MANOVA with singular variance matrix

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Abstract. Classical multivariate analysis of variance for \( p \) response variables is extended to cover high-dimensional data. For example, data often comprise many response variables that may be related. Therefore, inference based on all the response variables may be inefficient. However, the relationship between the response variables is usually not known. This leads to the assumption that the \( p \) response variables span a linear space of some fixed dimension, say \( r < p \); equivalently the \( p \times p \) variance matrix is singular of rank \( r \). We will assume that the rank is given. Following the classical approach of doing inference in linear models, parameters are first estimated and thereafter tests are constructed. Estimators and tests are based on the likelihood method. The present model differs from the classical multivariate analysis of variance model and consists of a deterministic part and a random part. It is noticed that the classical approach is a special case of the one which will be considered in this article.

1. Introduction

Often data comprise observations on many response variables of which many may be closely related. Therefore, inference based on all the response variables may be inefficient. In situations like this, usually the principal component method is applied to reduce the dimensionality of the response variables. However, after reducing the dimensionality, it is not known how to carry out the statistical inference on the unknown parameters. For example, in the one-sample problem, it is not known how to estimate the mean vector.
or how to carry out tests of hypotheses regarding the parameters of the mean vector. In this paper, we assume that the rank of the variance matrix is the same as the number of selected principal components. Thus, we shall assume that the variance matrix of the observations is a singular matrix of a given rank, the rank of the matrix is either determined by the past experience or from a practical consideration such as principal component analysis. We shall consider the multivariate analysis of variance model. Estimators and tests of hypotheses along with their distributions will be given.

Linear models with the singular variance matrix have been studied by many authors including Khatri (1968), Mitra & Rao (1968), Rao (1973), Alalouf (1978), Feuerverger & Fraser (1980) and Nordström (1985). Except Khatri (1968) these authors assume that the variance matrix $\Sigma$ is known. In this paper this matrix is unknown and therefore has to be estimated. Suppose that the variance matrix of the response variables, $\Sigma : p \times p$, is of rank $r$ which is a dimensionality reduction if $r < p$, and based on this assumption a MANOVA is performed. The results of this paper will be illustrated by a numerical example. The organization of the paper is as follows. In Section 2 we consider a numerical example to demonstrate the singularity of the variance matrix. Some of its eigenvalues are very small and we suggest that instead of considering initial eight-dimensional data, we will be better off with less than eight. In Section 3 we obtain the maximum likelihood estimators of the parameters and likelihood ratio tests are used to complete the analysis. In Section 4 confidence intervals for mean parameter $\Sigma$ are found and Section 5 is dedicated to the detection of outliers.

2. A numerical example

We consider the data of Russell et al. (1967), presented in Srivastava & Carter (1983, p. 128). The data present the clinical analysis of soil characteristics for three contours and four depths of soil. The experiment area was divided into four blocks and samples were taken randomly at various depths and contours. For our purpose we just consider four combinations of depths and contours. Moreover, the four areas are reduced to two by merging the original areas pairwise. The data are presented in Table 2.1.

According to Table 2.1 the multivariate data are obtained from a complete balanced block design. To be even more specific, there are two blocking factors of which one (Area) has two levels and the other (Group) has four levels. Furthermore, there are 16 independent observations of which each comprises 8 response variables.
Table 2.1. Soil data where the following abbreviations are used: $V_1$=pH; $V_2$=Total nitrogen (%); $V_3$=Bulk density (gm/cm$^3$); $V_4$=Total phosphorus (ppm); $V_5$=Exchangeable + soluble calcium (me/100 gm); $V_6$=Exchangeable + soluble magnesium (me/100 gm); $V_7$=Exchangeable + soluble potassium (me/100 gm); $V_8$=Exchangeable + soluble sodium (me/100 gm);

<table>
<thead>
<tr>
<th></th>
<th>5.40</th>
<th>5.65</th>
<th>5.14</th>
<th>5.14</th>
<th>5.14</th>
<th>5.10</th>
<th>4.70</th>
<th>4.46</th>
</tr>
</thead>
<tbody>
<tr>
<td>$V_2$</td>
<td>0.19</td>
<td>0.16</td>
<td>0.26</td>
<td>0.17</td>
<td>0.16</td>
<td>0.09</td>
<td>0.10</td>
<td>0.11</td>
</tr>
<tr>
<td>$V_3$</td>
<td>0.92</td>
<td>1.04</td>
<td>0.95</td>
<td>1.10</td>
<td>1.12</td>
<td>1.22</td>
<td>1.52</td>
<td>1.47</td>
</tr>
<tr>
<td>$V_4$</td>
<td>215</td>
<td>208</td>
<td>300</td>
<td>248</td>
<td>174</td>
<td>129</td>
<td>117</td>
<td>170</td>
</tr>
<tr>
<td>$V_5$</td>
<td>16.4</td>
<td>12.2</td>
<td>13.0</td>
<td>11.9</td>
<td>14.2</td>
<td>8.6</td>
<td>8.7</td>
<td>9.5</td>
</tr>
<tr>
<td>$V_6$</td>
<td>7.6</td>
<td>5.2</td>
<td>5.7</td>
<td>7.0</td>
<td>8.1</td>
<td>6.9</td>
<td>8.2</td>
<td>9.2</td>
</tr>
<tr>
<td>$V_7$</td>
<td>0.72</td>
<td>0.71</td>
<td>0.68</td>
<td>1.09</td>
<td>0.70</td>
<td>0.81</td>
<td>0.39</td>
<td>0.70</td>
</tr>
<tr>
<td>$V_8$</td>
<td>1.14</td>
<td>0.94</td>
<td>0.60</td>
<td>1.01</td>
<td>2.17</td>
<td>2.67</td>
<td>3.32</td>
<td>3.76</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Group</th>
<th>Area</th>
<th>1</th>
<th>1</th>
<th>1</th>
<th>1</th>
<th>2</th>
<th>2</th>
<th>2</th>
<th>2</th>
<th>1</th>
<th>1</th>
<th>2</th>
<th>2</th>
</tr>
</thead>
</table>

