SELECTION PROCEDURES FOR A PROBLEM IN ANALYSIS OF VARIANCE. (U)

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Selection Procedures for A Problem in Analysis of Variance*

by

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1. Introduction

For a completely randomized block design with one observation per cell, we express the observable random variables $X_{i\alpha}$ \((i = 1, \ldots, k; \alpha = 1, \ldots, n)\) as

\[(1.1) \quad X_{i\alpha} = \mu + \beta_{i} + \tau_{i} + \xi_{i\alpha}, \quad \sum_{i=1}^{k} \tau_{i} = 0,\]

where $\mu$ is the mean-effect, $\beta_{1}, \ldots, \beta_{n}$ are the block effects (nuisance parameters for the fixed effects model), $\tau_{1}, \ldots, \tau_{k}$ are the treatment effects, and $\xi_{i\alpha}$ are the error components. We assume that the errors within each block are jointly normally distributed.

We assume that the quality of a treatment is judged by the largeness of the $\tau_{i}$'s. A 'population' $\pi_{i}$ is called the best if $\tau_{i}$ is the largest. In general, it may be complicated to derive suitable tests for appropriate hypotheses, in which the experimenter may really be interested. We apply the subset selection approach (using certain basic hypotheses) and thus obtain more appropriate information regarding the treatments. A subset selection procedure is designed to select a subset so as to include the best population. Selection of any subset that contains the best is called a correct selection (CS).

Roughly speaking, any two populations that are in the same selected subset, will be considered as "equivalently good". If all populations are selected, we claim that all treatments are homogeneous. In general, for achieving the objective of the experimenter, one should establish a suitable set of basic hypotheses. Depending on the objective one should proceed to consider different

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ways of formulating the basic hypotheses. In this paper, we discuss a
method based on subset selection rules for the purpose of making a claim
of the type: \( \tau_i = \tau^* > \tau_j + \Delta \) for all \( i \in I \) and \( j \in J \), where \( I \) and \( J \) form
a partition of \( \{1,2,\ldots,k\} \). The process of making such a claim will be
called hypothesis identification. This is achieved by setting up certain
basic hypotheses regarding the \( \tau_i \)'s and using a subset selection procedure
to test these basic hypotheses. It should be pointed out that in identifying
an appropriate hypothesis, we assume that the constant \( \Delta \) in the claim is
specified by the experimenter, say, based on past experience. Associated with
the tests of the basic hypotheses using a selection rule, there are error
probabilities and the infimum of the probability of a correct selection for
the rule employed. These are related to the power function of these tests.
The sum of the average (over the basic hypotheses tested) of the error
probabilities and one minus the infimum of the probability of a correct
selection is called the identification risk. The main theorem of the
paper discusses the derivation of an optimal selection rule in the sense of
minimizing the identification risk. For a more general theory of
multiple decisions from ranking and selection approach, one can refer to a
recent monograph by Gupta and Huang (1981). A general survey of the entire
field is provided in Gupta and Panchapakesan (1979).

Let \( Y \) be a random observable vector with probability distribution depending
upon a parameter \( \tau' = (\tau_1,\ldots,\tau_k) \in \mathbb{R} \). Consider a family of hypotheses testing
problems as follows:

\[
(1.2) \quad H_0: \ \tau \in \Omega_0 \ vs \ H_i: \ \tau \in \Omega_i, \ 1 \leq i \leq k, 
\]

where \( \Omega_0 = \{\tau_1 = \ldots = \tau_k\} \) and \( \Omega_i = \{\tau_i > \max_{j \neq i} \tau_j\}, \ i = 1,2,\ldots,k \). A test
of the hypotheses (1.2) will be defined to be a vector \( (\delta_1(\gamma),\ldots,\delta_k(\gamma)) \),
where the elements of the vector are ordinary test functions; when \( \gamma \) is observed
we reject $H_0$ in favor of $H_i$ with probability $\delta_i(y)$, $1 \leq i \leq k$. The power function of a test $(\delta_1, \ldots, \delta_k)$ is defined to be the vector $(\beta_1(\bar{y}), \ldots, \beta_k(\bar{y}))$, where $\beta_i(\bar{y}) = E_i \delta_i(\bar{y})$, $1 \leq i \leq k$. For $i \in \Omega_i$, $\beta_i(\bar{y})$ is the probability of a correct selection $P(\text{CS})$ and $\delta_i(y)$ is the individual selection probability of selecting the best population $\pi_i$. Let $S_y$ be the set of all the tests $(\delta_1, \ldots, \delta_k)$ such that

\[(1.3) \quad E_i \delta_i(\bar{y}) \leq y, \quad i \in \Omega_0, \quad 1 \leq i \leq k, \]

where $y$ is the upper bound on the error probabilities associated with the treatment effects.

