{r} X_k'V_k^{-1}Y_k).$$  

When $V_k$ is unknown, the estimate of
\( \beta \) has the same form as above with \( V_k \) replaced by \( \hat{V}_k \), where \( \hat{V}_k \) is any consistent estimator of \( V_k \). The estimated covariance matrix of \( \hat{\beta} \) is \( \text{var}(\hat{\beta}) = \hat{V}_\beta = (\sum_{k=1}^{n} X'_k \hat{V}_k^{-1} X_k)^{-1} \). This is used for testing the hypothesis of the form \( K \beta = 0 \).

The statistic \( T_c = (K\hat{\beta})(KV_\beta K')^{-1}(K\hat{\beta}) \) approximately has a chi-square distribution with \( r \) degrees of freedom. This because as \( n \to \infty \), \( \hat{V} \to V \) in probability.

Thus under the null hypothesis, \( K\hat{\beta} \to N_r(0, KV_\beta K') \) and so \( T_c \to \chi^2_r \) in distribution.

### 2.3.2 Likelihood ratio method

Let \( \Theta = (\beta, \sigma^2, \rho) \) be the vector of unknown parameters of the model. Let \( L(\beta, \sigma^2, \rho) \) be the likelihood function obtained by evaluating the joint density of \( Y_1, Y_2, ..., Y_n \) at their specified values \( y_1, y_2, ..., y_n \). The likelihood ratio test for testing the linear hypothesis \( H_0 : K\beta = 0 \), where \( K \) is a \( r \times m \) matrix of full rank \( r \leq m \), rejects \( H_0 \) in favor of \( H_a \) if

\[
\Lambda = \frac{\max_{\theta \in \Theta} L(\beta, \sigma^2, \rho)}{\max_{\theta \in \Theta_0} L(\beta, \sigma^2, \rho)} < C
\]

with \( \Theta = (\beta, \sigma^2, \rho) \) and \( \Theta_0 = \{ (\beta, \sigma^2, \rho) : K\beta = 0 \} \).

Let \( \hat{\theta} = (\hat{\beta}, \hat{\sigma}^2, \hat{\rho}) \) be the MLE of \( \theta \in \Theta \) and \( \hat{\theta}_0 = (\hat{\beta}_0, \hat{\sigma}_0^2, \hat{\rho}_0) \) be the MLE of \( \theta \) under \( H_0 \), that is when \( K\beta = 0 \). The method of estimation of \( \theta \) under the entire parameter space \( \Theta \) and under the null hypothesis \( H_0 \) is discussed in the Appendix to this chapter.
Using these estimates, the likelihood ratio statistic reduces to

$$\Lambda = \left( \frac{\hat{\sigma}^2}{\sigma_0^2} \right)^{n_1} \left( \frac{1 - \hat{\rho}^2}{1 - \rho_0^2} \right)^{n_2}$$

In applications of the likelihood ratio method, we must obtain the sampling distribution of the likelihood ratio statistic $\Lambda$. However, it is well known that when the sample size is large, the sampling distribution of $-2\ln \Lambda$ is well approximated by a $\chi^2$ distribution with degrees of freedom $(\nu - \nu_0)$. Here the degrees of freedom $(\nu - \nu_0) = (\text{dimension of } \Theta) - (\text{dimension of } \Theta_0)$. Thus in the present situation $-2\ln \Lambda \sim \chi^2$ with $r$ degrees of freedom approximately.

### 2.3.3 Score test

Here we develop a score test for testing $H_0 : K\beta = 0$ vs. $H_a : K\beta \neq 0$. Let $\alpha = (\beta', \sigma^2, \rho)$ and let $\hat{\alpha}_0 = (\hat{\beta}', \hat{\sigma}^2, \hat{\rho})$ be the MLE's of $\alpha$ under the restriction of the null hypothesis. Let $l$ denote the loglikelihood function of $Y_1, Y_2, \ldots, Y_n$, which can be written as

$$l(\beta, \sigma^2, \rho) = \text{constant} - \frac{1}{2} \sum_{k=1}^{n} \log | \sigma^2 V_k |$$

$$- \frac{1}{2\sigma^2} \sum_{k=1}^{n} (Y_k - X_k\beta)'V_k^{-1}(Y_k - X_k\beta) \quad (2.23)$$
The components of the score vector are

\[
S = \begin{pmatrix}
\frac{\partial \ell}{\partial \beta} \\
\frac{\partial \ell}{\partial \sigma^2} \\
\frac{\partial \ell}{\partial \rho}
\end{pmatrix}
\]

\[
= \left( \frac{1}{\sigma^2} \sum_{k=1}^{n} (X_k'V_k^{-1}Y_k - X_k'V_k^{-1}X_k\beta) \right)
\]

\[
= -\frac{nt}{2\sigma^2} + \frac{1}{2\sigma^4} \sum_{k=1}^{n} (Y_k - X_k\beta)'V_k^{-1}(Y_k - X_k\beta)
\]

\[
= -\frac{n\rho}{1 - \rho^2} - \frac{1}{2\sigma^2} \sum_{k=1}^{n} (Y_k - X_k\beta)'(2\rho C_1 - C_2)(Y_k - X_k\beta)
\]

We need to evaluate the score vector \( S \) at \( \hat{a}_0 \), where \( \hat{a}_0 = (\hat{\beta}_0, \hat{\sigma}_0^2, \hat{\rho}_0) \), \( \hat{\beta}_0, \hat{\sigma}_0^2, \hat{\rho}_0 \)

being the estimates of \( \beta, \sigma^2 \), and \( \rho \) under the null model.

It is true that \( S_{\hat{a}_0} = \begin{pmatrix}
\frac{\partial \ell}{\partial \beta} |_{\hat{a}_0} \\
0 \\
0
\end{pmatrix} \)

Let the information matrix of \( \alpha = (\beta, \sigma^2, \rho) \) be partitioned as

\[
I = \begin{pmatrix}
I_{\beta\beta} & 0 & 0 \\
0 & I_{\sigma^2\sigma^2} & I_{\sigma^2\rho} \\
0 & I_{\rho\sigma^2} & I_{\rho\rho}
\end{pmatrix}
\]

where \( I_{\alpha_i\alpha_j} = E \left(-\frac{\partial^2 \ell}{\partial \alpha_i \partial \alpha_j}\right) \) for \( \alpha_i, \alpha_j = \beta, \sigma^2 \) and \( \rho \). The score statistic for \( H_0 : K\beta = 0 \) can be expressed in terms of the quadratic form as

\[
\Lambda = \left( \frac{\partial \ell}{\partial \beta} \right)' (I_{\beta\beta})^{-1} \left( \frac{\partial \ell}{\partial \beta} \right)
\]
where \( I_{\beta\beta} = -\frac{1}{\delta^2} \sum_{k=1}^{n} X'_k V_k^{-1} X_i \mid \beta_0 \). Estimation of \( \alpha \) under \( H_0 : K\beta = 0 \) is discussed in the Appendix. For large \( n \), the distribution of \( \Lambda \) is approximated by a \( \chi^2 \)-distribution with \( r \) degrees of freedom.

### 2.3.4 Approximate \( F \) test

We now discuss a new approach to testing hypotheses of the form \( H_0 : K\beta = 0 \) under the reduced model. Consider the model

\[
Y_k = X_k \beta + \epsilon_k \quad k = 1, \ldots, n
\]

We now require \( t_k = t \) for all \( k \) to develop this method. If we assume that \( \text{cov}(\epsilon_k) = \sigma^2 I \), to test the above hypothesis we have the usual \( F \) statistic given by

\[
F = \frac{(K\hat{\beta})'(K(X'X)^{-1}K')^{-1}(K\hat{\beta})}{r\hat{\sigma}^2}
\]

with \( r \) and \( n - r \) degrees of freedom, \( r \) being the rank of \( K \). However, we have dependent errors with \( \text{cov}(\epsilon_k) = V = \frac{\sigma^2}{1 - \rho^{|i-j|}} \) instead of \( \text{cov}(\epsilon_k) = \sigma^2 I \). In this situation the natural question that arises is about the distribution of the \( F \) statistic. Several authors including Box (1954) and Geisser and Greenhouse (1958) have tried to provide an answer to this question. We know that the sum of squares due to the hypothesis can be expressed as a quadratic form as \( Y'HY \). Let the sum of squares due to error be expressed as \( Y'EY \). In general to use the ratio \( \frac{Y'HY}{Y'EY} \) as a test statistic, we require the two quadratic forms to be independent.
Stated in other words, we require $HVE = 0$. This condition may not hold true in general. However in some special situations, like the one-way ANOVA model, split plot designs, the above condition holds true for all the standard hypothesis. The distributions of the $F$ statistics are considered in detail for the split plot design in Chapter 1. We now derive the expressions for the correction factor $e$ for the degrees of freedom under this autoregressive structure for $V$.

We may recall from Chapter 1 that

$$
e = (t - 1)^{-1} \frac{[tr(V - \frac{1}{t}JV)]^2}{tr(V - \frac{1}{t}JV)^2}.$$ 

For our choice of $V$

$$tr(V - \frac{1}{t}JV) = tr(V) - \frac{1}{t}tr(JV)$$

$$tr(V - \frac{1}{t}JV)^2 = tr(V) - \frac{2}{t}tr(VJV) + \frac{1}{t^2}tr(JVJV)$$

where

$$tr(V) = \frac{\sigma^2 t}{1 - \rho^2}$$
$$tr(JV) = \frac{\sigma^2[t(1 - \rho^2) - 2\rho(1 - \rho^4)]}{(1 - \rho^2)(1 - \rho)^2}$$
$$tr(VV) = \frac{\sigma^4}{(1 - \rho^2)^4} [t(1-\rho^4) - 2\rho^2(1 - \rho^2)]$$
$$tr(VJV) = \frac{\sigma^4 [t(1 + \rho)^2 + 2t\rho^{-1} + 2\rho^2 1 - \frac{\rho^2}{1 - \rho^2} - 4\rho(1 + \rho) \frac{1 - \rho}{1 - \rho^2}]}{(1 - \rho^2)^2(1 - \rho)^2}$$
$$tr(JVJV) = \left[ \frac{\sigma^2[t(1 - \rho^2) - 2\rho(1 - \rho^4)]}{(1 - \rho^2)(1 - \rho)^2} \right]^2$$

In order to evaluate $e$ we need the estimates of $\sigma^2$ and $\rho$. The estimation of these is described in the Appendix.
2.4 Appendix

2.4.1 Model and estimators

Consider the model

\[ Y_k = X_k \beta + \epsilon_k \quad k = 1, \ldots, n, \]

where \( \epsilon_k \) is a \( t \times 1 \) vector, \( X_k \) is the \( t \times m \) design matrix for the mean vector and \( Y_k \) is a \( t \times 1 \) vector of dependent variables on individual \( k \). It is assumed that \( E(\epsilon_k) = 0 \) and \( \text{cov}(\epsilon_k) = \sigma^2 V \), where

\[
V = \frac{1}{1 - \rho^2} \begin{bmatrix}
1 & \rho & \rho^2 & \ldots & \rho^{t-1} \\
\rho & 1 & \rho & \ldots & \rho^{t-2} \\
& \ddots & \ddots & \ddots & \ddots \\
& & \rho^{t-1} & \rho^{t-2} & \rho^{t-3} & \ldots & 1
\end{bmatrix}
\]

It can be verified that \( |V| = \frac{1}{1 - \rho^2} \) and \( V^{-1} = I + \rho^2 C_1 - \rho^2 C_2 \) where

\[
C_1 = \begin{bmatrix}
0 & 0 & \ldots & .0 & 0 \\
0 & 1 & \ldots & .0 & 0 \\
& \ddots & \ddots & \ddots & \ddots \\
& & \ddots & \ddots & \ddots \\
0 & 0 & \ldots & 1 & 0 \\
0 & 0 & \ldots & 0 & 0
\end{bmatrix}
\]
The likelihood function for the model is given by

\[ l = (\frac{1}{2\pi})^{\frac{n}{2}} \prod_{k=1}^{n} |\sigma^2 V|^{-\frac{1}{2}} \exp\{-\frac{1}{2\sigma^2} \sum_{k=1}^{n} (Y_k - X_k\beta)'V^{-1}(Y_k - X_k\beta)\} \quad (2.24) \]

and the loglikelihood function is given by

\[ l(\beta, \sigma^2, \rho) = \text{constant} - \frac{nt}{2}\ln\sigma^2 + \frac{n}{2}\ln(1 - \rho^2) \]

\[ - \frac{1}{2\sigma^2}\sum_{k=1}^{n} (Y_k - X_k\beta)'(I + \rho^2C_1 - \rho C_2)(Y_k - X_k\beta). \quad (2.25) \]

The equation (2.25) can be written as

\[ l(\beta, \sigma^2, \rho) = \text{constant} - \frac{nt}{2}\ln\sigma^2 + \frac{n}{2}\ln(1 - \rho^2) - \frac{A_1}{2\sigma^2} - \frac{\rho^2A_2}{2\sigma^2} + \frac{\rho A_3}{2\sigma^2} \quad (2.26) \]

where

\[ A_1 = \sum_{k=1}^{n} (Y_k - X_k\beta)'(Y_k - X_k\beta), \]

\[ A_2 = \sum_{k=1}^{n} (Y_k - X_k\beta)'C_1(Y_k - X_k\beta), \]

\[ A_3 = \sum_{k=1}^{n} (Y_k - X_k\beta)'C_2(Y_k - X_k\beta). \]
The maximum likelihood method yields the following non-linear maximum likelihood (ML) equations which can be solved iteratively for the unknown parameters $\beta, \sigma^2$ and $\rho$ respectively.

\[
\hat{\beta} = \frac{\left[ \sum_{k=1}^{n} X'_k X_k + \rho^2 \sum_{k=1}^{n} X'_k C_1 X_k - \rho \sum_{k=1}^{n} X'_k C_2 X_k \right]^{-1}}{\sum_{k=1}^{n} X'_k Y_k + \rho^2 \sum_{k=1}^{n} X'_k C_1 Y_k + \rho \sum_{k=1}^{n} X'_k C_2 Y_k} \quad (2.27)
\]

\[
\hat{\sigma}^2 = \frac{1}{nt} \left[ A_1 + \rho^2 A_2 - \rho A_3 \right] \quad (2.28)
\]

\[
2(t - 1) A_2 \rho^3 - (t - 2) A_3 \rho^2 - 2(A_1 + t A_2) \rho + t A_3 = 0. \quad (2.29)
\]

We note that only the equations (2.27) and (2.29) have to be solved iteratively.