Let the data be collected in $Y: 8 \times 16$ where the first 4 columns consist of the data from group 1, the next 4 columns consist of data from group 2, etc. The design matrix $X$ is given by

$$
X = \begin{pmatrix}
1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 \\
1 & 1 & -1 & -1 & 1 & -1 & -1 & 1 & 1 & -1 & -1 & 1 & -1 & -1 & 1 & -1 \\
1 & 1 & 1 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & -1 & -1 & -1 & -1 & -1 \\
0 & 0 & 0 & 0 & 1 & 1 & 1 & 1 & 0 & 0 & 0 & 0 & -1 & -1 & -1 & -1 & -1 \\
0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 1 & 1 & 1 & -1 & -1 & -1 & -1 & -1
\end{pmatrix}
$$

and the following model is assumed to hold:

$$
Y = \Xi X + \Sigma^{1/2} E,
$$

where the columns of $E$ are independent identically distributed (i.i.d.) random
vectors with the distribution \( N_6(0, I) \), and \( \Sigma^{1/2} \) is a positive semi-definite square root of \( \Sigma \). The sums-of-squares matrix due to error, i.e. \( S \), is calculated by (3.7):

\[
S = \begin{pmatrix}
0.20 & -0.01 & -0.01 & -14.5 & -0.38 & -1.42 & -0.09 & -0.80 \\
-0.01 & 0.01 & -0.03 & 8.00 & 0.42 & 0.12 & -0.00 & -0.05 \\
-0.01 & -0.03 & 0.26 & -18.1 & -2.34 & -1.33 & 0.02 & -0.03 \\
-14.5 & 8.00 & -18.1 & 8920 & 168.0 & 80.6 & 18.7 & -58.0 \\
-0.38 & 0.42 & -2.34 & 42.4 & 20.4 & -0.73 & 0.97 & 0.35 & 0.24 \\
-1.42 & 0.12 & -1.33 & 80.6 & 20.4 & 26.8 & 0.97 & 10.1 \\
-0.09 & -0.00 & 0.02 & 18.7 & -0.73 & 0.97 & 0.35 & 0.24 \\
-0.80 & -0.05 & -0.03 & -58.0 & 4.01 & 10.1 & 0.24 & 10.8 \\
\end{pmatrix}
\]

An unbiased estimator of the sample variance matrix is \( W = \frac{1}{n}S \) with eigenvalues

\[
811.31 \quad 5.02 \quad 1.50 \quad 0.40 \quad 0.017 \quad 0.013 \quad 0.0027 \quad 0.000039.
\]

Since the smallest eigenvalue is very close to zero, it is reasonable to assume that the rank of \( S \) is at most 7. It follows from Lemma 3.1, given in Section 3, that the rank of \( W \) is equal to the rank of \( \Sigma \). The second smallest eigenvalue is 0.003 which is also close to zero. Thus, it may be reasonable to assume that the rank of \( \Sigma \) is 6. The problem of deciding on the rank of \( \Sigma \) is rather difficult. For example, one may argue that since all the four smallest eigenvalues are rather small, the rank of \( \Sigma \) is only four. We may also note that the first eigenvalue contains 99% of the variation in the data, and thus one may argue that the rank of \( \Sigma \) is only one. The fewer components are chosen, the fewer linear combinations of the mean can be tested for significance; the difference can be detected only if it lies in the direction of the chosen components. So, there is a difference between the so-called principal component analysis and the problem of drawing inference on its mean components. Any inference based on one component only or on very few components may be very misleading. Here our objective is to select the rank of the variance matrix \( \Sigma \). For this, we should drop from considerations only those components for which the corresponding eigenvalues of the sample variance matrix are zero or very close to zero. In Table 2.2, however, we examine the effect of various variance assumptions on the rank of \( \Sigma \).

We are going to test that there is no area effect in our example, i.e.,

\[
H : \Gamma' Ec = 0 \quad \text{versus} \quad A : \Gamma' Ec \neq 0,
\]

(2.1)

where

\[
e' = (0 \quad 1 \quad 0 \quad 0 \quad 0)
\]

and \( \Gamma \) is as in (3.2). Let \( C(Z) \) denote the column vector space generated by
the columns of $Z$. Clearly $\mathcal{C}(c) \subseteq \mathcal{C}(X)$ as $r(X) = 5$ and we will utilize Theorem 3.3. However, note that $r(c) = 1$ and therefore the exact distribution for $\lambda$ in the theorem is available, i.e.

$$F = \frac{n - r(X) + 1 - r - 1 - \lambda}{\lambda} \sim F_{r, n - r(X) + 1 - r}, \quad (2.2)$$

where $r = r(\Sigma)$. In Table 2.2 the results of this test under various assumptions on $r(\Sigma)$ are presented.