For each $i$, $(1 \leq i \leq k)$, we would like to have $\beta_i(\bar{y})$ large when $\bar{y} \in \Omega_i$ subject to (1.3). For $i \in \Omega_i$, if we make $\beta_i(\bar{y})$ large, then $\beta_j(\bar{y})$ should be small for $j \neq i$.

It should be pointed out that in the formulation and proof of the optimal selection procedure, results from Neyman-Pearson theory are used.

2. Formulation of an Optimal Selection Procedure

Assume that

$$x'_\alpha = (x_{1\alpha}, \ldots, x_{k\alpha}),$$

$\alpha = 1, \ldots, n$, are independently and identically distributed random vectors with the following distribution:

\[(2.1) \quad (2\pi\sigma^2)^{-\frac{1}{2}kn}|\Lambda|^{-\frac{1}{2}}\exp\left[-\frac{1}{2\sigma^2}(x - \theta)'\Lambda^{-1}(x - \theta)\right],\]

where $x' = (x_1, \ldots, x_k); \ldots; x_{1n}, \ldots, x_{kn}$ and $\theta' = (\theta_{11}, \ldots, \theta_{1k}; \ldots; \theta_{kn}, \ldots, \theta_{kn})$, $\theta_{i\alpha} = \mu + \theta_{i\alpha} + \tau_i$, $i = 1, \ldots, k$; $\alpha = 1, 2, \ldots, n$ and $\Lambda$ is a known positive definite $kn \times kn$ correlation matrix defined as follows:
\[ \Lambda = (\lambda_{ij})_{kn \times kn} \]

\[
\begin{bmatrix}
\Lambda_1 & 0 & \ldots & 0 \\
0 & \Lambda_1 & \ldots & 0 \\
\vdots & \vdots & \ddots & \vdots \\
0 & 0 & \ldots & \Lambda_1
\end{bmatrix}, \quad \text{where}
\]

\[ \Lambda_1 = \begin{bmatrix} \lambda & \cdot & \cdot \\ \cdot & \lambda & \cdot \\ \cdot & \cdot & \cdot \end{bmatrix}_{k \times k}. \]

We rewrite the original model as the general linear model as follows:

\[
\bar{X} = \theta + \xi, \quad \xi \sim N(0, \sigma^2\Lambda).
\]

Since we are interested in the difference between all pairs of \( \tau_i \)'s, we transform the linear model to the following: For any \( i \), let

\[
Y_i = C \tau_i + \eta, \quad \eta \sim N(0, \sigma^2\Sigma_i),
\]

where \( \tau_i = (\tau_{i1}, \ldots, \tau_{ik}) \), \( \tau_{ij} = \tau_i - \tau_j, j \neq i \),

\[
Y_i = (Y_{i11}, \ldots, Y_{i1k}; Y_{i21}, \ldots, Y_{i2k})_{1 \times (k-1)n}
\]

\[
Y_{ij} = X_{ij} - X_{ji}, i \neq j; i, j = 1, \ldots, k; \ell = 1, \ldots, n,
\]

\[
Y_i = A_i \bar{X}, \quad \eta = A_i \xi
\]

\[
A_i = \begin{bmatrix}
A_{i1} \\
& A_{i1} & 0 \\
& & \ddots & \ddots \\
& & & 0 & \ddots \\
& & & & \ddots & A_{i1}
\end{bmatrix}_{(k-1) \times kn}.
\]
\[ \Sigma_i = A_i \Lambda_i A_i^t = \begin{bmatrix} A_i^t \Lambda_i A_i^t & 0 \\ 0 & \cdots & 0 \\ A_i^t \Lambda_i A_i^t & \cdots & 0 \end{bmatrix} (k-1)nx(k-1)n, \]