### 2.4.2 Estimation under the null hypothesis $K \beta = 0$

Consider the model

\[
Y_k = X_k \beta + \epsilon_k \quad k = 1, \ldots, n \quad (2.30)
\]

We need to estimate the parameters of the model under $H_0 : K \beta = 0$, where $K$ is a matrix of order $r \times m$ with $\text{rank}(K) = r$. To do that choose a matrix $L$ of order $m - r \times m$ such that $M = \begin{pmatrix} K \\ L \end{pmatrix}$ is a nonsingular matrix, and $MM^{-1} = I$. The model (2.30) can be written as

\[
Y_k = X_k M^{-1} M \beta + \epsilon
\]
\[ \begin{align*}
  &= X_k M^{-1} \begin{pmatrix} K \\ L \end{pmatrix} \beta + \epsilon \\
  &= (X_k^{(1)} | X_k^{(2)}) \begin{pmatrix} K \beta \\ L \beta \end{pmatrix} + \epsilon \quad \text{(writing } X_k M^{-1} = (X_k^{(1)} | X_k^{(2)})) \\
  &= X_k^{(1)} K \beta + X_k^{(2)} L \beta + \epsilon
\end{align*} \]

Under \( H_0 \) the model reduces to

\[ Y_k = Z_k \beta^* + \epsilon, \]

where \( Z_k = X_k^{(2)} \) and \( \beta^* = L \beta \). Now we need the estimates of \( \beta^*, \sigma^2 \) and \( \rho \). The likelihood function under the null hypothesis yields the following non-linear maximum likelihood (ML) equations which can be solved for the unknown parameters \( \beta^*, \rho \) and \( \sigma^2 \). The equations are:

\[ \hat{\beta}^* = \left[ \sum_{k=1}^{n} Z_k' \hat{Z}_k + \hat{\rho}^2 \sum_{k=1}^{n} Z_k' C_1 Z_k - \hat{\rho} \sum_{k=1}^{n} Z_k' C_2 Z_k \right]^{-1} \left[ \sum_{k=1}^{n} Z_k Y_k + \hat{\rho}^2 \sum_{k=1}^{n} Z_k' C_1 Y_k + \hat{\rho} \sum_{k=1}^{n} Z_k' C_2 Y_k \right] \]

\[ \hat{\sigma}^2 = \frac{1}{nt} \left[ B_1 + \hat{\rho}^2 B_2 - \hat{\rho} B_3 \right] \]

\[ 2(t - 1)B_2 \rho^3 - (t - 2)B_3 \rho^2 - 2(B_1 + t B_2) \rho + t B_3 = 0 \]

where

\[ B_1 = \sum_{k=1}^{n} (Y_k - Z_k \beta)' (Y_k - Z_k \beta), \]

36
\[ B_2 = \sum_{k=1}^{n} (Y_k - Z_k \beta)' C_1(Y_k - Z_k \beta), \]
\[ B_3 = \sum_{k=1}^{n} (Y_k - Z_k \beta)' C_2(Y_k - Z_k \beta). \]
Chapter 3

Analysis of Repeated Measures With Covariates

In this chapter we study methods of analyzing repeated measures data in the presence of covariates. In the medical field we often come across repeated measures data along with covariates that influence the response variable. It is important to conduct the analysis of repeated measures data to account for covariate effects, when data contain covariates that might affect the analysis. In clinical trials, baseline measurement can be thought of as a useful covariate for analyzing response patterns at successive visits. In the following we describe two situations, cited by Patel (1986), where one or more covariates occur naturally. In therapies for the treatment of chronic stable angina, treadmill walking time (a covariate) is recorded just before the administration of a dose, and then at some post-dose time. The
effectiveness of the visit doses is evaluated relative to the corresponding baseline walking times. In another situation the effectiveness of a drug in the treatment of atherosclerosis is probably influenced by factors such as diet, exercise and smoking (covariates). If the influencing factors are measured we need to construct a model which makes use of this information.

The situations described above differ in the nature of covariate information available. In the first example, the value of the covariate is specific to the subject and does not vary with time. However, in the second situation the value of each covariate is being measured at each time point along with the response variable. Thus depending on the type of covariate information available, we have

- Covariates fixed over time
- Covariates changing over time, or time varying covariates

In a repeated measures data analysis the main problems of interest are

(i) To test for the interaction between time and the group effects (parallel profiles)

(ii) To compare the profiles of the means (over time) of different groups (coincident profiles)

(iii) To test the difference between the means of the response variables at different time points (horizontal profiles)

In this chapter, we review the methods of analyzing repeated measures data in the presence of covariates. In the section 3.1 we discuss the methods of solving these
problems when the covariates are fixed over time. In section 3.2 we review various methods of analyzing the repeated measures data with time varying covariates. These methods are illustrated using numerical examples and the SAS software. The SAS programs are also provided for convenience.

3.1 Covariates fixed over time

Depending on the nature of the data and other considerations about the covariance structures of the repeated measures, one of the following two approaches are usually adopted in practice.

3.1.1 The multivariate approach

Let \( Y_{ij} = (y_{ij1}, \ldots, y_{ijt}) \) be a vector of \( t \) measurements on the \( j^{th} \) individual from the \( i^{th} \) group. We also have a set of \( m \) covariates, \( x_{1ij}, \ldots, x_{mij} \), on each individual. Let us assume \( m = 1 \) for simplicity of presentation and denote the value of the covariate by \( x_{ij} \). We have the following model for vector \( Y_{ij} \),

\[
Y_{ij} = \mu + \alpha_i + \lambda_i x_{ij} + \epsilon_{ij}, \quad j = 1, \ldots, n_i, \quad i = 1, \ldots, g
\]

where \( \mu, \alpha_i, \) and \( \lambda_i \) are all \( t \times 1 \) vectors, and \( \epsilon_{ij} \sim N_t(0, \mathbf{V}) \), \( \mathbf{V} \) a positive definite matrix.

We can rewrite (3.1) as

\[
Y_{ij}' = \mu' + \alpha_i' + x_{ij} \lambda_i' + \epsilon_{ij}'
\]
\[ X' = X'B + \epsilon_{ij}, \]

where \( X' \) is the \( 1 \times (2g + 1) \) vector \((1, 0, ..., 0, 1, 0, ..., 0, x_{ij}, ..., 0)\) and \( B \) is the \((2g + 1) \times t \) matrix such that \( B' = (\mu, \alpha_1, ..., \alpha_g, \lambda_1, ..., \lambda_g) \).

Next define \( Y_{nxt} = (Y_{11}, ..., Y_{1n_1}, Y_{21}, ..., Y_{2n_2}, ..., Y_{gn_p})', n = \sum_{i=1}^{g} n_i \) and \( X_{n \times (2g+1)} = (X_{11}, ..., X_{1n_1}, X_{21}, ..., X_{2n_2}, ..., X_{gn_p})' \). Then

\[ Y = XB + E, \quad (3.2) \]

where \( E \) is a \( n \times t \) matrix of errors. The rows of \( E \) are assumed to be independent and each distributed as \( N_i(0, \Sigma) \). Under this model, our testing problems (i)-(iii) can be tested using the general linear hypothesis \( H_0 : LB = 0 \). Any standard statistical package like SAS, can be utilized to do the above MANOVA.

### 3.1.2 The univariate approach

An alternative approach to the method described above is to adopt the analysis of a split plot design, but with covariates. Let \( y_{ijk} \) be an observed value of the response variable on the \( j^{th} \) subject from the \( i^{th} \) group at the \( k^{th} \) occasion. Then we have the model

\[ y_{ijk} = \mu + \alpha_i + \beta_k + (\alpha\beta)_{ik} + \lambda_k z_{ij} + \gamma_{j(i)} + \epsilon_{ijk}, \quad (3.3) \]

\[ k = 1, ..., t, \quad j = 1, ..., n_i, \quad i = 1, ..., g \]

with \( \epsilon_{ijk} \sim N(0, \sigma^2), \gamma_{j(i)} \sim N(0, \sigma^2_\gamma) \). These assumptions on the random effects amounts to having an equicorrelation structure for \( \text{cov}(Y_{ij}) = \Sigma = \sigma^2 V(\rho) \), that is,
\( V(p) = (1 - p)I + p11' \), for the correlation matrix of an individual. Hence we can apply the usual linear model theory. For that, write \( y' = (Y_{11}, \ldots, Y_{1n_1}, \ldots, Y_{21}, \ldots, Y_{g n_g}) \),

with \( \beta' = (\mu, \alpha_1, \ldots, \alpha_g, \beta_1, \ldots, \beta_t, (\alpha\beta)_{11}, \ldots, (\alpha\beta)_{g1}, \lambda_1, \ldots, \lambda_g) \). Then we have

\[
y = X\beta + \epsilon,
\]

where \( X \) is the appropriately chosen design matrix and \( \epsilon \sim N_n(0, \Omega), \Omega = I_n \otimes \sigma^2 V(p) \), \( n = \sum_{i=1}^g n_i \). Under this model, testing of various hypotheses can be easily carried out by testing \( H_0 : L\beta = 0 \) for specific choices of \( L \). Again softwares like SAS, can be utilized to do this testing.

The two approaches (the multivariate and the univariate) are different on the following two accounts:

(a) On one hand we assume a completely arbitrary correlation structure for the multivariate approach, and on the other hand we use a very simple equicorrelation structure for the univariate approach.

(b) We use the theory of multivariate statistical analysis for the first approach, and the ANOVA F-tests for the second.

Both of these approaches fail when the correlation structure is not of one of the two extremes considered above. To overcome the shortcomings of these methods the following alternative linear model approach can be adopted.
Suppose the linear model

\[ y = X\beta + \epsilon, \quad (3.5) \]

is used to model the repeated measures data with \( Y \) as a \( nt \times 1 \) vector of observations observed at \( t \) occasions on \( n = \sum_{i=1}^{g} n_i \) patients, \( X \) as a \( nt \times m \) design matrix and \( \beta \) as a \( m \times 1 \) vector of parameters. Suppose \( \text{cov} (\epsilon) = \Omega = I_n \otimes V_{1 \times 1} \), where \( V = V(\theta_1, ..., \theta_k) \), for some unknown parameters \( \theta_1, ..., \theta_k \).

Assuming multivariate normality for \( \epsilon \), the maximum likelihood estimates of \( \beta, \theta_1, ..., \theta_k \) can be derived. Then the likelihood ratio test, for testing any linear hypothesis of interest \( H_0 : L\beta = 0 \), can be utilized appealing to the asymptotic distribution of the test.

Another route taken is through generalized least squares estimates (GLE). For a known \( V \), since \( V \) is positive definite, GLE of \( \beta \), \( \hat{\beta} = (X'V^{-1}X)^{-1}X'V^{-1}Y \). Using this \( \hat{\beta} \) the ANOVA test for testing \( H_0 : L\beta = 0 \) can be derived. But \( V \) is usually unknown. Using some consistent estimators of \( \theta_1, ..., \theta_k \), \( V \) is estimated and the estimate \( \hat{V} = V(\theta_1, ..., \theta_k) \) is used in place of \( V \) in the formula for \( \hat{\beta} \) and the same ANOVA tests can still be used. But now the distribution of the test statistic will have to be approximated.

The maximum likelihood approach, when \( V \) is known, yields the same estimate of \( \beta \) as the GLE. Further, the likelihood ratio test for testing \( L\beta = 0 \) will still hold through if a \( \sqrt{n} \)-consistent estimate of \( V \), instead of the maximum likelihood
estimate, is utilized. This fact is useful since in many problems with structured covariances, finding the maximum likelihood estimates of \( \theta_1, \ldots, \theta_k \) is generally hard, if not impossible. However, consistent estimates of \( \theta_1, \ldots, \theta_k \) and hence of \( V \) can be obtained easily.

Another serious drawback of the multivariate and ANOVA methods is that they cannot be applied when data are unbalanced, that is, when different number of observations are available on different individuals. Further, the multivariate tests have been known to be less powerful, when the variance covariance matrix of the observation vector has a certain structure. Thus, either if data are unbalanced or the covariance matrix has a structure other than the equicorrelation structure, the maximum likelihood approach or the two stage GLE approach seems to be the right approach. Further, this general approach can be utilized when the covariates are time varying. Modification of this approach to accommodate unbalanced data will be discussed in section 3.2.4.

3.1.3 Example 1

In this section, we illustrate methods described in section 3.1.1 and 3.1.2 to analyze repeated measures data with fixed covariates. The data set in Table 1 is courtesy of Center for Pediatric Research, Eastern Virginia Medical School, Norfolk, Virginia and it has been discussed and analyzed by Pickering, Marrow, Herrera et al., (1995). The purpose of the study was to see the effect of maternal immunization on the
serum anti-rotavirus antibody concentration. Thirty two women were random­
ized into three groups and received a single oral dose of either rhesus rotavirus
monovalent reassortant vaccine (10^4 pfu), rhesus rotavirus tetravalent vaccine (10^4
pfu), or placebo. Measurements on the antibody concentration were taken. One
measurement was taken before the vaccination and six other measurements were
collected after a week, one month, two months, three months, four months and five
months after the vaccination. Measurement taken before the vaccination is consid­
ered a baseline measurement or a covariate. Since there are many missing values
for months four and five we have omitted them from our analysis.

Prior to carrying out statistical analysis it is important to verify if the data
satisfy the assumption of multivariate normality. This was done using tests based
on Mardia’s multivariate skewness and kurtosis measures. See Khattree and Naik
(1995) for SAS programs for computing these tests. For this data set with its
small p-values, (p-value(skewness) = 0 and p-value(kurtosis) = 0), there is an
indication that the assumption of multivariate normality is violated. We make a
log-transformation on these data, and then again test for multivariate normality.
We found that p-values for the tests based on skewness and kurtosis respectively are
0.1190 and 0.4180. This indicates that the log transformed data can be assumed
to have come from a multivariate normal distribution. Consequently, we apply our
methods to the log-transformed data.

Analysis using multivariate technique. Here we present our analysis using

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multivariate technique. The following considerations are in order.

1. The effect of covariates may be different in various treatment groups. This can be tested by testing for statistical significance of the interactions of the treatment with covariates.

2. Covariates may influence the measurements taken at different time points very differently. This can be tested by examining the significance of the interaction between covariates and the time factor.

We summarize the results of the SAS output in the following table:

<table>
<thead>
<tr>
<th>Hypothesis</th>
<th>Wilks' $\lambda$</th>
<th>Num DOF</th>
<th>Den DOF</th>
<th>p-value</th>
</tr>
</thead>
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It can be observed that the multivariate tests show that the effect of covariate over different treatment groups do not differ significantly ($p$-value=0.3741). However, the effect of the covariate itself is significant ($p$-value=0.0040). Therefore, it is important to include the covariate in the model. We also notice that there is no overall group effect.

Next we would like to examine whether the covariate influences the measurements taken at different time points differently. This can be achieved using REPEATED statement of SAS. The results are summarized in the following table:
Hypothesis       Wilks' $\lambda$  Num DOF  Den DOF  p-value

Time*Covariate  0.62301  3       17       0.0408

We see that the covariate does affect the measurements taken at different time points differently if 0.05 level of significance is used. However, if the level of significance is assumed to be 0.01 then the conclusion would be that the covariate does not affect differently.

Analysis using univariate approach. We present our results of the analysis using univariate model in the following table:

<table>
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<tr>
<th>Hypothesis</th>
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<th>Den DOF</th>
<th>$F$ value</th>
<th>p-value</th>
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</thead>
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We can see that the univariate tests indicate that the groups do not differ significantly ($p$-value=0.8099). The effect of the time factor is significant at 5% level of significance ($p$-value=0.0135). The interaction between time and group is
not significant ($p$-value=0.2489). We also see that the covariate does not affect different vaccination groups differently ($p$-value=0.7381).

### 3.2 Time varying covariates

As we mentioned in the introduction, in many repeated measures study, the covariates vary over time. Again assuming, for simplicity, that there is only one covariate, the data in this case can be represented by $y_{ijk}$, $x_{ijk}$, $k = 1, ..., t$, $j = 1, ..., n_i$, $i = 1, ..., g$. Analysis of these data seems to be a challenging problem. Only recently attempts have been made, in the literature, to analyze these data. We review these methods below.

#### 3.2.1 The multivariate approach

Patel (1986) proposed the following multivariate model for analyzing repeated measures designs with time varying covariates. Let $Y'_{ij}$ be a $1 \times t$ vector of response variables and $X'_{ij}$ be a $1 \times t$ vector of covariates, both taken over $t$ occasions on an individual. Next, define $Y_{nxt} = (Y_{11}, ..., Y_{1n_1}, Y_{21}, ..., Y_{2n_2}, ..., Y_{gn_g})'$, $n = \sum_{i=1}^{g} n_i$ and $X_{nxt} = (X_{11}, ..., X_{1n_1}, X_{21}, ..., X_{2n_2}, ..., X_{gn_g})'$. Then the following model

$$Y_{nxt} = A_{nxt}\xi_{mxt} + X_{nxt}\Gamma_{txt} + E_{nxt},$$

where $A$ is a design matrix, $\xi$ is a matrix of unknown parameters, $\Gamma$ is a diagonal matrix with the unknown diagonal elements, $\gamma_1, ..., \gamma_t$, and $E$ is a $n \times t$ error matrix,
can be utilized to represent the time varying covariates. Note that the model (3.6) is different from the usual multivariate analysis of covariance model in the sense that its parameter matrix $\Gamma$ is known (to be zero) except the diagonal entries. This makes it hard to handle the analysis of this model in a routine way.