<table>
<thead>
<tr>
<th>$r(\Sigma)$</th>
<th>8</th>
<th>7</th>
<th>6</th>
<th>5</th>
<th>4</th>
<th>3</th>
<th>2</th>
<th>1</th>
</tr>
</thead>
<tbody>
<tr>
<td>$\lambda$</td>
<td>0.16</td>
<td>0.16</td>
<td>0.20</td>
<td>0.21</td>
<td>0.63</td>
<td>0.65</td>
<td>0.96</td>
<td>0.98</td>
</tr>
<tr>
<td>$F$</td>
<td>2.64</td>
<td>3.76</td>
<td>4.09</td>
<td>5.16</td>
<td>1.18</td>
<td>1.61</td>
<td>0.19</td>
<td>0.22</td>
</tr>
<tr>
<td>$P$</td>
<td>0.18</td>
<td>0.082</td>
<td>0.055</td>
<td>0.027</td>
<td>0.39</td>
<td>0.25</td>
<td>0.83</td>
<td>0.65</td>
</tr>
</tbody>
</table>

Thus, it follows from Table 2.2 that when we do not put any rank restrictions on $\Sigma$, i.e. $r(\Sigma) = 8$, the hypothesis $H$ in (2.1) can not be rejected. However, under assumptions that $\Sigma$ is singular of rank 6 the hypothesis is rejected at 5.5% level of significance. Our conclusions are that there exist some differences between the areas which we certainly can not draw if just performing a test when $\Sigma$ is of full rank and which is the standard test in MANOVA. For comparison, each variable has been analyzed separately by performing a univariate ANOVA. The results are presented in Table 2.3.

<table>
<thead>
<tr>
<th>$V1$</th>
<th>$V2$</th>
<th>$V3$</th>
<th>$V4$</th>
<th>$V5$</th>
<th>$V6$</th>
<th>$V7$</th>
<th>$V8$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$P$</td>
<td>0.0002</td>
<td>1.00</td>
<td>0.15</td>
<td>0.64</td>
<td>0.32</td>
<td>0.51</td>
<td>0.09</td>
</tr>
</tbody>
</table>

Hence, from Table 2.3 it follows that there seem to be some differences between the groups although they are not very pronounced. Vaguely speaking, one may say that these differences were found in the multivariate approach under the assumption that the covariance matrix was not of full rank.
3. Tests and estimators of parameters in MANOVA with singular covariance

In this section, we derive the maximum likelihood estimators of the parameters and apply the likelihood ratio test for the general hypothesis.

The general MANOVA model can be written

$$\mathbf{Y} = \Xi \mathbf{X} + \Sigma^{\frac{1}{2}} \mathbf{E},$$  \hspace{1cm} (3.1)

where $\mathbf{Y}$: $p \times n$, $\Xi$: $p \times k$, $\mathbf{X}$: $k \times n$, the rank of $\Sigma$ is denoted $r(\Sigma) = r \leq p$, $n - r(\mathbf{X}) \geq r$, the columns of $\mathbf{E}$ are i.i.d. $N_p(0, I)$ and $\Sigma^{\frac{1}{2}}$ is the positive semi-definite square root of $\Sigma$. It is assumed that $\Xi$ and $\Sigma$ are unknown while $\mathbf{X}$ is a matrix of known constants. As it has been noted by several authors (e.g., Rao 1973, p. 297; Nordström 1985) the singularity in $\Sigma$ implies certain restrictions on the parameter space as well as the observation matrix.

By performing a one-one transformation of (3.1) this can be explored in some detail:

$$\begin{pmatrix} \Gamma_0' \\ \Gamma_0 \end{pmatrix} \mathbf{Y} = \begin{pmatrix} \Gamma_0' \\ \Gamma_0 \end{pmatrix} \Xi \mathbf{X} + \begin{pmatrix} \Gamma_0' \Sigma^{\frac{1}{2}} \mathbf{E} \\ 0 \end{pmatrix},$$  \hspace{1cm} (3.2)

where $(\Gamma : \Gamma_0)$ spans the whole space, $\Gamma_0' \Gamma_0 = 0$, $\Gamma_0' \Xi = 0$, $\Gamma_0' \Gamma = I_r$, $\Gamma : p \times r$, $\Xi = \Gamma \Lambda \Gamma'$, where $\Lambda = (\lambda_i)$ is the diagonal matrix of positive and ordered eigenvalues of $\Xi$, $\Gamma_0 : p \times (p - r)$ and $\Gamma_0' \Gamma_0 = I_{p-r}$. Throughout, a matrix $\Lambda_0$ will be any matrix which satisfies $C(\Lambda_0) = C(\Lambda)^\perp$.

From (3.2) it follows with probability 1 that

$$\Gamma_0' \mathbf{Y} = \Gamma_0' \Xi \mathbf{X}.$$  \hspace{1cm} (3.3)

Equation (3.3) may be viewed as a linear system of equations in $\Xi$. From Rao & Mitra (1971, Theorem 2.3.2, p. 24) a necessary and sufficient condition for the equation (3.3) to have a solution is that

$$\Gamma_0'(\Gamma_0')^+ \Gamma_0' \mathbf{Y} = \Gamma_0' \mathbf{Y},$$  \hspace{1cm} (3.4)

where $\mathbf{A}^+$ denotes the Moore–Penrose inverse of $\mathbf{A}$. That is $\mathbf{A}^+$ satisfies the four conditions (i) $\mathbf{AA}^+ \mathbf{A} = \mathbf{A}$, (ii) $\mathbf{A}^+ \mathbf{AA}^+ = \mathbf{A}^+$, (iii) $\mathbf{A}^+ \mathbf{A} = (\mathbf{A}^+ \mathbf{A})^\prime$ and (iv) $\mathbf{AA}^+ = (\mathbf{AA}^\prime)^\prime$. It may be noted that in the above equation we may use any generalized inverse of $\mathbf{A}$, say $\mathbf{A}^-$, satisfying only the condition $\mathbf{AA}^- \mathbf{A} = \mathbf{A}$. However, the Moore–Penrose inverse is unique. Since