\[ A_i = \begin{bmatrix} -1 & 0 \cdots & 0 & 1 & 0 \cdots & 0 \\ 0 & -1 & 0 \cdots & 0 & 1 & 0 \cdots & 0 \\ \vdots & \vdots & \ddots & \vdots & \vdots & \ddots & \vdots \\ 0 \cdots & 0 & 1 & -1 & 0 \cdots & 0 \\ 0 \cdots & 0 & 1 & -1 & 0 \cdots & 0 \\ 0 \cdots & 0 & 1 & 0 \cdots & 0 & -1 \end{bmatrix}^{i-1} \begin{bmatrix} 0 \cdots & 0 & 1 & 0 \cdots & 0 \\ 0 \cdots & 0 & 1 & -1 \end{bmatrix}^{i+1} (k-1)x(k-1) \]

\[ C_i = [I, \cdots, I]_{(k-1)x(k-1)n} \]

where each of the identity matrix in \( C_i \) is \((k-1)x(k-1)\). The maximum likelihood estimator of \( \Sigma_i \) is as follows:

\[ \hat{\Sigma}_i = (C_i \Sigma_i^{-1} C_i)^{-1} C_i \Sigma_i^{-1} Y_i. \]

Since,

\[ A_i \Lambda_i A_i^t = (1-\lambda) \begin{bmatrix} 2 & 1 \\ 1 & 2 \end{bmatrix} (k-1)x(k-1) \]

\[ (A_i \Lambda_i A_i^t)^{-1} = (1-\lambda)^{-1} \frac{1}{k} \begin{bmatrix} k-1 & -1 \\ -1 & k-1 \end{bmatrix} = V_i \]

\[ C_i \Sigma_i^{-1} C_i = n (A_i \Lambda_i A_i^t)^{-1} = \frac{n}{k(1-\lambda)} \begin{bmatrix} k-1 & -1 \\ -1 & k-1 \end{bmatrix} \]

\[ (C_i \Sigma_i^{-1} C_i)^{-1} = \frac{1-\lambda}{n} \begin{bmatrix} 2 & 1 \\ 1 & 2 \end{bmatrix} (k-1)x(k-1), \]
\[ C_i^{-1} = [I \ldots I] \begin{bmatrix} V_1 & 0 \\ 0 & V_i \end{bmatrix} = [V_1, \ldots, V_i] \]

\[ (C_i^{-1}C)^{-1}C_i^{-1} = \frac{1 - \lambda}{n} \begin{bmatrix} 2 & 1 \\ 1 & 2 \end{bmatrix} [V_1, \ldots, V_i] = \frac{1}{n} [I, \ldots, I]. \]

Hence,

\[ \bar{x}_i = (C_i^{-1}C)^{-1}C_i^{-1} Y_i \]

\[ \frac{1}{n} \begin{bmatrix} \sum_{k=1}^{n} Y_{ik} \\ \vdots \\ \sum_{k=1}^{n} Y_{ik} \end{bmatrix} = \begin{bmatrix} Y_{i1} \\ \vdots \\ Y_{ik} \end{bmatrix} = \begin{bmatrix} \bar{x}_i - \bar{x}_1 \\ \vdots \\ \bar{x}_i - \bar{x}_k \end{bmatrix}, \]

where \( \bar{x}_i = \frac{1}{n} \sum_{j=1}^{n} X_{ij}, \ 1 \leq i \leq k. \)

The joint density of \( Y_{i1}, \ldots, Y_{ik}; \ldots; Y_{1n}, \ldots, Y_{kn} \) is the following:

\[ p_{\bar{x}_i}(\bar{x}_i) = (2\pi \sigma^2)^{-\frac{k}{2}} \exp\left[-\frac{1}{2\sigma^2} (\bar{x}_i - C_{\bar{x}_i})'C_{\bar{x}_i}^{-1}(\bar{x}_i - C_{\bar{x}_i})\right] \]

where

\[ C_{\bar{x}_i} = A_1 \land A_1' = (1 - \lambda) \begin{bmatrix} J & 0 \\ 0 & J \end{bmatrix}(k-1)nx(k-1)n \]

\[ J = \begin{bmatrix} 2 & 0 \\ 0 & 2 \end{bmatrix} (k-1)x(k-1). \]

\[ \bar{x}_i^{-1} = \begin{bmatrix} V_1 & 0 \\ 0 & V_i \end{bmatrix}. \]
Now, we specify the $\Omega_i$'s as follows (Note that this is a different specification from that given earlier):

$$\Omega_i = \{ \tau_i \mid \tau_i \geq \max_{j \neq i} \tau_j + \Delta \sigma \}, \quad 1 \leq i \leq k,$$

and

$$\tilde{\Omega} = \bigcup_{i=1}^{k} \Omega_i.$$

Assume that $\sigma$ is known. Let

$$\Delta_i = (\Delta \sigma, \ldots, \Delta \sigma)_{1 \times (k-1)}, \quad i = 1, \ldots, k, \quad \Delta > 0.$$