As usual rows of $E$ are assumed to be independently distributed with a common multivariate normal distribution with a zero mean vector and a $t \times t$ covariance matrix $V$. Patel (1986) has provided the maximum likelihood estimators of the parameters. He has also provided an iterative algorithm describing the computation of the unknown parameters and the likelihood ratio test for any general linear hypothesis of the form $L_{k \times m} \xi_{m \times 1} M_{t \times e} = 0$.

The likelihood function under model (3.6) is

$$L = (2\pi)^{-\frac{1}{2}nt}|V|^{-\frac{1}{2}t}exp \left(-\frac{1}{2}tr(V^{-1}R)\right),$$

where $R = (Y - A\xi - X\Gamma)'(Y - A\xi - X\Gamma)$. The maximum likelihood estimators of $\xi$ and $\Gamma$ are obtained by minimizing $\phi = ln |R|$. The idea is that, if $\xi$ and $\Gamma$ are known then the MLE of $V$ would be $\frac{R}{n}$. In that case the log-likelihood function reduces to $ln |R|$ besides a constant. Hence the procedure iterates between $\hat{V} = \frac{\hat{R}}{n}$ and minimizing $ln |R|$ with respect to $\xi$ and $\Gamma$, $\hat{R}$ being $R$ when $\xi$ and $\Gamma$ are replaced by their estimates, say $\hat{\xi}$ and $\hat{\Gamma}$.

Note that

$$\frac{\partial \phi}{\partial \xi} = 0 \quad \Rightarrow \quad \hat{\xi} = (A'A)^{-1}A'(Y - X\hat{\Gamma})$$

(3.8)
and \( \frac{\partial \phi}{\partial \Gamma} = -2\text{diag}\{R^{-1}X'(Y - A\hat{\xi} - X\hat{\Gamma})\} = 0 \)

implies

\[
\text{diag}\{\hat{R}^{-1}X'(I - A(A'A)^{-1}A')Y\} = \text{diag}\{\hat{R}^{-1}X'X\hat{\Gamma}\}. \tag{3.9}
\]

Patel (1986) has suggested the following iterative steps to solve the ML equations:

Step 1. Taking \( R = I \), compute \( \hat{\Gamma} \) from (3.9). This estimate of \( \Gamma \) is same as that obtained by minimizing \( \text{tr}(R) \) w.r.t. \( \Gamma \).

Step 2. Compute \( \hat{\xi} \) from (3.9)

Step 3. Compute \( \hat{R} \) using \( \hat{\xi} \) and \( \hat{\Gamma} \). Then compute \( \hat{\phi} = \log |\hat{R}| \).

Step 4. Compute a revised estimate of \( \Gamma \) from (3.9) using \( \hat{R} \) obtained in step 3.

Step 5. Repeat steps 2, 3 and 4 until the absolute difference between two successive values of \( \hat{\phi} \) is less than a pre-determined number.

Patel (1986) has also given similar algorithm for computing the ML estimates under the null hypothesis \( H_0 : L\xi M = 0 \). Using these MLE's the likelihood ratio test for testing \( H_0 \) can be obtained. Recall that the likelihood function under model (3.6) is given by

\[
L = (2\pi)^{-\frac{1}{2}nt}|V|^{-\frac{1}{2}n}\exp\left[-\frac{1}{2}\text{tr}(V^{-1}R)\right].
\]
We compute the maximum of $L$ as

$$max \ L = (2\pi)^{-\frac{1}{2}nt} |\hat{\mathbf{V}}|^{-\frac{1}{2}n} \exp(-\frac{1}{2}nt), \tag{3.10}$$

where $\hat{\mathbf{V}} = \hat{\mathbf{R}}/n$. Similarly, writing the maximum likelihood estimator of $\mathbf{V}$ under $H_0$ as $\hat{\mathbf{V}} = \hat{\mathbf{R}}/n$, we get the maximum of $L$ under $H_0 : L \xi M = 0$ as

$$max \ L_{H_0} = (2\pi)^{-\frac{1}{2}nt} |\hat{\mathbf{V}}|^{-\frac{1}{2}n} \exp(-\frac{1}{2}nt). \tag{3.11}$$

From (3.10) and (3.11) the likelihood ratio test for testing $H_0$ is to reject the null hypothesis if

$$\lambda = |\hat{\mathbf{V}} \hat{\mathbf{V}}^{-1}|^{-\frac{1}{2}n} \leq C_\alpha, \tag{3.12}$$

where $C_\alpha$ is a constant to be determined satisfying $pr(\lambda \leq C_\alpha | H_0) = \alpha$. We use the standard result, that is, $-2 \log \lambda$ has asymptotically a chi-squared distribution with $mt - bc$ degrees of freedom, to test the hypothesis.

### 3.2.2 Example 2

In this section we illustrate the use of Patel’s algorithm to analyze repeated measures data with time varying covariates. We use data given in Table 2, courtesy of Dr. Barbara Hargrave of Biological Sciences department, Old Dominion University. The effects of phenylephrine induced increase in arterial pressure on the secretion of atrial natriuretic peptide (ANP) in the ovine fetus are studied by Hargrave and Castle (1995). In this study 16 chronically cannulated fetal sheeps were divided into
two groups. Arterial pressure was increased by infusing phenylephrine to the fetus from each of the two groups. Systematic mean arterial pressure (MAP), plasma ANP concentrations and plasma renin activity (PRA) were measured at three time points (5 min, 15 min, and 30 min) after infusion. We take PRA as the response variable and MAP as time varying covariate. Here we have \( g=2, t=3, n_1=6 \) and \( n_2=10 \) and one covariate. We illustrate the computation of \( \lambda \) given by (3.12) for testing various hypotheses by choosing different \( L \) and \( M \).

Let \( L = I_{2 \times 2} \) and

\[
M = \begin{bmatrix}
1 & 1 \\
-1 & 0 \\
0 & -1
\end{bmatrix}.
\]

Then the hypothesis of possible differences between different time points can be tested using the hypothesis \( L \xi M = 0 \). The ML estimates of \( n \Sigma \), under the nonnull and null hypotheses respectively are

\[
\hat{R} = \begin{bmatrix}
19807.802 & -4186.271 & 4663.295 \\
-4186.271 & 48285.522 & 30775.957 \\
4663.295 & 30775.957 & 24974.933
\end{bmatrix}
\]

and

\[
\tilde{R} = \begin{bmatrix}
20212.801 & -4704.252 & 5694.256 \\
-4704.252 & 48330.394 & 30810.573 \\
5694.256 & 30810.573 & 28866.521
\end{bmatrix}.
\]
Table 3.2: Plasma renin activity data

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<td>20</td>
<td>5</td>
<td>-17</td>
<td>13</td>
<td>-17</td>
</tr>
<tr>
<td>old</td>
<td>20</td>
<td>15</td>
<td>-38</td>
<td>54</td>
<td>7.7</td>
</tr>
<tr>
<td>old</td>
<td>20</td>
<td>30</td>
<td>50</td>
<td>205</td>
<td>0.2</td>
</tr>
</tbody>
</table>

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The value of $\lambda=0.0025$ and $-2\ln\lambda=11.9920$. Similarly we can test other hypotheses of interest with the appropriate choices of $L$ and $M$. We summarize our results below:

<table>
<thead>
<tr>
<th>Hypothesis</th>
<th>$\lambda$</th>
<th>D.O.F</th>
<th>Approximate $\chi^2$ value</th>
<th>p-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Group</td>
<td>0.1042</td>
<td>2</td>
<td>4.5225</td>
<td>0.1042</td>
</tr>
<tr>
<td>Time</td>
<td>0.0025</td>
<td>2</td>
<td>11.9920</td>
<td>0.0025</td>
</tr>
<tr>
<td>Time*Group</td>
<td>0.5993</td>
<td>2</td>
<td>1.0241</td>
<td>0.5993</td>
</tr>
</tbody>
</table>

From the above table we see that only the time effect is significant.

### 3.2.3 SUR model approach

Verbyla (1988) pointed out that model (3.6) can be expressed as Zellner's seemingly unrelated regression (SUR) model and a two stage estimation procedure can be utilized.

To illustrate, let $\xi = [\xi_1 : \ldots : \xi_t]$, $X = [X_1 : \ldots : X_t]$ and the elements of $\Gamma$ be denoted by $\gamma_1, \ldots, \gamma_t$. Then model (3.6) can be written as

$$E(Y) = A[\xi_1 : \ldots : \xi_t] + [X_1 \gamma_1 : \ldots : X_t \gamma_t]$$

$$= [A\xi_1 + X_1 \gamma_1 : \ldots : A\xi_t + X_t \gamma_t]$$

$$= [A_1 \theta_1 : \ldots : A_t \theta_t],$$

where $A_k = [A : X_k]$ and $\theta_k = \begin{pmatrix} \xi_k \\ \gamma_k \end{pmatrix}$.  

55
Thus writing \( E(Y) = Y = [Y_1 : \ldots : Y_t] \) we have

\[
E[Y_1 : \ldots : Y_t] = [A_1\theta_1 : \ldots : A_t\theta_t],
\]

which is a SUR model with the covariance matrix of \( \text{vec}(Y') = I \otimes V \).

Now consider the maximum likelihood estimation based on (3.13) and the covariance \( I \otimes V \) under the assumption of normality. Then the log of the likelihood function can be written as

\[
\log l = -\frac{n}{2} \ln |V| - \frac{1}{2} \sum_{i=1}^{t} \sum_{j=1}^{t} \text{tr}\{v_{ij}(Y_i - A_i\theta_i)'(Y_j - A_j\theta_j)\},
\]

where \( v_{ij} \) is the \((i,j)\)th element of \( V^{-1} \). Differentiating \( \log l \) with respect to \( \theta_k \) and equating the derivative to zero, we get

\[
\hat{\theta}_k = (A_k' A_k)^{-1} A_k' Y_k + (A_k' A_k)^{-1} A_k' \sum_{i \neq k} (Y_i - A_i\theta_i) v_{ik}(v_{kk})^{-1}.
\]

The solutions \( \hat{\theta}_k, k = 1, \ldots, t \) can be obtained using the algorithm provided by Verbyla and Venables (1988). Equation (3.15) can be regarded as coming from the conditional model with the conditional expectation and variance respectively as

\[
E[Y_k/Y_{i,i \neq k}] = A_k\theta_k + \sum_{i \neq k} (Y_i - A_i\theta_i) B_{ik},
\]

\[
\text{var}(Y_k/Y_{i,i \neq k}) = I_n \otimes (v_{kk})^{-1},
\]

where \( B_{ik} = -v_{ik}(v_{kk})^{-1} \).

An algorithm for estimation is as follows:

**Step 1.** Estimate the initial \( \theta_k \) marginally by

\[
\hat{\theta}_k^0 = (A_k' A_k)^{-1} A_k' Y_k.
\]
Step 2. To find the \( j^{th} \) iterate, begin with \( \theta_1 \), and regress \( Y_t \) on \( A_t, R_{t-1}^j, \ldots, R_{t-1}^1 \),

where \( \theta_k^{j-1} \) is the \((j-1)^{st}\) step estimate of \( \theta_k \) and \( R_{k}^{j-1} = Y_k - A_k \theta_k^{j-1} \).

Step 3. For the \( j^{th} \) iterate of \( \theta_k \), regress \( Y_k \) on \( A_k, R_{1}^{j-1}, \ldots, R_{k-1}^{j-1}, R_{k+1}^{j-1}, \ldots, R_{t}^{j-1} \).

Step 4. Continue until convergence.

Once the estimates of \( \theta_1, \ldots, \theta_k \) are obtained using the algorithm, the maximum likelihood estimator of \( V \) can be derived using the residuals as \( \hat{V} = (Y - [A_1 \hat{\theta}_1 : \ldots : A_t \hat{\theta}_t])'(Y - [A_1 \hat{\theta}_1 : \ldots : A_t \hat{\theta}_t])/n \). Verbyla (1988) has also provided an algorithm to estimate the parameters under the null hypothesis \( H_0 : \Lambda^\prime M = 0 \). Let the ML estimator of \( V \) under \( H_0 \) be \( \hat{V} \). Then the likelihood ratio test can be constructed in the usual manner to test any linear hypothesis. The advantage of SUR model is in its simplicity of estimation. Only regression calculations are required making its implementation quite simple.

In the following, we use the statistic given in (3.12) for testing hypothesis of the form \( L^\prime \zeta M = 0 \). The estimators of the covariance matrix under the null and the non-null hypotheses are obtained under SUR model set up. We apply this approach on the data set given in Example 2. The results are presented in the following table.
<table>
<thead>
<tr>
<th>Hypothesis</th>
<th>$\lambda$</th>
<th>D.O.F</th>
<th>Approximate $\chi^2$ value</th>
<th>p-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Group</td>
<td>0.0596</td>
<td>2</td>
<td>5.6402</td>
<td>0.0596</td>
</tr>
<tr>
<td>Time</td>
<td>0.0255</td>
<td>2</td>
<td>7.3360</td>
<td>0.0255</td>
</tr>
<tr>
<td>Time*Group</td>
<td>0.2406</td>
<td>2</td>
<td>2.8488</td>
<td>0.2406</td>
</tr>
</tbody>
</table>

Notice from the above table, as before, that only the time effect is significant.

### 3.2.4 An alternative linear model approach

Although SUR model approach seems like a good alternative to the multivariate approach, both of these approaches have serious shortcomings. None of these approaches will handle unbalanced data. Suppose the $i^{th}$ individual has $t_i$, $i = 1, ..., n$ repeated measurements. Then the analysis of data is more complex and both of the approaches fail. However, an alternative modeling which can handle the unbalanced data will be described below. This alternative approach is also noted by Verbyla and Cullis (1990).

Consider model (3.6)

$$Y_{nxt} = A_{nxm} \xi_{mxt} + X_{nxt}t_{xt} + E_{nxt},$$

where $\Gamma = diag(\gamma_1, ..., \gamma_5)$. Then the $i^{th}$ row of $Y$ can be modeled as

$$y_{i,xt} = a'_i[\xi_1 : ... : \xi_5] + [x_{i1}, ..., x_{it}] diag(\gamma_1, ..., \gamma_5) + \epsilon'_i$$

$$= [a'_i \xi_1 : ... : a'_i \xi_5] + [\gamma_1, ..., \gamma_5] diag(x_{i1}, ..., x_{it}) + \epsilon'_i.$$
Taking transpose of both sides of (3.19) we can write

\[
y_i = \begin{bmatrix}
a_i' & 0 & \ldots & 0 \\
0 & a_i' & \ldots & 0 \\
\vdots & \vdots & \ddots & \vdots \\
0 & 0 & \ldots & a_i'
\end{bmatrix} \begin{bmatrix}
\xi_i \\
\xi_i \\
\vdots \\
\xi_i
\end{bmatrix} + \begin{bmatrix}
x_{i1} & 0 & \ldots & 0 \\
0 & x_{i2} & \ldots & 0 \\
\vdots & \vdots & \ddots & \vdots \\
0 & 0 & \ldots & x_{it}
\end{bmatrix} \begin{bmatrix}
\gamma_i \\
\gamma_i \\
\vdots \\
\gamma_i
\end{bmatrix} + \epsilon_i
\]

where \( A_i = \text{diag}(a_1', ..., a_i') \), \( D_i = \text{diag}(x_{i1}, ..., x_{it}) \), \( \eta' = [\xi_i', ..., \xi_i'] \) and \( \gamma' = (\gamma_1, ..., \gamma_i) \).