$$\Gamma_0' \Gamma_0 \Gamma_0' = \Gamma_0', \quad \Gamma_0' \Gamma_0 \Gamma_0 = \Gamma_0,$$

and $\Gamma_0' \Gamma_0$ and $\Gamma_0' \Gamma_0$ are symmetric matrices, it follows that $\Gamma_0 : (p - r) \times p$ is the Moore–Penrose inverse of $\Gamma_0'$. It can also be shown that the Moore–Penrose inverse of $\mathbf{X}$ is given by

$$\mathbf{X}^+ = \mathbf{X}'(\mathbf{XX}')^+,$$
where it is known that \((XX')^+ = X^+X^+\), \((X')^+ = (X^+)^\prime\) and \((X^+)^\prime = X\).

Then, the left-side of (3.4) becomes

\[
\Gamma_0'YX^+X = (\Gamma_0'\Xi X)X^+X = \Gamma_0'\Xi X,
\]

which is equal to \(\Gamma_0'Y\), from (3.3). Thus, the necessary and sufficient condition is satisfied. From equation (2.3.5) of Rao & Mitra (1971, p. 24), a general solution written in terms of Moore–Penrose inverses is given by

\[
\Xi = \Gamma_0\Gamma_0'YX^+ + U - \Gamma_0\Gamma_0'UXX^+, \tag{3.5}
\]

where \(U: p \times k\) is an arbitrary matrix of parameters, which is of the same dimension as \(\Xi\). Since

\[
\Gamma_0\Gamma_0'\Gamma a\Gamma' = I_p,
\]

the general solution can be rewritten as

\[
\Xi = \Gamma_0\Gamma_0'YX^+\Gamma a\Theta + \Gamma_0\eta(I - XX^+), \tag{3.6}
\]

where \(\Theta = \Gamma_0'U : r \times k\), and \(\eta = \Gamma_0'U : (p - r) \times k\). It may be noted that if the \(k \times n\) matrix \(X\) is of full rank \(k \leq n\), then \(XX^+ = I_k\) and the third term on the right-side of (3.4) vanishes. Thus, the \((p - r) \times k\) matrix \(\eta\) is an arbitrary matrix due to the rank deficiency in \(X\), while the \(r \times k\) matrix \(\Theta\) is a matrix of parameters that can be estimated from the random part of the model (3.2).

The solution (3.5) has been obtained from the deterministic part of the model (3.2) and does not affect the random part of (3.2) as can be seen by substituting the solution into (3.2). The random part of the model is given by

\[
\Gamma'Y = \Theta X + \Gamma'\Sigma^{1/2}E. \tag{3.6}
\]

In (3.6) the dispersion matrix \(\Gamma'\Sigma\Gamma\) is non-singular as we shall show later. Since \(\Gamma\) is unknown, it is not obvious that estimation can be carried out. However, as will be seen from the following lemma, the known sums-of-squares matrix

\[
S = Y(I - X'(XX')^{-1}X)Y' \tag{3.7}
\]

gives some insight in \(C(\Gamma) = C(\Sigma)\) which turns out to be crucial for estimation and testing.

In the subsequent \(W_p(\bullet, n)\) represents the Wishart distribution with \(n\) degrees of freedom.

**Lemma 3.1.** Let \(S \sim W_p(\Sigma, n - r(X))\). Then, with probability 1, \(C(S) \subseteq C(\Sigma)\) and if \(n - r(X) \geq r = r(\Sigma)\), \(C(S) = C(\Sigma)\).

**Proof.** We may note that \(X^+X\) is a symmetric matrix. That is, \(X^+X = (X'X^+)'\). Hence

\[
S = Y(I_n - X^+X)Y' = Y(I_n - X^+X)(I_n - X^+X)'Y'.
\]
Thus,
\[ C(S) \subseteq C(Y(I_n - X^+X)) = C(\Sigma X + \Sigma^2 E)(I_n - X^+X) \]
\[ = C(\Sigma^{1/2} E(I_n - X^+X)) \subseteq C(\Sigma^{1/2}) = C(\Sigma). \]

Hence, the rank of \( S \), \( r(S) \leq r(\Sigma) = r \), that is the rank of the covariance matrix \( S \) is always less or equal to the rank of \( \Sigma \). However, to prove that the rank of \( S \) is exactly equal to \( r = r(\Sigma) \) when \( r \leq n \), we use the assumption that \( S \) is Wishart distributed. Since \( \Sigma \) is of rank \( r \) we can write \( \Sigma = BB' \), where \( B \) is a \( p \times r \) matrix of rank \( r \) (see Srivastava & Khatri, 1979, Definition 2.1.1, p. 43). Hence, under the assumption of Wishartness, \( S \) can be written \( S = BZZ'B' \), where \( Z = (z_1, \ldots, z_n) \) is an \( r \times n \) matrix and \( z_i \)'s are i.i.d. \( N_r(0, I) \). Since \( n \geq r \), the rank of the matrix \( Z \) is \( r \), with probability one (see Srivastava & Khatri 1979, p. 73). Hence,
\[ r(S) = r(BZ) = r(B) = r, \]
from Corollary 1.5.3 of Srivastava & Khatri (1979, p. 10).