Thus

$$\frac{p_{\Delta_i}(y_i)}{p_{0}(y_i)} = \exp \left( \frac{1}{\sigma^2} \right) \left( \frac{y_i - (C_{\Delta_i})' \varepsilon_i^{-1} (x_{\Delta_i} - C_{\Delta_i}) + \varepsilon_i^{-1} y_i}{2\sigma^2} \right) \Delta_i C' \varepsilon_i^{-1} C_{\Delta_i}.$$}

Hence, we can rewrite

$$\frac{p_{\Delta_i}(y_i)}{p_{0}(y_i)} \geq d' \quad \text{as} \quad y_{i1} + \ldots + y_{ik} \geq d'' \sigma.$$

Let a selection rule $\delta^0 = (\delta^0_1, \ldots, \delta^0_k)$ be defined by

$$\delta^0_i(y_i) = \begin{cases} 1 & \text{if} \quad p_{\Delta_i}(y_i) \geq d' p_{0}(y_i), \\ 0 & \text{otherwise} \end{cases},$$

such that

$$(2.2) \quad \sum_{i=1}^{k} \delta^0_i(y_i) = \gamma, \quad i \in \Omega_0. \quad \text{Then}$$

$\delta^0$ maximizes
Note that $\delta_i^0(Y_i)$ is also based on the maximum likelihood estimators $\tilde{\tau}_i$ of $\tau_i$. Since for any $\delta \in S(y)$,

$$P(CS|\delta) = \int \delta_i(Y_i)p_\tau(Y_i) d\nu(Y_i)$$

$$\geq \min_{1 \leq i < k} \inf_{\tau \in \Omega_i} \int \delta_i(Y_i)p_\tau(Y_i) d\nu(Y_i).$$

We have

$$\inf_{\tau \in \tilde{\Omega}} P(CS|\delta) = \min_{1 \leq i < k} \inf_{\tau \in \Omega_i} \int \delta_i(Y_i)p_\tau(Y_i) d\nu(Y_i).$$

For any $\delta \in S(y)$, it follows that

$$\int (\delta_i - \delta_i^0)(p_{\tilde{\Delta}_i} - dp_0) \leq 0$$

which implies

$$\int \delta_i^0 p_{\tilde{\Delta}_i} \geq \int \delta_i p_{\tilde{\Delta}_i}.$$

Since $\delta_i^0(Y_i)$ is nondecreasing in $Y_i$, hence

$$\inf_{\tau \in \tilde{\Omega}} P(CS|\delta^0) = \min_{1 \leq i < k} \inf_{\tau \in \Omega_i} \int \delta_i^0(Y_i)p_{\tilde{\Delta}_i}(Y_i) d\nu(Y_i)$$

$$\geq \min_{1 \leq i < k} \int \delta_i(Y_i)p_{\tilde{\Delta}_i}(Y_i) d\nu(Y_i)$$

$$\geq \min_{1 \leq i < k} \inf_{\tau \in \Omega_i} \int \delta_i(Y_i)p_{\tau}(Y_i) d\nu(Y_i)$$

$$= \inf_{\tau \in \tilde{\Omega}} P(CS|\delta).$$

We rewrite $\delta^0$ as follows:

$$\delta_i^0(Y_i) = \begin{cases} 1 & \text{if } Y_{i1} + \ldots + Y_{ik} \geq d\nu_0, \\ 0 & \text{otherwise} \end{cases}$$
Thus, the optimal subset selection rule is as follows:

\[ \delta^0_i(x) = \begin{cases} 1 & \text{if } \bar{x}_i \geq \frac{1}{k-1} \sum_{j \neq i} \bar{x}_j + d_0, \\ 0 & \text{otherwise} \end{cases} \]

where \( d = \frac{d''}{k-1} \).