Further, the above equation can be rewritten in the form

\[
y_i = \begin{bmatrix}
A_i \quad D_i
\end{bmatrix} \begin{bmatrix}
\eta \\
\gamma
\end{bmatrix} + \epsilon_i
\]

\[
y_{i\times1} = B_{i\times1} \begin{bmatrix}
\eta \\
\gamma
\end{bmatrix} + \epsilon_i
\]

where \( B_i = [A_i : D_i] \), and \( \beta = \begin{bmatrix}
\eta \\
\gamma
\end{bmatrix} \).

Now let \( y' = (y_1', ..., y_n') \) and \( \epsilon' = (\epsilon_1', ..., \epsilon_n') \). Then

\[
y = \begin{bmatrix}
B_1 \\
\vdots \\
B_i \\
\vdots \\
B_t
\end{bmatrix} \beta + \epsilon
\]

or

\[
y = B\beta + \epsilon,
\]

(3.21)
which is an usual linear model with \( \text{cov}(\epsilon) = \Omega = I_n \otimes V \), where \( V \) is a \( t \times t \) covariance matrix of the repeated measures, and \( B = (B_1', ..., B_n')' \).

Then the general least squares estimator of \( \beta \) is

\[
\hat{\beta} = (B'\Omega^{-1}B)^{-1}B'\Omega^{-1}y.
\] (3.22)

Writing \( B' = (B_1' : ... : B_n') \) and \( \Omega^{-1} = I \otimes V^{-1} \), we have

\[
\hat{\beta} = \left( \sum_{i=1}^{n} B_iV^{-1}B_i \right)^{-1}\left( \sum_{i=1}^{n} B_iV^{-1}y_i \right).
\] (3.23)

We may use any two stage procedure to estimate \( \beta \) and \( V \). Further, any hypothesis of interest can be tested using the general linear hypothesis \( H_0 : L\beta = 0 \).

For any specific structured covariance matrix \( V \), this modeling approach can be taken. The SAS statistical procedure PROC MIXED can be adopted to find the estimates and to test any hypothesis of interest. See Khattree and Naik (1995) for some data analysis using PROC MIXED.

As it was pointed out earlier, this modeling technique enables us to deal with unbalanced data as well. Suppose \( y_i \) is vector of order \( t_i \times 1 \) containing the repeated measurements on the \( i^{th} \) individual. Let \( y' \) and \( \epsilon' \) be as defined before. The order of \( y \) and \( \epsilon \) is now \( \sum_{i=1}^{n} t_i \times 1 \). Then the model is

\[ y = B\beta + \epsilon, \]

where \( \text{cov}(\epsilon) = \text{diag}(V_1, ..., V_n) \), and \( V_i \) is of order \( t_i \times t_i \).

Assuming a specific structure for \( V_i \) becomes almost inevitable in this case. It is common to assume that \( V_i \) depends only on a few parameters, say \( \theta_1, ..., \theta_k \). See
Jennrich and Schluchter (1986) for a list of possible structures that can be used. Common structures for $V_i$ are equicorrelation ($\theta_1 = \sigma^2, \theta_2 = \rho$) and autoregressive ($\theta_1 = \sigma^2, \theta_2 = \rho$) structures. The maximum likelihood estimation method is adopted to estimate the unknown parameters $\theta_1, \ldots, \theta_k$ and $\beta$. The form of the MLE of $\beta$ remains as in (3.23) except that the matrix $V$ has to be replaced by the matrix $V_i$ that is of order $t_i \times t_i$ and $B_i$ is of order $t_i \times (m + 1)t_i$, for $i = 1, \ldots, n$. These (unbalanced) data can also be analyzed by PROC MIXED procedure of SAS software.
Chapter 4

Analysis Of Multivariate Repeated Measurements

In this chapter we consider a set of $t$ repeated measurements on $p$ variables (or characteristics) on each of the $n$ individuals. Thus data on each individual is a $p \times t$ matrix. The $n$ individuals themselves may be divided and randomly assigned to $g$ groups. Analysis of these data using a MANOVA model is considered. The well known Satterthwaite type approximation to the distribution of a quadratic form in normal variable is extended to the distribution of a multivariate quadratic form in multivariate normal variates. The multivariate tests using this approximation are developed for testing the usual hypotheses.

When measurements on a variable (or a characteristic) are made at several occasions or under different treatment conditions on the same experimental unit, we
have repeated measures (or longitudinal) data. Repeated measures data routinely occur in many diverse fields like Medicine, Psychology, and Education. Analysis of these data needs special care since the measurements made at different occasions on the same individual may quite likely be correlated. A typical set of repeated measures data taken on \( n(=n_1\ldots+n_g) \) individuals forming \( g \) groups over \( t \) occasions (time points) are shown in Figure 4.1. The measurement taken on the \( j^{th} \) individual belonging to the \( i^{th} \) group at the \( t^{th} \) occasion is denoted by \( y_{ijtk} \), where

\[ k = 1,\ldots,t, \quad j = 1,\ldots,n_i, \quad i = 1,\ldots,g. \]

The problems of interest are to test for (i) the time effect, (ii) the group effect, and (iii) the time*group interaction. Let us assume that the vector of \( t \) measurements on each individual is a sample from a \( t \)-variate normal distribution with a certain positive definite covariance matrix, say \( \mathbf{V} \). The above three problems can be solved using profile analysis, a standard technique in the multivariate statistical analysis. For example, see Morrison (1976). In many practical problems where the repeated measures occur, the covariance matrix \( \mathbf{V} \) is found to have some structure. In that case, the multivariate tests are found to be less powerful and use of univariate analysis of variance or regression models is recommended.

If \( \mathbf{V} \) has the simple structure \( \sigma^2\mathbf{I} \), it is clear that the above three problems, (i)-(iii), can be written in the form of testing of hypothesis problems in a two-way analysis of variance model. Suppose \( \mathbf{V} = \sigma^2\mathbf{V}(\rho) \), where \( \mathbf{V}(\rho) = (1 - \rho)\mathbf{I} + \rho\mathbf{1}\mathbf{1}' \), \( \mathbf{I} \) is an identity matrix and \( \mathbf{1} \) is a vector of ones. Then the analysis of repeated
measures data is essentially same as that of a typical split-plot design. The usual tests of hypotheses in the split-plot design will address the problems (i)-(iii) (see Winer (1971)).

Many authors, for example, Baldessari (1965), Huynh and Feldt (1970), and Rouanet and Lepine (1970) have characterized the class of all covariance structures for $V$, such that the split-plot type of analysis will remain invariant. A typical member of this class, for a fixed vector $a' = (a_1, ..., a_t)$, is

$$V = \sigma^2(I + a1' + 1a') = \sigma^2 H(a).$$

(4.1)

This structure is termed as type H structure. Recently, Chaganty and Vaish (1995) studied this characterization more closely and produced the following form for $H(a)$:

$$H^*(a) = I + \frac{1}{t}(a1' + 1a') - \frac{1}{t}(1 + \bar{a})11',$n

where $a' = (a_1, ..., a_t)$ is such that

$$\frac{1}{t} \sum_{i=1}^{t} (a_i - \bar{a})^2 \leq \bar{a}$$

(4.2)

and $\bar{a}$ is the mean of the components of the vector $a$. This characterization by Chaganty and Vaish (1995) is most explicit in the sense that it gives easy condition (4.2) on the elements of $a$ such that $V$ is positive (semi) definite. The likelihood ratio test for testing for type H structure, using the results of Mauchly (1940), is given by Huynh and Feldt (1970). For certain efficiency studies involving these type of structures see Jensen (1982).
### Table 4.1: Repeated measures data

<table>
<thead>
<tr>
<th>Time</th>
<th>Group</th>
<th>Subject</th>
<th>$y_{111}$</th>
<th>$y_{112}$</th>
<th>...</th>
<th>$y_{11t}$</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>1</td>
<td>2</td>
<td>$y_{121}$</td>
<td>$y_{122}$</td>
<td>...</td>
<td>$y_{12t}$</td>
</tr>
<tr>
<td></td>
<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
</tr>
<tr>
<td></td>
<td>$n_1$</td>
<td></td>
<td>$y_{1n_11}$</td>
<td>$y_{1n_12}$</td>
<td>...</td>
<td>$y_{1n_1t}$</td>
</tr>
<tr>
<td></td>
<td>1</td>
<td>2</td>
<td>$y_{211}$</td>
<td>$y_{212}$</td>
<td>...</td>
<td>$y_{21t}$</td>
</tr>
<tr>
<td></td>
<td>2</td>
<td>2</td>
<td>$y_{221}$</td>
<td>$y_{222}$</td>
<td>...</td>
<td>$y_{22t}$</td>
</tr>
<tr>
<td></td>
<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
</tr>
<tr>
<td></td>
<td>$n_2$</td>
<td></td>
<td>$y_{2n_21}$</td>
<td>$y_{2n_22}$</td>
<td>...</td>
<td>$y_{2n_2t}$</td>
</tr>
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<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
</tr>
<tr>
<td></td>
<td>1</td>
<td>$g$</td>
<td>$y_{g11}$</td>
<td>$y_{g12}$</td>
<td>...</td>
<td>$y_{g1t}$</td>
</tr>
<tr>
<td></td>
<td>$g$</td>
<td>2</td>
<td>$y_{g21}$</td>
<td>$y_{g22}$</td>
<td>...</td>
<td>$y_{g2t}$</td>
</tr>
<tr>
<td></td>
<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
</tr>
<tr>
<td></td>
<td>$n_g$</td>
<td></td>
<td>$y_{gn_g1}$</td>
<td>$y_{gn_g2}$</td>
<td>...</td>
<td>$y_{gn_gt}$</td>
</tr>
</tbody>
</table>

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A strategy for analysis of repeated measures data may be as follows: Test for the type H covariance structure; if accepted adopt the univariate split-plot type of analysis for solving (i)-(iii), else adopt the profile analysis. When the hypothesis of type H structure for $V$ is rejected, one can still use the F-statistics of the split-plot analysis. However, the distributions of these test statistics will no longer be exact. Box (1954) and Geisser and Greenhouse (1958) have developed usable approximations to the distributions of these F-statistics using the result of Satterthwaite (1941). Satterthwaite (1941) approximated the distribution of a quadratic form in normal variables, by a scale multiple of a chi square distribution such that the expected value and the variance of the quadratic form are equal to those of the approximating quantity. This approximation is called Satterthwaite approximation in the literature. Practical implementation of this method requires estimation of the degrees of freedoms of the approximate chi square or the F distributions. Various procedures to estimate the degrees of freedoms are available in the literature. For example, see Greenhouse and Geisser (1959) and Huynh and Feldt (1976). Several authors have studied the effect of (a) heterogeneity of covariance, (b) estimating the degrees of freedom by different methods, and (c) sample size considerations, on the distributions of the F-statistics involved, using simulation experiments. For example, see Huynh and Feldt (1980), Keselman and Keselman (1990), and Quintana and Maxwell (1994). The complete analysis of repeated measures data, including the estimation of the degrees of freedom, has been successfully implemented.
in several statistical softwares. A detailed explanation of the analysis of repeated measures data using the SAS software can be found in Khattree and Naik (1995).

When the observations on $n$ experimental units are made on a set of $p$ variables (or characteristics) at $t$ occasions, we have what can be termed as a set of multivariate repeated measures data. Analysis of these data is further complicated by the existence of correlation among the measurements on different variables along with the correlation among measurements taken at different occasions. A typical set of multivariate repeated measures data would be in the same form as Figure 4.1, except that, $y_{ijk}$ now is a $p \times 1$ vector of measurements on $p$ characteristics. For clarity we denote this vector by $Y_{ijk}$. Analysis of these type of data is considered in this article. Considering a mixed effects MANOVA (Multivariate Analysis of Variance) model we formulate the usual testing of hypothesis problems. The approximate distributions of various SSCP (sums of squares and crossproducts) matrices are derived and used to test the hypotheses. The derivation of the approximate distribution of a SSCP matrix (a multivariate quadratic form) is in the spirit of Satterthwaite (1941), Box (1954) and Geisser and Greenhouse (1958).

4.1 General covariance structure

Suppose, as before, a set of repeated measurements at $t$ occasions are taken on $p$ variables on each of $n(= n_1 + ... + n_g)$ individuals belonging to $g$ groups. The
problems of interest are (i) test for the difference between groups, (ii) test for the
difference between the occasions and (iii) test for group and occasion interaction.

It may also be of interest to test certain hypothesis about the mean vector of \( p \)
variables. We adopt the multivariate analysis of variance (MANOVA) technique to
analyze the data so that our questions of interest can be answered.

Let \( Y_{ijk}, k = 1, \ldots, t, j = 1, \ldots, n_i, i = 1, \ldots, g, \) be a \( p \times 1 \) vector of measurements
on the \( j^{th} \) individual in the \( i^{th} \) group at the \( k^{th} \) occasion. Let \( Y_{ij}' = (Y_{ij1}, \ldots, Y_{ijg}). \)
Then \( Y_{ij} \) is \( pt \times 1 \) random observational vector corresponding to the \( j^{th} \) individual
in the \( i^{th} \) group. Let \( \text{cov}(Y_{ij}) = \Omega, \) for \( j = 1, \ldots, n_i, i = 1, \ldots, g, \) where \( \Omega \) is a
positive definite matrix. Define the \( n \times pt \) matrix \( Y \) as \( Y = (Y_{11}, \ldots, Y_{gn})'. \) Then
the multivariate repeated measures data can be modeled as

\[
Y = XB + E,
\]

where \( X \) is a known design matrix, \( B \) is the matrix of unknown parameters, and
\( E \) is the matrix of errors such that the rows of \( E \) are independently distributed as
multivariate normal with zero mean vector and variance covariance matrix \( \Omega. \) Any
hypothesis about the effect of the time factor, the group factor, or the interaction
between them or any linear hypothesis about the expected values of \( Y_{ijk} \) can be
formulated in the form of the general linear hypothesis

\[
LBM = 0 \quad (4.3)
\]

for known and full rank matrices \( L \) and \( M. \) Using Wilks’ \( \Lambda \) or any other standard
multivariate tests the null hypothesis (4.3) can be tested. This approach for analyzing the multivariate repeated measures data is commonly adopted in practice (see Timm, 1980). Suppose \( \Omega = I_t \otimes \Sigma \), for a \( p \times p \) positive definite matrix \( \Sigma \), to indicate that the measurements taken over time are uncorrelated. Then the multivariate repeated measures data can be analyzed using two-way MANOVA model

\[
Y_{ijk} = \mu + \alpha_i + \beta_k + (\alpha \beta)_{ik} + \epsilon_{ijk}, \tag{4.4}
\]

where \( \mu, \alpha_i, \beta_k, (\alpha \beta)_{ik} \) are all \( p \times 1 \) vectors with the usual meaning as in the two-way MANOVA model. The hypotheses of interest are

(i) \( \alpha_1 = \ldots = \alpha_t \) (no group effect), (ii) \( \beta_1 = \ldots = \beta_t \) (no time or occasion effect) and (iii) \( (\alpha \beta)_{ik} = \ldots = \alpha_t \) (no group and time interaction). Under the assumption \( \Omega = I_t \otimes \Sigma \), the errors \( \epsilon_{ijk} \) are independently distributed with zero mean vector and variance covariance matrix \( \Sigma \). Testing of hypothesis problems (i)-(iii), can be tackled using the usual MANOVA technique.

Next, to accommodate simple correlation among repeated measures, let us consider mixed effects MANOVA model (similar to the split-plot design model of the univariate analysis of the usual repeated measures data). The MANOVA table similar to the ANOVA table for split-plot design model, is shown in Table (4.2).