**Corollary 3.1L.** If \( S = HLH' \) where \( H'H = I_r \), then \( \Gamma = HQ \) where \( Q \) is an \( r \times r \) orthogonal matrix.

From the above result, it follows that \( \Gamma_0'S\Gamma_0 = 0 \) with probability one and thus \( \Gamma_0 \) may be considered as known. Next we show that the matrix \( \Gamma'_\Sigma^{1/2} E \) occurring in (3.6) is of rank \( r = r(\Sigma) \). We note that since \( \Sigma^{1/2} = \Gamma A^{1/2} \Gamma' \), and \( \Gamma' \Gamma = I_r \),
\[ \Gamma'_\Sigma^{1/2} E = \Lambda^{1/2} \Gamma'E, \]
where the columns of \( E \) are i.i.d. \( N_r(0, I) \). Thus, the \( n \) columns of \( \Gamma'E \) are i.i.d. \( N_r(0, I) \) and hence as above the rank is \( r \). Thus
\[ r(\Gamma'_\Sigma^{1/2} E) = r(\Gamma'E) = r \]
as \( \Lambda^{1/2} \) is nonsingular. From this it follows that
\[ (\Gamma'_Y - \Theta X)(\Gamma'_Y - \Theta X)' \]
is nonsingular with probability one. In the model (3.6), the matrices of unknown parameters are \( \Gamma, \Theta \) and \( A \) as
\[ \Gamma'_\Sigma^{1/2} = \Lambda^{1/2} \Gamma'. \]

Thus, the likelihood function for these parameters is given by
\[ L(\Theta, \Gamma, A) = c_0 |A|^{-\frac{n}{2}} \text{etr}\left( -\frac{1}{2} A^{-1}(\Gamma'_Y - \Theta X)(\Gamma'_Y - \Theta X)' \right), \]
ca. \( c_0 = (2\pi)^{-\frac{nr}{2}} \),
(3.8)
where \( \text{etr}(A) \) stands for the exponential of the trace of the matrix \( A \). From Srivastava & von Rosen (2002) we get the following theorem.
Theorem 3.1. Assume that $n - k \geq r(\Sigma)$. Then the maximum likelihood estimators of the parameters are given by
\[
\hat{\Theta} = H, \\
\hat{\Lambda} = L/n, \\
\hat{\Sigma} = S/n, \\
\hat{\Theta} = H'YY'(XX')^+ + U_0(I - XX^+), \\
\hat{\Xi} = YY^+X, \\
\hat{\Xi} = YY^+ + HU_0(I - XX^+),
\]
where $U_0$ is an arbitrary matrix.

Corollary 3.1. Assume that the $k \times n$ matrix $X$ is of full rank. Then the maximum likelihood estimators of $\Theta$ and $\Xi$ are given by
\[
\hat{\Theta} = H'YY'(XX)^{-1}, \\
\hat{\Xi} = YY'(XX)^{-1},
\]
respectively, and all the other estimators remain the same as in Theorem 3.1.

Since $\Gamma_0^*Y = \Gamma_0^*\Xi X$ with probability 1, the unknown mean parameters that may be tested for redundancy are $\Gamma^*\Xi$. We may wish to test the hypothesis
\[
H : \Gamma^*\Xi C = 0 \text{ versus } A : \Gamma^*\Xi C \neq 0, \tag{3.9}
\]
where $C : k \times l$. The restrictions in (3.9) are equivalent to [see (3.5)]
\[
\Theta C = 0
\]
which may be written as
\[
\Theta = \Theta_1 C_0',
\]
where $\Theta_1$ is a new parameter matrix. Therefore, instead of (3.6) we have the model
\[
\Gamma^*Y = \Theta_1 C_0'X + \Gamma^*\Sigma \frac{1}{2} E.
\]
Thus, the next theorem is a consequence of Theorem 3.1.

Theorem 3.2. Let
\[
S_H = Y(I - X'C_0(C_0'XX'C_0)^{-}C_0'X)Y',
\] $H_H : p \times r$ be a semiorthogonal matrix of eigenvectors which correspond to the $r = r(\Sigma)$ largest eigenvalues $l_{Hi}$ of $S_H$, i.e. $H_H' H_H = I_r$, and $(H_H)_{ii} =$
(l_{Hr}). Assume that \( n - r(\mathbf{X} : \mathcal{C}) + r(\mathcal{C}) \geq r \). The maximum likelihood estimators of the parameters in (3.1) under \( H \) in (3.9) are given by

\[
\hat{\Lambda} = \frac{1}{n} \mathbf{L}_H, \\
\hat{\Gamma} = \mathbf{H}_H, \\
\hat{\Sigma} = \frac{1}{n} \mathbf{H}_H \mathbf{L}_H \mathbf{H}_H', \\
\hat{\Xi} = \mathbf{Y} \mathbf{X}' \mathbf{C}_0 (\mathbf{C}_0' \mathbf{X} \mathbf{X}' \mathbf{C}_0)' + \mathbf{U} (\mathbf{I} - (\mathbf{C}_0' \mathbf{X}) (\mathbf{C}_0' \mathbf{X})'),
\]

where \( \mathbf{U} \) is an arbitrary matrix.

Observe that the test in (3.9) is in fact a between individuals test, e.g. a comparison of different treatment groups. The \( \mathbf{\Gamma} \) matrix is only used for reducing the dimension and this reduction is the same for each treatment group. If we are curious about what kind of linear combinations are tested, Corollary 3.1L may be utilized and with probability 1 we are studying \( \mathbf{H}' \Xi \mathbf{C} = 0 \).