Here the \( p \times nt \) matrix \( Y \) is defined as

\[
Y = (Y_{111}, \ldots, Y_{11t}, \ldots, Y_{1n_11}, \ldots, Y_{1n_1t}, \ldots, Y_{gn_p1}, \ldots, Y_{gn_pt}) \quad \text{and} \quad N = nt. \quad \text{The ma-}
\]
<table>
<thead>
<tr>
<th>Source</th>
<th>D.O.F.</th>
<th>SSCP Matrix</th>
<th>Dist. Under $H_0$</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Between Groups</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Groups</td>
<td>$g - 1$</td>
<td>$Q_1 = YA_1Y'$</td>
<td>$W_p(g - 1, \Sigma)$</td>
</tr>
<tr>
<td>Individuals</td>
<td>$n - g$</td>
<td>$Q_2 = YA_2Y'$</td>
<td>$W_p(n - g, \Sigma)$</td>
</tr>
<tr>
<td><strong>Within Groups</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Time</td>
<td>$t - 1$</td>
<td>$Q_3 = YA_3Y'$</td>
<td>$W_p(t - 1, \Sigma)$</td>
</tr>
<tr>
<td>Time*Group</td>
<td>$(g - 1)(t - 1)$</td>
<td>$Q_4 = YA_4Y'$</td>
<td>$W_p((t - 1)(g - 1), \Sigma)$</td>
</tr>
<tr>
<td>Error</td>
<td>$(t - 1)(n - g)$</td>
<td>$Q_5 = YA_5Y'$</td>
<td>$W_p((t - 1)(n - g), \Sigma)$</td>
</tr>
<tr>
<td>Total</td>
<td>$N - 1$</td>
<td>$Y(I_N - \frac{1}{N}J_N)Y'$</td>
<td></td>
</tr>
</tbody>
</table>

Table 4.2: Manova table for mixed effects model
trix quadratic forms $Q_1 - Q_5$ are

\[
Q_1 = t \sum_{i=1}^{g} n_i (\bar{Y}_{i..} - \bar{Y}_{..})(\bar{Y}_{i..} - \bar{Y}_{..})' = Y A_1 Y'
\]

\[
Q_2 = t \sum_{i=1}^{g} \sum_{j=1}^{t} n_i (\bar{Y}_{ij} - \bar{Y}_{i..})(\bar{Y}_{ij} - \bar{Y}_{i..})' = Y A_2 Y'
\]

\[
Q_3 = n \sum_{k=1}^{t} (\bar{Y}_{..k} - \bar{Y}_{..})(\bar{Y}_{..k} - \bar{Y}_{..})' = Y A_3 Y'
\]

\[
Q_4 = \sum_{i=1}^{g} n_i \sum_{k=1}^{t} (\bar{Y}_{i,k} - \bar{Y}_{..} - \bar{Y}_{..,k} + \bar{Y}_{...})(\bar{Y}_{i,k} - \bar{Y}_{i..} - \bar{Y}_{.,k} + \bar{Y}_{...})' = Y A_4 Y'
\]

\[
Q_5 = \sum_{i=1}^{g} \sum_{j=1}^{t} \sum_{k=1}^{t} (Y_{ijk} - \bar{Y}_{ij} - \bar{Y}_{i,k} + \bar{Y}_{i..})(Y_{ijk} - \bar{Y}_{ij} - \bar{Y}_{i,k} + \bar{Y}_{i..})' = Y A_5 Y'
\]

with appropriate choice of matrices $A_1 - A_5$. These matrices are symmetric of order $N \times N$ and can be easily derived. For example, see Geisser and Greenhouse (1958).

It is clear from the works of Khatri (1962), Arnold (1979), Reinsel (1982), and Mathew (1989) that the above matrix quadratic forms are independent of each other and under the appropriate null hypothesis each has a scale multiple of a Wishart distribution with a certain degrees of freedom.

Thomas (1983) considered a class of structures for $\Omega$, members of which are sufficient to keep the multivariate analysis of variance (MANOVA) obtained under the mixed effects model invariant preserving the independence and distributions of the SSCP matrices of Table 4.1. Pavur (1987) characterized the class of all covariance structures for $\Omega$ under which the MANOVA remains invariant. Vaish (1994) in a recent Ph.D. thesis pointed out using counter examples that Pavur (1987)'s characterization may contain some matrices that are not non negative.
definite. Further, he has provided a more elegant characterization of $\Omega$, which is:

$$\Omega = (I_2 - \frac{1}{t}J_t) \otimes \Sigma + \frac{1}{t}(1_t \otimes I_p)H' + \frac{1}{t}H(1'_t \otimes I_p) - \frac{1}{t}1_t1'_t \otimes \tilde{H},$$

(4.5)

where $H$ is a $pt \times p$ arbitrary matrix of rank $(H) \leq p$, and $\tilde{H} = \frac{1}{t} \sum_{i=1}^{t} H_i$. Further, the $p \times p$ matrices $H_i$'s are such that $H = (H'_1, ..., H'_t)'$.

The structure (4.5) is slightly more general than the structure

$$\Omega = \begin{bmatrix}
\Sigma_1 & \Sigma_2 & \cdots & \Sigma_2 \\
\Sigma_2 & \Sigma_1 & \cdots & \Sigma_2 \\
\cdots & \cdots & \cdots & \cdots \\
\Sigma_2 & \Sigma_2 & \cdots & \Sigma_1
\end{bmatrix}$$

considered by Arnold (1979) and Mathew (1989). It may be noted that this structure is a multivariate analogue of the well known equicorrelation structure. Similarly, the covariance structure

$$\Omega = I_t \otimes \Sigma_1 + 1_t1'_t \otimes \Sigma_2$$

considered by Reinsel (1982) is also a particular case of (4.5).

Boik (1988) independently (of Pavur (1987)) showed that the structure considered by Thomas (1983) is necessary and sufficient for MANOVA to remain invariant. He also developed the likelihood ratio test for testing for this structure. Further, Boik (1988) considered the Satterthwaite type approximations (multivariate Satterthwaite approximation) to the distributions of various sums of squares and
crossproducts (SSCP) matrices of the MANOVA table, when the covariance matrix \( \Omega \) is any general positive definite matrix. While approximating the distribution of a SSCP matrix by a scale multiple of Wishart distribution, Boik (1988) assumed that the expected value and the trace of the covariance matrix of the SSCP matrix are equal to those of approximating matrix.

Tan and Gupta (1983) and recently Khuri, Mathew, and Nel (1994) also have considered the problems relating to multivariate Satterthwaite approximation to a SSCP matrix. Their approximation is different from Boik (1988) in the sense that they use generalized variance (determinant of the covariance matrix) instead of trace, for their approximations.

**4.2 Covariance structure \( V \otimes \Sigma \)**

All the work done in the literature thus far, about multivariate repeated measures, has the basic assumption that \( \text{cov}(Y_{ij}) = \Omega \), where \( \Omega \) is a positive definite matrix. The structures similar to type H structure on \( \Omega \) were found, under which the MANOVA remained invariant. As we have noted in the previous section, the multivariate Satterthwaite approximations for SSCP matrices of the MANOVA table (Table 4.1) were derived under the basic assumption that \( \text{cov}(Y_{ij}) = \Omega \). In this section we start with the assumption that \( \text{cov}(Y_{ij}) = V \otimes \Sigma \), where \( V \) is a \( t \times t \) positive definite matrix. This structure has several advantages over the general covariance
structure. First, it is well known that (Crowder and Hand (1993), Jones (1993)) the correlation matrix of the repeated measures usually has a simple structure as opposed to a general structure. In our formulation it is easier to accommodate any structure for the correlation matrix of repeated measures (via $V$). Next, the number of unknown parameters of the variance covariance matrix, in our formulation, is much less, $(t(t + 1)/2 + p(p + 1)/2)$, as opposed to $(pt(pt + 1)/2)$ in the case of general covariance structure. Furthermore, as we will see later, the multivariate Satterthwaite approximations to various SSCP matrices is much simpler in the present situation.

Note that the $i^{th}$ row of the $p \times nt$ matrix $Y$, where $Y = (Y_{11i}, \ldots, Y_{1ti}, \ldots, Y_{1_{n1}i}, \ldots, Y_{1_{nt}i}, \ldots, Y_{p_{n1}i}, \ldots, Y_{p_{nt}i})$, has the covariance matrix proportional to $\Delta = I_n \otimes V$. In fact it is $\sigma_{ii}\Delta$, where $\sigma_{ii}$ is the $i^{th}$ diagonal element of $\Sigma$.

Suppose $V = I_t$, that is, the repeated measures are uncorrelated. Then using the two-way MANOVA model (4.4) analysis of the multivariate repeated measures data can be carried out as before. Now suppose $V = V(\rho) = (1 - \rho)I_t + \rho J_t$. Using a result which is multivariate analogue of Cochran's theorem for quadratic forms it can be shown that the MANOVA in Table 1 will remain invariant for this case (see Theorem 4.6.3 of Vaish (1994)). Further, a characterization of the class of structures for $V$ such that the MANOVA remains invariant yields type H structure (4.1) for $V$. Many interesting results about the structure $V \otimes \Sigma$ are summarized.
4.2.1 Test for type H structure

In this section we construct the likelihood ratio test for testing

\[ H_0 : \text{cov}(Y_{ij}) = H(a) \otimes \Sigma \quad \text{vs} \quad H_a : \text{cov}(Y_{ij}) = V \otimes \Sigma. \]

Make a transformation on the vector of measurements \( Y_{ij} \) to \( U_{ij} \), such that \( \text{cov}(U_{ij}) = W \otimes \Sigma \), where \( W = C \Sigma C' \). The choice of \( C \) is such that \( C'1 = 0 \) and \( CC' = I_{t-1} \).

One possible choice for \( C \) is the appropriately chosen submatrix of the Helmert’s matrix. It is true that testing the above hypothesis is equivalent to testing

\[ H_0 : \text{cov}(U_{ij}) = I \otimes \Sigma \quad \text{vs} \quad H_a : \text{cov}(U_{ij}) = W \otimes \Sigma. \quad (4.6) \]

A likelihood ratio test for testing (4.6) is given by

\[ \lambda = \frac{|\hat{W}|^{n_p/2} |\hat{\Sigma}|^{n(t-1)/2}}{|\hat{\Sigma}_0|^{n(t-1)/2}}, \quad (4.7) \]

where \( \hat{W} \) and \( \hat{\Sigma} \) are respectively the maximum likelihood estimates (MLEs) of \( W \) and \( \Sigma \) under the non-null case \( (H_a) \) and \( \hat{\Sigma}_0 \) is the MLE of \( \Sigma \) under the restriction of the null hypothesis \( (H_0) \). Then using standard results \(-2ln \lambda\) has a \( \chi^2 \) distribution with degrees of freedom equal to \( t(t-1)/2 \).

4.2.2 Derivation of the MLEs

We first derive the maximum likelihood estimators of \( W \) and \( \Sigma \) under \( (H_a) \). We have a random sample \( U_{ij}, j = 1, ..., n_i, i = 1, ..., g \) from a \( p(t-1) \)-variate normal
distribution with a mean $\mu_i$ and variance covariance matrix $W \otimes \Sigma$. Here $\mu_i$ is a $p(t-1) \times 1$ vector given by $\mu'_i = (\mu'_{i1}, \ldots, \mu'_{i(t-1)})$, where $\mu_{ik}$ is a $p \times 1$ vector representing the expected value of the transformed variable corresponding to the $i^{th}$ group and the $j^{th}$ time point.

The log-likelihood function is given by

$$
\ln l = \ - \frac{p(t-1)n}{2} \ln(2\pi) - \frac{n}{2} \ln |W \otimes \Sigma| \\
- \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} (U_{ij} - \mu_i)'W^{-1} \otimes \Sigma^{-1}(U_{ij} - \mu_i).
$$

Let $B = \sum_{i=1}^{g} \sum_{j=1}^{n_i} (U_{ij} - \overline{U}_i)(U_{ij} - \overline{U}_i)'$, with $\overline{U}_i = \sum_{j=1}^{n_i} U_{ij}/n_i$.

Then the likelihood function can be written as

$$
\ln l = \ - \frac{np(t-1)}{2} \ln(2\pi) - \frac{n}{2} \ln |W \otimes \Sigma| - \frac{1}{2} tr(W^{-1} \otimes \Sigma^{-1})B \\
- \frac{1}{2} \sum_{i=1}^{g} n_i(\overline{U}_i - \mu_i)'W^{-1} \otimes \Sigma^{-1}(\overline{U}_i - \mu_i).
$$

(4.8)

Next, we partition the $p(t-1) \times 1$ vector $(U_{ij} - \overline{U}_i)$ into $(t-1)$ blocks of $p \times 1$ vectors such that $(U_{ij} - \overline{U}_i) = (U_{ij1} - \overline{U}_{i1}, \ldots, U_{ij(t-1)} - \overline{U}_{i(t-1)})'$. Using this partition of $(U_{ij} - \overline{U}_i)$, we rewrite the likelihood (4.8) and maximize it simultaneously with respect to $\Sigma$ and $W$. We get the following likelihood equations for estimating $\Sigma$ and $W$.

$$
\dot{\Sigma} = \frac{1}{n(t-1)} \sum_{k=1}^{t-1} \sum_{l=1}^{t-1} \sum_{i=1}^{g} \sum_{j=1}^{n_i} (U_{ijk} - \overline{U}_{i,k})(U_{ijl} - \overline{U}_{i,l})
$$

(4.9)

$$
\dot{W} = \frac{1}{np} A_i
$$

(4.10)
where $\hat{w}_{kl}$ is the $(k,l)$-th element of $\hat{W}^{-1}$. The $(k,l)$-th element of matrix $A$ is given by

$$A_{kl} = \sum_{i=1}^{g} \sum_{j=1}^{n_i} (U_{ijk} - U_{i.k})' \hat{\Sigma}^{-1} (U_{ijl} - U_{i.l}).$$

The above equations are to be solved iteratively to get the estimates of $\Sigma$ and $W$. There is no general consensus as to when iterative methods should be stopped and the current values declared to be ML estimators. In our illustrative example, we have selected the following stopping rule. Compute two matrices: (a) matrix of difference between two successive solutions of (4.9), and (b) matrix of difference between two successive solutions of (4.10). Continue the iterations until the maximums of the absolute values of the elements of the matrices in (a) and (b) are smaller than the pre-specified quantities. A computer program using IML of SAS software is available with the author to compute these estimates.

In a manner analogous to what is described above, we can get the estimate of $\Sigma$ under restriction of the null hypothesis $(H_0)$. The Maximum likelihood estimate of $\Sigma$ under the null hypothesis is given by

$$\hat{\Sigma}_0 = \frac{1}{n(t-1)} \sum_{k=1}^{t-1} \sum_{i=1}^{g} \sum_{j=1}^{n_i} (U_{ijk} - \bar{U}_{i,k})(U_{ijl} - \bar{U}_{i,l})'.$$  \hspace{1cm} (4.11)

If $H_0$ is accepted, one can use the MANOVA of (4.2) to analyze the data. Otherwise multivariate Satterwaite type approximation can be applied to find the approximation to the distributions of the SSCP matrices. This approximation will be described in the following sections.
4.2.3 A simulation study

In section 4.2.1 we derive a test for testing

\[ H_0 : \text{cov}(Y_{ij}) = H(a) \otimes \Sigma \quad \text{vs} \quad H_a : \text{cov}(Y_{ij}) = V \otimes \Sigma, \]

using the likelihood ratio principle and derive the MLE’s in section 4.2.2. Since the choice of the method of data analysis rests entirely on the outcome of this testing procedure, it is important that the test we are using is powerful enough. To study the power of our test given by (4.7) we conduct a small simulation study section.