We end this section by presenting the likelihood ratio test for \( H \) in (3.9). Let \( U_{p,m,n} \) be the standard test statistic in MANOVA, i.e.

\[
U_{p,m,n} = \frac{|\text{SSE}|}{|\text{SSE} + \text{SST}|},
\]

where \( \text{SSE} \sim W_p(\Sigma, n) \) is the sums-of-squares matrix due to error and \( \text{SST} \sim W_p(\Sigma, m) \) is the sums-of-squares matrix due to hypothesis (see e.g. Srivastava & Carter 1983, p. 97).

**Theorem 3.3.** Suppose \( \mathcal{C}(\mathcal{C}) \subset \mathcal{C}(\mathbf{X}) \). The likelihood ratio test rejects \( H \), given by (3.9), if

\[
\lambda = U_{r(\Sigma), r(\mathcal{C}), n - r(\mathbf{X}) \leq c},
\]

where

\[
\lambda = \frac{|\hat{\Gamma}' \hat{\Sigma} \hat{\Gamma}|}{|\hat{\Gamma}_H' \hat{\Sigma}_H \hat{\Gamma}_H|} = \frac{\prod_{i=1}^{l} l_i}{\prod_{i=1}^{l} \hat{l}_{H_i}},
\]

when \( l_1 > \cdots > l_r \) and \( l_{H1} > \cdots > l_{Hr} \) are the ordered eigenvalues of \( \mathbf{S} \) and \( \mathbf{S}_H \), respectively, \( \hat{\Gamma} \) and \( \hat{\Sigma} \) are as in Theorem 3.1, and \( \hat{\Gamma}_H \) and \( \hat{\Sigma}_H \) are as in Theorem 3.2. The constant \( c \) is chosen so that a prespecified significance level is obtained.
Proof. Now,
\[
\lambda^{-1} = \frac{[\tilde{\Gamma}_H^* Y(I - X'C_0(C_0'XX'C_0)^{-1}C_0'X)Y^T\tilde{\Gamma}]}{[\tilde{\Gamma}^*\Sigma\tilde{\Gamma}]}
\]
\[
= \frac{\frac{\tilde{\Gamma}_H^* Y(1 - X'XX')X'Y^T\tilde{\Gamma}}{[\tilde{\Gamma}^*\Sigma\tilde{\Gamma}]}}{\frac{\tilde{\Gamma}_H^* YY'(XX')^{-1}C(C'(XX')^{-1}C)'(XX')^{-1}YY'T}{[\tilde{\Gamma}^*\Sigma\tilde{\Gamma}]}}
\]
(3.10)
because \( C(C) \subseteq C(X) \). However, since for some orthogonal matrix \( Q \), \( \tilde{\Gamma} = \tilde{\Gamma}_HQ \),
\[
[\tilde{\Gamma}_H^*\Sigma\tilde{\Gamma}_H] = [\tilde{\Gamma}^*\Sigma\tilde{\Gamma}].
\]
Moreover, \( C(\tilde{\Gamma}_H) = C(\Gamma') \) implies that
\[
\frac{\tilde{\Gamma}_H^* (\tilde{\Gamma}_H^*\Sigma\tilde{\Gamma}_H)^{-1}\tilde{\Gamma}_H^*}{\Gamma'(\Gamma'S\Gamma)^{-1}\Gamma'} = \Gamma'(\Gamma'S\Gamma)^{-1}\Gamma'.
\]
Thus, from (3.10), with probability 1,
\[
\lambda^{-1} = \frac{[\Gamma'S\Gamma + \Gamma'YY'(XX')^{-1}C(C'(XX')^{-1}C)'(XX')^{-1}YY'T]}{[\Gamma'S\Gamma]}.
\]
Since
\[
\Lambda^{-\frac{1}{2}}\Gamma'S\Gamma\Lambda^{-\frac{1}{2}} \sim W_{r(\Sigma)}(I,n-r(X))
\]
and under \( H \)
\[
\Lambda^{-\frac{1}{2}}\Gamma'YY'(XX')^{-1}C \sim N_{r(\Sigma),r(C)}(0,I,C'(XX')^{-1}C),
\]
and both expressions are independent of \( \Gamma \) as well as \( \Lambda \), the theorem is established. Here \( N_{p,q}(\mu,\Sigma,\Psi) \) denotes the matrix normal distribution, i.e. \( X \sim N_{p,q}(\mu,\Sigma,\Psi) \) is equivalent to \( \text{vec}X \sim N_{pq}(\text{vec}\mu,\Psi \otimes \Sigma) \).

Remark. If we do not require \( C(C) \subseteq C(X) \), it follows with probability 1 that
\[
\lambda = \frac{[\tilde{\Gamma}_H^*\Sigma\tilde{\Gamma}]}{[\tilde{\Gamma}_H^*\Sigma\tilde{\Gamma}_H\tilde{\Gamma}_H]} \sim U_{r(\Sigma),r(N),n-r(X)},
\]
where \( N \) is a matrix satisfying \( C(N) = C(X) \cap C(C) \).

4. Confidence intervals for \( \Xi \)

In this section we will construct four different types of confidence intervals: a single confidence interval for an arbitrary bilinear combination of the elements in \( \Xi \), i.e. \( a'b \) for any given \( a \) and \( b \); simultaneous within subjects confidence intervals, \( a'b \), i.e. for all \( a \) and any given \( b \); simultaneous between subjects confidence intervals, \( a'b \), i.e. for a given \( a \) and all choices of \( b \); simultaneous confidence intervals, \( a'b \), i.e. for all choices of \( a \) and \( b \).
Throughout, $a \in C(\Sigma)$ and $b \in C(X)$, because we are dealing with matrices which are not supposed to be of full rank.

For the first case we start with the ratio

$$R = \frac{(a'(YX'(XX')^r - \Xi)b)^2}{a'Sab'(XX')^r - b}.$$

The ratio is meaningful since $a \in C(\Sigma)$ and $b \in C(X')$. Then

$$\frac{a'Sa}{a'_\Sigma a} \sim \chi^2(n - r(X))$$

and

$$\frac{(a'(YX'(XX')^r - \Xi)b)^2}{a'Sab'(XX')^r - b} \sim \chi^2(1)$$

which are mutually independent random variables. Thus,

$$(n - r(X))R \sim F(1, n - r(X)).$$

In the next theorem we present a confidence interval based on these quantities.

**Theorem 4.1.** Let $a \in C(\Sigma)$ and $b \in C(X)$. For the MANOVA model (3.1),

$$P(a'\Xi b \in a'YX'(XX')^r - b$$

$$\pm \left\{ \frac{1}{n - r(X)} 1_{P} = 1 - \alpha \right\} \right.$$

To obtain simultaneous within subjects confidence intervals we start again with $R$ but now we rely on the Cauchy-Schwarz inequality because we are interested in

$$\max_{a \in C(\Sigma)} R.$$

Since $\Sigma = \Gamma\Delta\Gamma'$, we study

$$\max_{a \in C(\Sigma)} R = \max_{q} \frac{(q'\Gamma'(YX'(XX')^r - \Xi)b)^2}{q'\Gamma'\Sigma\Gamma q'(XX')^r - b}$$

$$= \frac{b'(YX'(XX')^r - \Xi)b}{b'(XX')^r - b} \sim W_r(\Lambda, n - r(X))$$

and

$$\Gamma'(YX'(XX')^r - \Xi)b(b'(XX')^r - b)^{-1/2} \sim N_r(0, \Lambda).$$

Therefore the next theorem has been established.
Theorem 4.2. Let \( b \in C(X) \) and all \( a \in C(\Sigma) \). Simultaneous within individuals confidence intervals for the MANOVA model (3.1) are given by

\[
P(a'\Xi b \in a'YX'(XX')^{-}b) \\
\pm \left\{ \frac{1}{n-r(\Sigma)-r+1} F_{1-\alpha}(r, n-r(X) - r + 1) a'Sa b'(XX')^{-}b \right\}^{\frac{1}{2}} = 1 - \alpha.
\]

For the simultaneous confidence intervals between the individuals it is noted that

\[
\max_{b \in C(X)} R = \frac{a'(YX'(XX')^{-} - \Xi)(YX'(XX')^{-} - \Xi)'a}{a'Sa}
\]

which is \( F \)-distributed.

Theorem 4.3. Let \( a \in C(\Sigma) \) and all \( b \in C(X) \). For the MANOVA model (3.1), simultaneous between individuals confidence intervals are given by

\[
P(a'\Xi b \in a'YX'(XX')^{-}b) \\
\pm \left\{ \frac{r(X)}{n-r(X)} F_{1-\alpha}(r(X), n-r(X)) a'Sa b'(XX')^{-}b \right\}^{\frac{1}{2}} = 1 - \alpha.
\]

Finally for the simultaneous confidence intervals, i.e. when both \( a \) and \( b \) may vary, we note that

\[
\max_{a \in C(\Sigma), b \in C(X)} R \\
\frac{b'(YX'(XX')^{-} - \Xi)T(G'SG)^{-1}%(YX'(XX')^{-} - \Xi)b}{b'(XX')^{-}b} \equiv e_1,
\]

where \( e_1 \) is the largest eigenvalue of \( BW^{-1} \), where

\[
B = \Gamma'(YX'(XX')^{-} - \Xi)X'(XX')^{-}X(YX'(XX')^{-} - \Xi)'T \sim W_r(\Lambda, r(X)).
\]

The distribution of the largest root of \( W^{-1}B \) has been derived by Pillai (1955) and Roy (1957) and tables where given by Heck (1960). It can also be obtained from the relationship \( e_1 = \frac{e_1}{c_n} \), where \( c_n \) is tabulated by Srivastava (2002, Table B.7).

Theorem 4.4. Let \( c_\alpha \) be the \( \alpha \)-quantile of the distribution of \( e_1 \) in (4.1). Simultaneous confidence intervals for the MANOVA model (3.1) are given by

\[
P(a'\Xi b \in a'YX'(XX')^{-}b \pm \{c_{1-\alpha}/(1 - c_{1-\alpha}) a'Sa b'(XX')^{-}b \}^{\frac{1}{2}}) = 1 - \alpha.
\]
5. Outliers

The problem of finding outliers has received much attention over the years. For a brief review see Srivastava & von Rosen (1998). Consider (3.1) which can be rewritten as

\[ y_i = \Xi x_i + \Sigma^{1/2} \epsilon_i, \quad i = 1, 2, \ldots, n, \]

where \( \epsilon_i \sim N(0, \Sigma) \). A shift in the mean of the i\textsuperscript{th} observation vector may be expressed as

\[ E[y_i] = \Xi x_i + \delta. \tag{5.1} \]