We know that the power function \( \beta_\lambda(\theta) \) of the test statistic \( \lambda \) is defined as

\[ \beta_\lambda(\theta) = P_\theta(\text{Rejecting } H_0) = P_\theta(\lambda > \lambda_c) \tag{4.12} \]

as a function of \( \theta \), where \( \theta \) is the vector of all parameters of the covariance matrix \( V \otimes \Sigma \) and \( \lambda_c \) is such that under \( H_0 : \theta = \theta_0 \), \( \beta_\lambda(\theta_0) \) is a specified value, say 0.05.

In order to calculate the power of the test, we generate random samples from multivariate normal distribution \( N_{pt}(0, V \otimes \Sigma) \), where \( V \) is the autoregressive covariance structure and \( \Sigma \) is a positive definite matrix. Note that autoregressive structure is not a member of class of all matrices of type \( H \). For \( p=3 \),

\[ \Sigma = \begin{bmatrix} 2 & 1 & 2 \\ 1 & 4 & 3 \\ 2 & 3 & 5 \end{bmatrix}, \]

and for \( t=3 \) and 5 and for different choices of \( \rho \), we generate multivariate normal samples. To represent both small samples and large samples we have selected...
sample sizes of 12 and 30. We have selected $N=1000$ simulation runs of each case and the power of the test was calculated as

$$P_0(\text{Rejecting } H_0) = \frac{\#(-2\ln \lambda > x^2_\nu)}{1000}$$

(4.13)

where $x^2_\nu$ is the critical value under chi-square distribution with degrees of freedom $t(t - 1)/2$ and level of significance $\alpha=0.05$. We summarize the results of our study for different choices of $n$, $t$ and $\rho$, in the following table:

<table>
<thead>
<tr>
<th>$\rho$</th>
<th>$n = 12$</th>
<th>$n = 30$</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>0.0</td>
<td>0.2</td>
</tr>
<tr>
<td>$t = 3$</td>
<td>0.04</td>
<td>0.075</td>
</tr>
<tr>
<td>$t = 5$</td>
<td>0.059</td>
<td>0.168</td>
</tr>
</tbody>
</table>

We make the following observations from the above table:

We see that power of the test is close to one as we depart from $H_0$. In other words as $\rho$ increases power of the test increases quite rapidly. Not surprisingly, power is higher for larger values of $n$, $\rho$ being the same.

When $\rho=0$, it amounts to sampling from the null distribution, and we expect the power function to take values that meets the size condition. That is, the power should be close to the level of significance, which is in fact the case in the present context.
4.2.4 Exact null distribution of $Q_i$

In the following, under no structures on $V$, we derive the distribution of $Q_i = Y A_i Y'$, where $A_i$ is an appropriately defined symmetric matrix of order $nt \times nt$. First of all, it is easy to show that $A_1 A_2 = A_2 A_1 = A_3 A_5 = A_5 A_3 = A_5 A_4 = A_4 A_5 = 0$ (see Geisser and Greenhouse(1958)), where $A = I_n \otimes V$. Hence the matrix quadratic forms $Q_1$ and $Q_2$, $Q_1$ and $Q_5$, $Q_2$ and $Q_5$, $Q_3$ and $Q_5$, and $Q_4$ and $Q_5$, are all pairwise independent. Let the rank of $(A_i) = \nu_i$ (for example, $\nu_1 = g - 1$ and so on). We observe that each row of $Y$ is a multivariate normal vector of order $nt \times 1$ and has a covariance matrix proportional to $\Delta = I_n \otimes V$. Since by assumption $\Delta$ is a positive definite matrix, there exist $\Delta^{1/2}$ and $\Delta^{-1/2}$ such that $\Delta = \Delta^{1/2} \Delta^{1/2}$ and $\Delta^{-1} = \Delta^{-1/2} \Delta^{-1/2}$ and that $\Delta^{-1/2} \Delta^{1/2} = I_{nt}$.

Consider $Q_i = Y A_i Y' = Y \Delta^{-1/2} A_i \Delta^{1/2} \Delta^{-1/2} Y' = Z B_i Z'$ with $Z = Y \Delta^{1/2}$ and $B_i = \Delta^{1/2} A_i \Delta^{1/2}$. Now the rows of $Z$ form a random sample of size $nt$ from $N_p(0, \Sigma)$ distribution under certain null hypothesis. Since $B_i$ is a symmetric matrix with rank $\nu_i$, it can be written as $B_i = \Lambda_i \Gamma' \Gamma$, where $\Gamma' \Gamma = \Gamma' \Gamma = I_{nt}$ and $\Lambda_i = \text{Diag}(\lambda_1, ..., \lambda_{\nu_i}, 0, ..., 0)$, where $\lambda_1, ..., \lambda_{\nu_i}$ are the eigenvalues of $B_i$. Thus $Q_i = Z B_i Z' = Z \Gamma \Lambda_i \Gamma' \Gamma' = U \Lambda_i U' = \sum_{j=1}^{\nu_i} \lambda_j U_j U_j'$, where $U_j \sim N_p(0, \Sigma)$ under $H_0$ and $U_1, ..., U_{\nu_i}$ are all independent.

It is well known that $U_j U_j' \sim W_p(1, \Sigma)$, which is a pseudo-Wishart distribution, since the degrees of freedom is less than the dimension, and has no pdf. Thus the distribution of $Q_i$ is same as the distribution of linear combination of $\nu_i$ pseudo-
Wishart random matrices. In summary, we have the following:

\[ Q_1 \sim W_p((g - 1), \Sigma), \]

\[ Q_2 \sim W_p((n - g), \Sigma), \]

\[ Q_3 \sim \sum_{j=1}^{t-1} \lambda_j W_{p,j}(1, \Sigma), \]

\[ Q_4 \sim \sum_{j=1}^{t-1} \lambda_j W_{p,j}(g - 1, \Sigma), \]

\[ Q_5 \sim \sum_{j=1}^{t-1} \lambda_j W_{p,j}(n - g, \Sigma), \]

where \( \lambda_1, \ldots, \lambda_{t-1} \) are the eigenvalues of \((I - 1/t) \mathbf{V}\) and \(W_{p,j}(\nu_j, \Sigma)\), for \(j = 1, \ldots, t - 1\) are Wishart random matrices with \(\nu_j\) degrees of freedom, and mutually independent.

Suppose we want to test \(H_{01} : \alpha_1 = \cdots = \alpha_g\). Then the Wilks' \(\Lambda\) for testing \(H_{01}\) is

\[ \Lambda_1 = \frac{|Q_2|}{|Q_1 + Q_2|}. \]

The test is based on the usual asymptotic distribution (in some situations exact (Rao, 1973, p. 555)) of \(\Lambda_1\). For example,

\[-[(n - g) - \frac{(p - g)}{2}] \ln \Lambda_1 \sim \chi^2_{p(g-1)}\text{approximately}.\]

The approximate distributions of the test statistics for testing \(H_{02} : \beta_1 = \cdots = \beta_t\) and \(H_{03} : (\alpha \beta)_{ik}s\) are all equal, will be derived in the next section.
4.2.5 Approximate null distributions of $Q_3$, $Q_4$, and $Q_5$

In this section, we approximate the distribution of each of SSCP matrices, $Q_3$, $Q_4$, and $Q_5$, to a scale multiple of a Wishart matrix, $gW(h, \Sigma)$, for some constants $g$ and $h$. As in the univariate case the approximation is derived by equating the first two central moments. For that we first find the first two central moments of $Q_i$. It is well known (for example, see Eaton, 1983, p. 305) that for any $S \sim W_p(\nu, \Sigma)$, $E(S) = \nu \Sigma$ and $D(S) = 2 \nu \Sigma \otimes \Sigma$. Here $D(S)$ denotes the variance covariance matrix of all the random quantities in $S$. Using these formulae we have the following:

$E(Q_3) = [tr(V - 1/tJV)]\Sigma$ under $H_{02}: \beta_1 = \ldots = \beta_t$

$E(Q_4) = [(g - 1)tr(V - 1/tJV)]\Sigma$ under $H_{03}: (\alpha \beta)_{ik} = \gamma$ for all $i, k$.

$E(Q_5) = [(n - g)tr(V - 1/tJV)]\Sigma$

also

$D(Q_3) = [2tr(V - 1/tJV)^2] \Sigma \otimes \Sigma$ under $H_{02}$

$D(Q_4) = [2(g - 1)tr(V - 1/tJV)^2] \Sigma \otimes \Sigma$ under $H_{03}$

$D(Q_5) = [2(n - g)tr(V - 1/tJV)^2] \Sigma \otimes \Sigma$.

Suppose we want to approximate the distribution of $Q_3$ by a random matrix having the distribution $g_1W_p(h_1, \Sigma)$ so that the first two central moments of $Q_3$ and $g_1W_p(h_1, \Sigma)$ are the same. Then,

$$tr[(V - 1/tJV)]\Sigma = g_1h_1 \Sigma$$

(4.14)

82
\[ 2[\text{tr}(V - 1/tJV)^2] \Sigma \otimes \Sigma = 2g_2^2 h_1 \Sigma \otimes \Sigma. \quad (4.15) \]

From (4.14) and (4.15)

\[ g_1 = \frac{\text{tr}(V - 1/tJV)^2}{\text{tr}(V - 1/tJV)} \]
\[ h_1 = \frac{[\text{tr}(V - 1/tJV)]^2}{\text{tr}(V - 1/tJV)^2}. \]

Next, to approximate the distribution of \( Q_4 \) by a random matrix having the distribution \( g_2 W_p(h_2, \Sigma) \) so that the first two central moments of \( Q_4 \) and \( g_2 W_p(h_2, \Sigma) \) are the same we have,

\[ (g - 1) \text{tr}[(V - 1/tJV)] \Sigma = g_2 h_2 \Sigma \quad (4.16) \]
\[ 2(g - 1)[\text{tr}(V - 1/tJV)^2] \Sigma \otimes \Sigma = 2g_2^2 h_2 \Sigma \otimes \Sigma. \quad (4.17) \]

From (4.16) and (4.17) we have,

\[ g_2 = \frac{\text{tr}(V - 1/tJV)^2}{\text{tr}(V - 1/tJV)} = g_1 \]
\[ h_2 = (g - 1) \frac{[\text{tr}(V - 1/tJV)]^2}{\text{tr}(V - 1/tJV)^2} = (g - 1)h_1. \]

Therefore, \( Q_4 \sim g_1 W_p((g - 1)h_1, \Sigma) \) approximately.

Similarly, it can shown that \( Q_5 \sim g_1 W_p((n - g)h_1, \Sigma) \) approximately.

Now for testing \( H_{02} : \beta_1 = \ldots = \beta_i \) one can use the Wilks' \( \Lambda \), which is, \( \Lambda_2 = \frac{|Q_5|}{|Q_3 + Q_5|} \) and the fact that
4.2.6 Estimation of the degrees of freedom

Since in practice $V_{\lambda \lambda}$ is unknown the degrees of freedoms in the chi square approximations of (4.18) and (4.19) are unknown. One needs to estimate these so that the distributions in (4.18) and (4.19) can be utilized in applications. For estimating these degrees of freedoms, which are functions of $(V - 1/f)$, we simply need an estimate of $V$. One can use the maximum likelihood estimate of $V$ that is obtained by simultaneously solving the following equations:

$$
\hat{V} = \frac{1}{np} A, \quad (4.21)
$$

where $\hat{v}_{kl}$ is the $(k,l)$-th element of $\hat{V}^{-1}$ and $(k,l)$-th element of matrix $A$ is given by $A_{kl} = \sum_{i=1}^{g} \sum_{j=1}^{n_i} (Y_{ijk} - \bar{Y}_{i,k}) \hat{V}^{-1} (Y_{ijl} - \bar{Y}_{i,l})$.

In the next section we consider an example and illustrate the estimation of the degrees of freedom using the maximum likelihood estimate of $V$. 

\[\]
4.3 Example

To illustrate our methods, we use data from Table 7.2 in Timm (1980), which was also analyzed by Thomas (1983) and Boik (1988). These data were obtained by Timm from T. Zullo of the School of Dental Medicine at the University of Pittsburgh. The study concerns with the relative effectiveness of two orthopedic adjustments of the mandible. Nine subjects were assigned to each of two orthopedic treatments \( (g = 2, n_1 = 9, n_2 = 9) \), called activator treatments. The measurements were made on three characteristics \( (p=3) \) to assess the changes in the vertical position of the mandible at three time points \( (t=3) \) of activator treatment. Thus the data matrix \( Y = (Y_{ij}) \) is an \( 18 \times 9 \) matrix. The data is presented in the following table:

The choice of method to analyze our data, rests on the test of hypothesis about the covariance structure described in section (3). Hence to test the null hypothesis (4.6), first we transform \( Y_{ij} \) to \( U_{ij} \) using the transformation \( U_{ij} = (C \otimes I)Y_{ij} \), with

\[
C = \begin{bmatrix}
0.7071 & -0.7071 & 0.0000 \\
0.4082 & 0.4082 & -0.8165
\end{bmatrix}.
\]

Then we have \( \text{cov}(U_{ij}) = W \otimes \Sigma \).

The maximum likelihood estimates of \( \Sigma \) and \( W \), simultaneously solving (4.9) and (4.10), are given by
Table 4.3: Dental data

<table>
<thead>
<tr>
<th>Subj</th>
<th>S0r-Me (mm)</th>
<th>ANS-ME (mm)</th>
<th>Pal-MP angle (degrees)</th>
</tr>
</thead>
<tbody>
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\[
\hat{\Sigma} = \begin{bmatrix}
0.3861535 & 0.2324518 & -0.16931 \\
0.2324518 & 1.0859517 & -0.097388 \\
-0.16931 & -0.097388 & 0.4350231
\end{bmatrix}
\]

and

\[
\hat{\mathbf{W}} = \begin{bmatrix}
1.5931756 & 0.5033948 \\
0.5033948 & 0.5951812
\end{bmatrix}
\]

Estimate of \( \Sigma \) under \( H_0 \) using (4.11) is

\[
\hat{\Sigma}_0 = \begin{bmatrix}
0.4192387 & 0.2425412 & -0.240329 \\
0.2425412 & 0.7572016 & -0.145422 \\
-0.240329 & -0.145422 & 0.6873045
\end{bmatrix}
\]

The value of the statistic given in (4.7) is \( \lambda=0.0000139 \). Then the test statistic value, \( -2ln\lambda \), is equal to 22.37366. Comparing this with \( \chi_3^2(0.05) = 7.815 \), we clearly reject \( H_0 \) given in (4.6). Therefore, we use multivariate Satterwaiththe type approximation described in sections (3.3-3.4), to find approximation to the distributions of the SSCP matrices. In order to use these methods we need to estimate \( g_1 \) and \( h_1 \) which in turn requires the estimate of \( V \). We simultaneously solve (4.9)
Hypothesis Wilks' $\lambda$ D.O.F Approximate $\chi^2$ value p-value

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Table 4.4: Approximate MANOVA

and (4.10) to get the maximum likelihood estimate of $\boldsymbol{V}$ and it is found to be

$$
\hat{\boldsymbol{V}} = \begin{bmatrix}
2.4224556 & 2.2517394 & 2.2765762 \\
2.2517394 & 2.2246306 & 2.2018579 \\
2.2765762 & 2.2018579 & 2.2294622
\end{bmatrix}.
$$

Using this we get $g_1=0.0636441$ and $h_1=1.5332644$.

We summarize our results in Table 4.3.

For computing the above p-values one can use any easily available softwares. For example, we have used PROBF function of SAS Software. Note from Table 4.3 that only the effect of time factor is significant.
Chapter 5

Analysis of Unbalanced Multivariate Repeated Measurements

In chapter 4 we have considered a set of balanced (same number of measurements on each individual) multivariate repeated measurements. These data can be represented by $Y_{ijk}, k = 1, ..., t; j = 1, ..., n_i; i = 1, ..., g$, which is a $p \times 1$ vector of measurements on the $j^{th}$ individual in the $i^{th}$ group at the $k^{th}$ occasion. Let $Y'_{ij} = (Y'_{ij1}, ..., Y'_{ijt})$. Then $Y_{ij}$ is $pt \times 1$ random observational vector corresponding to the $j^{th}$ individual in the $i^{th}$ group. We have assumed that $cov(Y_{ij}) = V \otimes \Sigma$, where $V$ is a $t \times t$ positive definite matrix. It was seen that if $V = V(\rho) = (1 - \rho)I_t + \rho J_t$, then the analysis of multivariate repeated measurements can be carried out using...
the two-way MANOVA model. Further, it was noted that a characterization of the
class of structures for V such that the MANOVA remains invariant yields type H
structure. On the other hand, if V is completely arbitrary, we have suggested Sat-
terthwaite type approximation to the distribution of multivariate quadratic forms
of SS&CP matrices. Multivariate tests using this approximation were developed
for testing the usual hypothesis.

5.1 Autoregressive structure for V

We now assume autoregressive structure for V, that is, V = (ρ| |). This structure
is neither type H, nor completely arbitrary. With this structure on V, we derive the
approximate distributions of SS&CP matrices. In chapter 4, assuming no structure
on V we have shown the following:

\[ Q_1 \sim W_p((g - 1), \Sigma), \]
\[ Q_2 \sim W_p((n - g), \Sigma), \]
\[ Q_3 \sim W_p(h_1, \Sigma) \text{ approximately,} \]
\[ Q_4 \sim W_p((g - 1)h_1, \Sigma) \text{ approximately,} \]
\[ Q_5 \sim W_p((n - g)h_1, \Sigma) \text{ approximately,} \]

with

\[ g_1 = \frac{tr(V - 1/tJV)^2}{tr(V - 1/tJV)} \]

90
\[ h_1 = \frac{[\text{tr}(V - 1/tJV)]^2}{\text{tr}(V - 1/tJV)^2}. \]

Now, let \( V = (\rho^{i-j}) \). Next we derive expressions for \( g_1 \) and \( h_1 \) in terms of \( \rho \).

Notice that

\[
\text{tr}(V - 1/tJV) = \text{tr}(V) - 1/t \text{tr}(JV), \tag{5.1}
\]

\[
\text{tr}(V - 1/tJV)^2 = \text{tr}(VV) - 2\text{tr}(VJV) + 1/t^2 \text{tr}(JVJV). \tag{5.2}
\]

Expressions (5.1) and (5.2) and in turn \( g_1 \) and \( h_1 \) can be evaluated using the following:

\[
\text{tr}(V) = t,
\]

\[
\text{tr}(JV) = \frac{t(1 - \rho^2) - 2\rho(1 - \rho^i)}{(1 - \rho)^2},
\]

\[
\text{tr}(VV) = \frac{1}{(1 - \rho^2)^2} \left[ t(1 - \rho^4) - 2\rho^2(1 - \rho^{2i}) \right],
\]

\[
\text{tr}(VJV) = \frac{1}{(1 - \rho)^2} \left[ t(1 + \rho)^2 + 2t\rho^{i+1} + 2\rho^2 \frac{1 - \rho^{2i}}{1 - \rho^2} - 4\rho(1 + \rho) \frac{1 - \rho^i}{1 - \rho} \right]
\]

\[
\text{tr}(JVJV) = \left[ \frac{t(1 - \rho^2) - 2\rho(1 - \rho^i)}{(1 - \rho)^2} \right]^2.
\]

### 5.1.1 Distributions of \( Q_3, Q_4 \) and \( Q_5 \)

In order to find the distributions of the matrix quadratic forms \( Q_3, Q_4 \) and \( Q_5 \) we need to estimate \( g_1 \) and \( h_1 \). In the above discussion we have given expressions for
$g_1$ and $h_1$ as functions of $\rho$. However since $\rho$ is unknown, we need to estimate the same. In the following we present maximum likelihood estimate of $\rho$.

Consider a random sample $Y_{ij}, j = 1, ..., n; i = 1, ..., g$ from a $pt$-variate normal distribution with mean $\mu_i$ (can be a linear function in a smaller dimension space) and variance covariance matrix $V \otimes \Sigma$. Here $V$ is an autoregressive structure matrix and $\Sigma$ is any positive definite matrix. Then the log-likelihood function can be written as,

\[
\ln l = \text{constant} - \frac{n}{2} \ln |V \otimes \Sigma| - \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} (Y_{ij} - \mu_i)' V^{-1} \otimes \Sigma^{-1} (Y_{ij} - \mu_i),
\]

Let $B = \sum_{i=1}^{g} \sum_{j=1}^{n_i} (Y_{ij} - \overline{Y}_i)(Y_{ij} - \overline{Y}_i)'$, with $\overline{Y}_i = \frac{1}{n_i} \sum_{j=1}^{n_i} Y_{ij}$. Then

\[
\ln l = \text{constant} - \frac{n}{2} \ln |V \otimes \Sigma| + \frac{1}{2} \text{tr}(V^{-1} \otimes \Sigma^{-1} B) - \frac{1}{2} \sum_{i=1}^{g} n_i (\overline{Y}_i - \mu_i)' V^{-1} \otimes \Sigma^{-1} (\overline{Y}_i - \mu_i). 
\]

Clearly the maximum likelihood estimate of $\mu_i = \overline{Y}_i$. By plugging this value for $\mu_i$ in (5.3) we get

\[
\ln l = - \frac{npt}{2} \ln(2\pi) - \frac{n}{2} \ln |V \otimes \Sigma| - \frac{1}{2} \text{tr}(V^{-1} \otimes \Sigma^{-1} B). 
\]

Recall that $|V| = (1 - \rho^2)^{t-1}$ and $V^{-1} = \frac{1}{1 - \rho^2} [I + \rho^2 C_1 - \rho C_2]$. Differentiating (5.4) with respect to $\rho$ and equating to zero we have,

\[
-\rho^3 2np(t-1) + \rho^2 A_3 + \rho (2np(t-1) - 2A_1 - 2A_2) + A_3 = 0, 
\]
where

\[
A_1 = \sum_{i=1}^{g} \sum_{j=1}^{n_i} (Y_{ij} - \bar{Y}_{i, i})'(I \otimes \Sigma^{-1})(Y_{ij} - \bar{Y}_{i, i}), \\
A_2 = \sum_{i=1}^{g} \sum_{j=1}^{n_i} (Y_{ij} - \bar{Y}_{i, i})'((C_1 \otimes \Sigma^{-1})(Y_{ij} - \bar{Y}_{i, i}), \\
A_3 = \sum_{i=1}^{g} \sum_{j=1}^{n_i} (Y_{ij} - \bar{Y}_{i, i})'((C_2 \otimes \Sigma^{-1})(Y_{ij} - \bar{Y}_{i, i}).
\]

The cubic equation in \( \rho \) given by (5.5) can easily be shown to have a unique root in the interval (-1,1). Hence it is easy to compute the MLE \( \hat{\rho} \), of \( \rho \). Let \( \hat{V} = V(\hat{\rho}) \).

Next, to find the maximum likelihood estimator of \( \Sigma \), we need to maximize (5.4) with respect to \( \Sigma \). In order to do that, partition the \( pt \times 1 \) vector \( (Y_{ij} - \bar{Y}_{i, i}) \) into \( t \) blocks of \( p \times 1 \) vectors such that \( (Y_{ij} - \bar{Y}_{i, i}) = (Y_{ij_1} - \bar{Y}_{i, 1}, \ldots, Y_{ij_t} - \bar{Y}_{i, t})' \). Using this partition of \( (Y_{ij} - \bar{Y}_{i, i}) \), we rewrite the likelihood (5.4) and maximize with respect to \( \Sigma \). This process yields,

\[
\hat{\Sigma} = \frac{1}{nt} \sum_{k=1}^{t} \sum_{l=1}^{t} \hat{\nu}_{kl} \sum_{i=1}^{g} \sum_{j=1}^{n_i} (Y_{ijk} - \bar{Y}_{i, k})(Y_{ijkl} - \bar{Y}_{i, l}),
\]

(5.6)

where \( \hat{\nu}_{kl} \) is the \( (kl) \)th element of \( \hat{V}^{-1} \).

By solving equations (5.5) and (5.6) iteratively we find \( \hat{V} \) and \( \hat{\Sigma} \), and then the estimates of \( g_1 \) and \( h_1 \). Thus we have approximate distributions of \( Q_3, Q_4 \) and \( Q_5 \).

### 5.2 Unbalanced multivariate repeated measures

So far, we have considered balanced case of multivariate repeated measures. That is, we have considered a set of \( t \) repeated measurements on \( p \) variables (or char-
acteristics) on each of the \( n \) individuals. Thus data on each individual is a \( p \times t \) matrix. However, in practice we do come across data where we may not have measurements on the individuals at all the \( t \) time points. Stated in other words, we have \( t_{ij} \) repeated measurements on \( p \) variables on the \( j^{th} \) individual from the \( i^{th} \) group. Now data on each individual is a \( p \times t_{ij} \) matrix. Let \( Y_{ijk} \) be a \( p \times 1 \) vector of observations on the \( j^{th} \) individual from the \( i^{th} \) group at the \( k^{th} \) occasion. Let \( Y'_{ij} = (Y'_{ij1}, \ldots, Y'_{ijt_{ij}}) \). Then \( Y_{ij} \) is a \( pt_{ij} \times 1 \) random observational vector corresponding to the \( j^{th} \) individual from the \( i^{th} \) group. Let us assume that \( \text{cov}(Y_{ij}) = V_{ij} \otimes \Sigma \), where \( V_{ij} \) is a \( t_{ij} \times t_{ij} \) positive definite matrix and \( \Sigma \) is a \( p \times p \) positive definite matrix. It appears that the covariance matrix \( V_{ij} \) is different for every \( i, j \). However this is not true. If \( V \) is a positive definite matrix of order \( t \times t \), where \( t \) is the maximum of \( t_{ij} \), then \( V_{ij} \) is the \( t_{ij} \times t_{ij} \) submatrix of \( V \). In applications, we assume \( V_{ij} \) for every \( i \) and \( j \), to have some structure depending on the same set of parameters, say \( \theta_1, \ldots, \theta_k \). In other words, \( V_{ij} \) depends on \( i \) and \( j \) only through its dimension. Analysis of this type of data is not very easy. In the following section, we describe a general linear model approach to analyze unbalanced data, assuming certain structures for \( V_{ij} \).
5.3 A linear model approach

Consider the situation where we have a set of unbalanced multivariate repeated measurements. Let \( Y_{ij} \) be the \( t_{ij} \times p \) matrix of observations on the \( j^{th} \) individual from the \( i^{th} \) group. Let us assume the following model for \( Y_{ij} \):

\[
Y_{ij} = X_{ij}B_i + E_{ij}, \quad i = 1, \ldots, g; \ j = 1, \ldots, n_i, \tag{5.7}
\]

where \( X_{ij} \) is \( t_{ij} \times m \) design matrix and \( B_i \) is \( m \times p \) matrix of unknown parameters, with \( \text{cov}(\text{vec } E_{ij}') = V_{ij} \otimes \Sigma \). Here \( V_{ij} \) is a positive definite matrix depending on only a few parameters, say \( \theta_1, \ldots, \theta_k \). If \( t_{ij} = t \) for all \( i \) and \( j \) (balanced data) then \( X_{ij} \) is a \( t \times m \) matrix, say \( X_i \). Further if \( X_i \) an identity matrix, then the model would contain a matrix of unstructured means, different for different groups.

In the unbalanced case, the interpretation of \( X_{ij} \) can be made as follows. Suppose \( G_{ij} \) is a \( t_{ij} \times t \) matrix of 0's and 1's such that it has 1 at the \((k, l_k)\)th position and 0 everywhere else, assuming that observations are available at the time points \( l_1, \ldots, l_{t_{ij}} \). Then we take \( X_{ij} = G_{ij}X_i \).

Let us stack all the \( \sum_{j} t_{ij} \) observations from the \( i^{th} \) group one below the other.
and write

\[
Y_i = \begin{bmatrix}
Y_{i1} \\
\vdots \\
Y_{ini}
\end{bmatrix},
X_i = \begin{bmatrix}
X_{i1} \\
\vdots \\
X_{ini}
\end{bmatrix}, \quad \text{and} \quad E_i = \begin{bmatrix}
E_{i1} \\
\vdots \\
E_{ini}
\end{bmatrix}.
\]

Then we have the multivariate linear model

\[
\begin{bmatrix}
Y_1 \\
\vdots \\
Y_g
\end{bmatrix} = \begin{bmatrix}
X_1 & 0 & \ldots & 0 \\
0 & X_2 & \ldots & 0 \\
\vdots & \vdots & \ddots & \vdots \\
0 & 0 & \ldots & X_g
\end{bmatrix} \begin{bmatrix}
B_1 \\
\vdots \\
B_g
\end{bmatrix} + \begin{bmatrix}
E_1 \\
\vdots \\
E_g
\end{bmatrix}.
\]

By writing \( n = \sum \sum t_{ij} \) and with other obvious notations we have the following multivariate regression model

\[
Y_{nxp} = X_{nxmg} B_{mgxp} + E_{nxp},
\]

with \( \text{cov}(\text{vec}(Y')) = W = \text{diag}(V_{11}, \ldots, V_{1n_1}, \ldots, V_{g n_g}) \otimes \Sigma. \)

From the usual multivariate regression theory it can be shown that

\[
\hat{B} = (X'W^{-1}X)^{-1}X'W^{-1}Y
\]

\[
= \left[ \sum \sum X_{ij}' V_{ij}^{-1} X_{ij} \right]^{-1} \left[ \sum \sum X_{ij}' V_{ij}^{-1} Y_{ij} \right]
\]

and \( \hat{\Sigma} = (Y - X\hat{B})'W^{-1}(Y - X\hat{B})/(n - m). \)
Using this \( \hat{\mathbf{B}} \) (say \( \theta_1, \ldots, \theta_k \) are known) any one of the MANOVA tests, for example Wilks' \( \Lambda \) for testing \( H_0 : \mathbf{LB} \mathbf{M} = 0 \) can be used. But \( \mathbf{V}_{ij} \) and in turn \( \mathbf{W} \) is usually unknown. Using some consistent estimators of \( \theta_1, \ldots, \theta_k \), \( \mathbf{V}_{ij} \) is estimated and the estimate \( \hat{\mathbf{W}} \) is used in place of \( \mathbf{W} \) in the formula for \( \hat{\mathbf{B}} \). The same MANOVA tests can be used, even in this case for testing \( H_0 : \mathbf{LB} \mathbf{M} = 0 \). In the following sections we assume equicorrelation and autoregressive structures for \( \mathbf{V}_{ij} \) and discuss estimation of parameters of variance covariance matrix. These two structures are commonly used in practice. Note that in both of these cases, \( \mathbf{V}_{ij} \) is a function of only one unknown parameter \( \rho \). We derive the maximum likelihood estimate of this unknown parameter and utilize it to perform the MANOVA.