We define an observation \( y_i \) to be an outlier if the hypothesis \( H_0 : \delta = 0 \) is rejected according to a fixed significance level. However, since we will test each observation, many tests are performed and the problem is to find a proper critical value. Let \( e_i \) be the i\textsuperscript{th} column of \( I_n \). Then from (5.1) it follows that

\[ E[Y] = (\delta, \Xi) \begin{pmatrix} e_i' \\ X_i' \end{pmatrix} = \Xi X_i'. \tag{5.2} \]

If \( \delta = 0 \) then we say that we have model \( H \). Otherwise the model in (5.2) is denoted \( H_i \). The likelihood ratio criterion for testing \( H \) against \( H_i \) is according to Theorem 3.3, in obvious notations, given by

\[ \lambda_i = \frac{|\hat{\Gamma}_{H_i}' S_i \hat{\Gamma}_{H_i}|}{|\hat{\Gamma}_{H}' S \hat{\Gamma}_{H}|}, \]

where

\[ S = Y(I - R)Y', \]
\[ S_i = Y(I - R_i)Y', \]
\[ R = X'(XX')^{-1}X, \quad R_i = X_i'(X_i'X_i')^{-1}X_i'. \]

Now, with probability 1

\[ \lambda_i = \frac{|\Gamma'S\Gamma|}{|\Gamma'S_i\Gamma|}. \]

Observe that

\[ S_i = S - (1 - r_{ii})^{-1}d_id'_i, \]

where

\[ d_i = y_i - YX'(XX')^{-1}x_i, \]
\[ r_{ii} = x_i'(XX')^{-1}x_i. \]

Thus we may consider

\[ T_i = d_i'(\Gamma'(S\Gamma)^{-1}\Gamma')d_i/(1 - r_{ii}), \]

since

\[ \lambda_i = (1 - T_i)^{-1}. \]
If we want to have a numerical value of $T_i$ we can replace $\Gamma$ by $\widehat{\Gamma}_H$, for example. This will not change the distribution of $T_i$. Observe that $H$ is rejected for large values of $T_i$. An overall test statistic will be based on

$$ T = \max_{1 \leq i \leq n} T_i. $$

A conservative test based on a Bonferroni upper bound is given by

$$ P(T \geq t) \leq \sum_{i=1}^{n} P(T_i \geq t) $$

and

$$ \frac{n + r(X) - r(\Sigma)}{r(\Sigma)} \frac{T_i}{1 - T_i} \sim F_{r(\Sigma), n-r(X), r(\Sigma)}. $$

Thus,

$$ P(T \geq t) \leq P(F_{r(\Sigma), n-r(X), r(\Sigma)} \geq \frac{n + r(X) - r(\Sigma)}{r(\Sigma)} \frac{t}{1 - t}). $$

To sharpen the above result we may use a generalized Bonferroni inequality (Hunter 1976, Worsley 1982, Meng & von Rosen 2003):

$$ P(T \geq t) \leq n P(T_1 \geq t) - \sum_{i=1}^{n-1} P(T_i \geq t, T_{i+1} \geq t). $$

Hence, the joint distribution of $T_i$ and $T_{i+1}$ is needed. Let

$$ l_i = e_i(I - R)(1 - r_{ii})^{1/2}, $$

$$ L' = (l_i, I_j), \quad F = LL' $$

and then

$$ \Gamma'\Sigma\Gamma = \Gamma'UF^{-1}UT + \Gamma'WT, $$

where

$$ (\Gamma'\Sigma\Gamma)^{-1/2} \Gamma'U = (\Gamma'\Sigma\Gamma)^{-1/2} \Gamma'YL' \sim N_{r(\Sigma),2}(0, I, F), $$

$$ (\Gamma'\Sigma\Gamma)^{-1/2} \Gamma'WT(\Gamma'\Sigma\Gamma)^{-1/2} \sim W_{r(\Sigma)}(I, n - r(X) - 2). $$

Now

$$ T_i = u_i'\Gamma'(\Gamma'UF^{-1}UT + \Gamma'WT)^{-1}u_i, \quad (5.3) $$

where

$$ U = (u_i, u_j) = (Yl_i, Yl_j) $$

and

$$ T_{ij} = u_i'\Gamma'(\Gamma'UF^{-1}UT + \Gamma'WT)^{-1}u_j, $$

$$ \rho_{ij} = l_i'l_j. $$

In (5.3) $T_i$ is beta distributed (see Srivastava & Khatri, 1979). By applying Theorem 2.1 in Srivastava & von Rosen (1998) we get the joint density of $T_i$ and $T_j$. 
Theorem 5.1. Let the statistics $T_i, i = 1, 2, \ldots, n$, be defined by (5.3). Then the joint density of $T_i$ and $T_j, i \neq j$, is given by

$$
(1 - \rho_{ij})^{-\frac{1}{2}}(n - r(X)-3) \frac{\Gamma_r(\Sigma)(\frac{1}{2}(n - r(X)))}{\Gamma_r(\Sigma)(\frac{1}{2}(n - r(X) - 2))}\frac{1}{\Gamma_2(\frac{1}{2}r(\Sigma))}
$$

$$
\times \int_a^b (T_i T_j - T_{ij}) \frac{r(\Sigma)-3}{2} \{(1 - T_i)(1 - t_j) - (\rho_{ij} - T_{ij})^2\}^{\frac{1}{2}(n-k-r(\Sigma)-3)} dT_{ij},
$$

where

$$
a = \max(-\sqrt{T_i T_j}, \rho_{ij} - \sqrt{(1 - T_i)(1 - t_j)}),
$$

$$
b = \min(\sqrt{T_i T_j}, \rho_{ij} + \sqrt{(1 - T_i)(1 - t_j)}).
$$

References


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