### 5.3.1 Equicorrelation structure for \( \mathbf{V}_{ij} \)

Suppose we have an initial estimate of \( \mathbf{B} \). For example, an initial estimate of \( \mathbf{B} \) may be obtained by performing regression under \( \rho = 0 \). Then let \( \hat{\mathbf{E}}_{ij} = \mathbf{Y}_{ij} - \mathbf{X}_{ij} \hat{\mathbf{B}}_i \). We can assume that \( \hat{\mathbf{e}}_{ij} = \text{vec}(\hat{\mathbf{E}}_{ij}) \), \( i = 1, \ldots, g; \ j = 1, \ldots, n_i \), form a random sample from \( p_{t_{ij}} \)-variate normal distribution with zero mean vector and covariance matrix \( \mathbf{V}_{ij} \otimes \mathbf{\Sigma} \). Then the log-likelihood function can be written as

\[
\ln l = \text{constant} - \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \ln | \mathbf{V}_{ij} \otimes \mathbf{\Sigma} | \\
- \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \hat{\mathbf{e}}_{ij} \mathbf{V}_{ij}^{-1} \otimes \mathbf{\Sigma}^{-1} \hat{\mathbf{e}}_{ij} \tag{5.9}
\]
Next, partition the $p t_{ij} \times 1$ vector $\hat{\epsilon}_{ij}$ into $t_{ij}$ blocks of $p \times 1$ vectors such that $\hat{\epsilon}_{ij} = (\hat{\epsilon}_{ij1}, ..., \hat{\epsilon}_{ij t_{ij}})$. Using this partition rewrite the likelihood (5.9) as

\[
\ln l = \text{constant} - \frac{p}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \ln |V_{ij}| - \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \hat{t}_{ij} \ln |\Sigma| \\
- \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \sum_{k=1}^{t_{ij}} v_{ik}^* \hat{\epsilon}_{ijk} \Sigma^{-1} \hat{\epsilon}_{ijk}.
\]

(5.10)

Differentiating (5.10) with respect to $\Sigma$ and equating to zero we get

\[
\hat{\Sigma} = \frac{\sum_{i=1}^{g} \sum_{j=1}^{n_i} \sum_{k=1}^{t_{ij}} v_{ik}^* (\hat{\epsilon}_{ijk})(\hat{\epsilon}_{ijk})'}{\sum_{i=1}^{g} \sum_{j=1}^{n_i} t_{ij}}.
\]

(5.11)

Recall that $|V_{ij}| = (1 - \rho)^{t_{ij}-1}(1 + (t_{ij} - 1)\rho)$ and $v_{ik}^* = \frac{1 + (t_{ij} - 2)\rho}{[1 + (t_{ij} - 1)\rho](1 - \rho)}$ and $v_{ik} = \frac{\rho}{[1 + (t_{ij} - 1)\rho](1 - \rho)}$. Now using this we can rewrite (5.10) as

\[
\ln l = \text{constant} - \frac{p}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} (t_{ij} - 1) \ln(1 - \rho) - \frac{p}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \ln[1 + (t_{ij} - 1)\rho] \\
- \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} t_{ij} \ln |\Sigma| - \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} t_{ij} \frac{1 + (t_{ij} - 2)\rho}{[1 + (t_{ij} - 1)\rho](1 - \rho)} (\hat{\epsilon}_{ijk})'(\hat{\epsilon}_{ijk})^{-1} \\
- \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \sum_{k=1}^{t_{ij}} \frac{\rho}{[1 + (t_{ij} - 1)\rho](1 - \rho)} (\hat{\epsilon}_{ijk})'(\hat{\epsilon}_{ijk})^{-1} (\hat{\epsilon}_{ijk})
\]

(5.12)

Differentiating (5.12) with respect to $\rho$ and equating to zero yields,

\[
\frac{p}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \frac{(t_{ij} - 1)t_{ij}}{[1 + (t_{ij} - 1)\rho](1 - \rho)} \\
- \frac{\rho}{2(1 - \rho)^2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \sum_{k=1}^{t_{ij}} \sum_{l=1}^{t_{ij}} \frac{(t_{ij} - 1)[2 + (t_{ij} - 2)\rho]}{[1 + (t_{ij} - 1)\rho]} (\hat{\epsilon}_{ijk})'(\hat{\epsilon}_{ijk})^{-1} (\hat{\epsilon}_{ijk}) \\
+ \frac{1}{2(1 - \rho)^2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \sum_{k=1}^{t_{ij}} \frac{1 + (t_{ij} - 1)\rho^2}{[1 + (t_{ij} - 1)\rho]^2} (\hat{\epsilon}_{ijk})'(\hat{\epsilon}_{ijk})^{-1} (\hat{\epsilon}_{ijk}) = 0
\]

(5.13)
Solving (5.11) and (5.13) simultaneously, we get the estimates of $\Sigma$ and $\rho$. Now, with this new estimate of $\rho$ find a revised estimate of $B$. This two-stage estimation procedure of finding $\rho$ and $B$, is continued until the estimates are stabilized.

### 5.3.2 Autoregressive structure for $V_{ij}$

In this case, unlike in the equicorrelation case, the basic structure of $V_{ij}$ is distorted in the presence of missing observations. We may now consider two types of missing data. If the missing observations are only at the end (monotone data) then the basic form of $V_{ij}$ remains the same. That is $V_{ij} = (\rho^{i-j})$ where $i, j = 1, \ldots, t_{ij}$. However in the other forms of missing values, where the observations may be missing for any occasion the autoregressive structure leads to what is called a Markov structure. In the following we describe the method of estimating $\rho$ in both situations.

**Monotone data:** As noted before, monotone data refers to the unbalanced case where the data are missing from the last few occasions. As before, let us suppose that an initial estimate of $B$ is available. Then the form of the MLE of $\Sigma$ is same as that in (5.11). But now, $v_{kl}^*$ is the $(kl)^{th}$ element of $V_{ij}^{-1}$ given by

$$ V_{ij}^{-1} = \frac{1}{(1 - \rho^2)}[I_{t_{ij}} + \rho^2 C_{1ij} - \rho C_{2ij}] $$

where $C_{1ij}$ and $C_{2ij}$ are $t_{ij} \times t_{ij}$ matrices with similar structures as described in the Appendix of Chapter 2. Next, using, $|V_{ij}| = (1 - \rho^2)^{t_{ij}-1}$ and $V_{ij}^{-1}$ rewrite the
log-likelihood function given by (5.10) as

\[ L = \text{constant} - \frac{p}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} (t_{ij} - 1) \ln(1 - \rho^2) - \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} t_{ij} \ln|\Sigma| \]

\[ - \frac{1}{2(1 - \rho^2)} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \tilde{e}_{ij}^2 [(I_{ij} \otimes \Sigma^{-1} + \rho^2 C_{1ij} \otimes \Sigma^{-1} - \rho C_{2ij} \otimes \Sigma^{-1})] \tilde{e}_{ij} \] (5.14)

Differentiating (5.14) with respect to \( \rho \) and equating to zero we get,

\[-2\rho^3 pn + \rho^2 A_3 + 2\rho(pn - A_1 - A_2) + A_3 = 0, \] (5.15)

where

\[ n = \sum_{i=1}^{g} \sum_{j=1}^{n_i} (t_{ij} - 1) \]

\[ A_1 = \sum_{i=1}^{g} \sum_{j=1}^{n_i} \tilde{e}_{ij}^2 (I_{ij} \otimes \Sigma^{-1}) \tilde{e}_{ij} \]

\[ A_2 = \sum_{i=1}^{g} \sum_{j=1}^{n_i} \tilde{e}_{ij}^2 (C_{1ij} \otimes \Sigma^{-1}) \tilde{e}_{ij} \]

\[ A_3 = \sum_{i=1}^{g} \sum_{j=1}^{n_i} \tilde{e}_{ij}^2 (C_{2ij} \otimes \Sigma^{-1}) \tilde{e}_{ij} \]

The cubic equation (5.15) is similar to the one studied by Prabhala (1995) in the context of growth curve models. It has been proved that the equation has a unique root in the interval (-1,1). Solving (5.15) and (5.11) with appropriate \( v_{kl} \) simultaneously we get the estimates of \( \rho \) and \( \Sigma \).

**General unbalanced data:** We now consider a model that can be used for analyzing data that are missing from any of the intermediate occasions as well. Let \( Y_{ij} \) be the \( t_{ij} \times p \) matrix of observations on the \( j^{th} \) individual from the \( i^{th} \) group.
such that $Y_{ij} = G_{ij} \tilde{Y}_{ij}$. Here $\tilde{Y}_{ij}$ is a $t \times p$ matrix of measurements, $t = \max(t_{ij})$ and $G_{ij}$ is a matrix of order $t_{ij} \times t$ as defined in the beginning of section 5.3.

Let us assume the following model for $Y_{ij}$

$$Y_{ij} = X_{ij}B_i + E_{ij}, \quad i = 1, \ldots, g; \quad j = 1, \ldots, n_i,$$

(5.16)

where $X_{ij}$ is $t_{ij} \times m$ design matrix and $B_i$ is $m \times p$ matrix of unknown parameters.

Then $\text{cov}(\text{vec}(E_{ij})) = G_{ij}V'G_{ij} \otimes \Sigma$, where $V$ is a $t \times t$ matrix having an autoregressive structure. Then we can verify that $G_{ij}V'G_{ij}$ has a Markov structure, that is $G_{ij}V'G_{ij} = V_{ij} = (\rho^{t_{ij} - t_{ijk}}, k, k' = 1, \ldots, t_{ij})$, where $t_{ijk}$'s are the consecutive time points where the observations are made on the $j^{th}$ individual from the $i^{th}$ group.

Then $V_{ij}$ is of the form

$$V_i(\rho) = \begin{pmatrix}
1 & \rho^{t_{ij2}-t_{ij1}} & \rho^{t_{ij3}-t_{ij1}} & \cdots & \rho^{t_{ijt_{ij1}}-t_{ij1}} \\
\rho^{t_{ij2}-t_{ij1}} & 1 & \rho^{t_{ij3}-t_{ij2}} & \cdots & \rho^{t_{ijt_{ij2}}-t_{ij2}} \\
. & . & . & \cdots & . \\
. & . & . & \cdots & . \\
. & . & . & \cdots & . \\
\rho^{t_{ijt_{ij1}}-t_{ij1}} & \rho^{t_{ijt_{ij2}}-t_{ij2}} & \cdots & . & 1
\end{pmatrix}$$

The determinant and the inverse of this matrix are given below. Let $d_{ijk} = t_{ij(k+1)} - t_{ijk}$, and $f_{ijk} = \frac{1}{1 - \rho^2 t_{ijk}}$, $k = 1, \ldots, t_{ij} - 1$. Then

$$|V_{ij}| = (f_{ij1}f_{ij2}\ldots f_{ijt_{ij}-1})^{-1}$$

and $V_{ij}^{-1} = \ldots$
Let \( \hat{\mathbf{E}}_{ij} = \mathbf{Y}_{ij} - X_{ij} \mathbf{B}_i \). The log-likelihood function of \( \hat{\mathbf{e}}_{ij} = \text{vec}(\hat{\mathbf{E}}_{ij}) \), \( i = 1, \ldots, g \), \( j = 1, \ldots, n_i \), is

\[
\ln l = \text{constant} - \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \ln |\mathbf{V}_{ij} \otimes \bm{\Sigma}| - \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \hat{\mathbf{e}}_{ij} \mathbf{V}_{ij}^{-1} \otimes \bm{\Sigma}^{-1} \hat{\mathbf{e}}_{ij}. \tag{5.17}
\]

Following steps similar to what we did in the previous section, it is easy to see that

\[
\hat{\bm{\Sigma}} = \frac{\sum_{i=1}^{g} \sum_{j=1}^{n_i} \sum_{t_i=1}^{t_{ij}} \sum_{t_j=1}^{t_{ij}} v_{ki}(\hat{\mathbf{e}}_{ijk})(\hat{\mathbf{e}}_{ij})'}{\sum_{i=1}^{g} \sum_{j=1}^{n_i} t_{ij}}. \tag{5.18}
\]

Next, differentiating \( \ln l \) with respect to \( \rho \) and equating to zero, we get

\[
-\frac{p}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \sum_{k=1}^{t_{ij}-1} 2d_{ijk} \rho^{2d_{ijk}-1} \frac{1}{1 - \rho^{2d_{ijk}}} - \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \hat{\mathbf{e}}_{ij} \frac{\partial}{\partial \rho} [\mathbf{V}_{ij}^{-1} \otimes \bm{\Sigma}^{-1}] \hat{\mathbf{e}}_{ij} = 0
\]

that is,

\[
-\frac{p}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \sum_{k=1}^{t_{ij}-1} 2d_{ijk} \rho^{2d_{ijk}-1} \frac{1}{1 - \rho^{2d_{ijk}}} - \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \text{tr} \left( \frac{\partial}{\partial \rho} [\mathbf{V}_{ij}^{-1} \otimes \bm{\Sigma}^{-1}] \hat{\mathbf{e}}_{ij} \hat{\mathbf{e}}_{ij}' \right) = 0.
\]
i.e.,

\[ - \frac{p}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \sum_{k=1}^{t_{ij}} 2d_{ijk} \rho^{2d_{ijk}-1} - \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \sum_{k=1}^{t_{ij}} v_{kk} tr(\Sigma^{-1} \epsilon_{ijk} \epsilon'_{ijk}) \]

\[ + \frac{1}{2} \sum_{i=1}^{g} \sum_{j=1}^{n_i} \sum_{k=1}^{t_{ij}} v_{kk+1} tr(\Sigma^{-1} \epsilon_{ijk} \epsilon'_{ijk+1}) \]

(5.19)

where \( v_{kk} \) is the \( k^{th} \) diagonal element of \( \frac{\partial}{\partial \rho} V_{ij}^{-1} \) and \( v_{kk+1} \) is the \((k, k+1)^{st}\) off-diagonal element of \( \frac{\partial}{\partial \rho} V_{ij}^{-1} \). These elements are calculated below.

To evaluate \( \frac{\partial V_{ij}^{-1}}{\partial \rho} \), consider

\[ \frac{\partial f_{ijk}}{\partial \rho} = \frac{\partial}{\partial \rho} \left( \frac{1}{1 - \rho^{2d_{ijk}}} \right) = \frac{2d_{ijk} \rho^{2d_{ijk}-1}}{(1 - \rho^{2d_{ijk}})^2} = f_{ijk}^2 2d_{ijk} \rho^{2d_{ijk}-1}. \]

Thus the diagonal elements of \( \frac{\partial}{\partial \rho} V_{ij}^{-1} \) are:

\[ (f_{ij1}^2 2d_{ij1} \rho^{2d_{ij1}-1}, f_{ij2}^2 2d_{ij2} \rho^{2d_{ij2}-1}, ..., f_{ij(t_{ij}-1)2d_{ij(t_{ij}-1)}-1}^2 2d_{ij(t_{ij}-1)} \rho^{2d_{ij(t_{ij}-1)}-1}). \]

Also,

\[ \frac{\partial (f_{ijk} \rho^{d_{ijk}})}{\partial \rho} = f_{ijk} d_{ijk} \rho^{d_{ijk}-1} + \rho^{d_{ijk}} \frac{\partial f_{ijk}}{\partial \rho} \]

\[ = f_{ijk} d_{ijk} \rho^{d_{ijk}-1} + \rho^{d_{ijk}} f_{ijk}^2 2d_{ijk} \rho^{2d_{ijk}-1} \]

\[ = d_{ijk} \rho^{d_{ijk}-1} f_{ijk} \left( 1 + \frac{2\rho^{2d_{ijk}}}{1 - \rho^{2d_{ijk}}} \right) \]

\[ = d_{ijk} \rho^{d_{ijk}-1} f_{ijk} \left( 1 + \rho^{2d_{ijk}} \right) \]

\[ = d_{ijk} \rho^{d_{ijk}-1} f_{ijk}^2 (1 + \rho^{2d_{ijk}}). \]

Therefore, the off-diagonal elements of \( \frac{\partial}{\partial \rho} V_{ij}^{-1} \) are:
(d_{ij1} \rho^{d_{ij1} - 1} f_{ij1}^2 (1 + \rho^{2d_{ij1}}), d_{ij2} \rho^{d_{ij2} - 1} f_{ij2}^2 (1 + \rho^{2d_{ij2}}), ..., d_{ijs} \rho^{d_{ijs} - 1} f_{ijs}^2 (1 + \rho^{2d_{ijs}})).

Plugging these values in (5.19) we can solve for \( \rho \). As before an iterative scheme can be developed to find the ML estimates of \( B \), \( \rho \) and \( \Sigma \).
Bibliography


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108


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