Now, we wish to determine \( d \) and \( n \). We make the following transformation:

\[ z_{ik} = \left[ 1 \ldots 1 \right]_l x(k-1) = \begin{bmatrix} y_{i1} \\ \vdots \\ y_{ik} \end{bmatrix}, \quad \text{and} \]

\[ \tau = \tau_{i1} + \ldots + \tau_{ik} = (k-1)\tau_i - \sum_{j \neq i} \tau_j. \]

Since the distribution of

\[ \tilde{z}_i = \begin{bmatrix} y_{i1} \\ \vdots \\ y_{ik} \end{bmatrix} = (C'\Sigma_i^{-1}C)^{-1}C'\Sigma_i^{-1}y_i \]

is \( (2\pi)^{-\frac{k}{2}}|\Sigma_i|^{-\frac{1}{2}} \exp\left[-\frac{1}{2} (\tilde{z}_i - \bar{z}_i)'\Sigma_i^{-1}(\tilde{z}_i - \bar{z}_i)\right] \), where \( \bar{z}_i = \frac{1-\lambda}{n}J \).

Then the distribution of \( Z_{ik} \) is

\[ [2\pi \sigma^2(1-\lambda)k(k-1)]^{-\frac{1}{2}} \exp\left[-\frac{n}{2\sigma^2(1-\lambda)k(k-1)} (Z_{ik} - \tau)^2 \right]. \]

Hence

\[ E_\sigma \delta^0_{i}(y_i) = P(Z_{ik} \geq d''\sigma) \]

\[ = \phi\left[-\frac{d''n}{\sqrt{(1-\lambda)k(k-1)}}\right] = \gamma, \]

and
\[
\inf_{\delta \in \mathcal{D}} \mathbb{E}[\text{CS} | \delta^0] = \min_{1 \leq i < k} \int_{1 \leq i < k} \mathbb{P}_i (Z_{ik} > d^\nu) \mathbb{P}(Z_{ik} - (k-1)\Delta) \mathbb{P}(Z_{ik}) \mathbb{P}(Z_{ik})^{(k-1)\Delta}
\]

\[
= \min_{1 \leq i < k} \mathbb{P}_i \left( \frac{(Z_{ik} - (k-1)\Delta)^\nu}{\sqrt{(1-\lambda)k(k-1)}} \geq \frac{(d^\nu - (k-1)\Delta)^\nu}{\sqrt{(1-\lambda)k(k-1)}} \right)
\]

\[
(2.5) \quad \phi \left[ - \frac{(d^\nu - (k-1)\Delta)^\nu}{\sqrt{(1-\lambda)k(k-1)}} \right] = p^*.
\]

For given \( r, p^*, k, \lambda, \) and \( \Delta, \) we can find \( d^\nu \) and the smallest number of blocks, \( n, \) to satisfy equations (2.4) and (2.5). Note that this \( n \) is also the minimum sample size for the case of one observation per cell in the completely randomized block design.

We rewrite (2.4) and (2.5) as

\[
\phi \left[ - \frac{d\sqrt{n(k-1)}}{\sqrt{(1-\lambda)k}} \right] = \gamma
\]

and

\[
\phi \left[ - \frac{(d-\Delta)^\nu \sqrt{n(k-1)}}{\sqrt{(1-\lambda)k}} \right] = p^*.
\]

Let \( z_{p^*} \) and \( z_{\gamma} \) represent the upper percentage points corresponding to \( p^* \) and \( \gamma, \) respectively of the standard normal distribution. Then we have

\[
d = -\frac{z_{\gamma}}{z_{p^*} - z_{\gamma}},
\]

and

\[
n = \left\lfloor \frac{(1-\lambda)k(z_{p^*} - z_{\gamma})^2}{(k-1)\Delta^2} \right\rfloor,
\]

where \( \lfloor a \rfloor \) is the smallest integer greater than or equal to \( a. \)
Summarizing the previous results, we obtain the following theorem.

**Theorem:** Under model (1.1) with the stated assumption on $\xi_\alpha$, an optimal procedure for selecting a subset of the "best" or "worthwhile" treatments based on the observed data $x$ and satisfying the conditions (2.2) and (2.3) is: Select the population $\tau_i$ with probability $\delta_i^0(x)$ given by

$$
\delta_i^0(x) = \begin{cases} 
1 & \text{if } \bar{x}_i \geq \frac{1}{k-1} \sum_{j \neq i} \bar{x}_j + \sigma, \\
0 & \text{otherwise,}
\end{cases}
$$

where the smallest values of $d$ and $n$ are given by

$$
d = -\frac{z_{1-\gamma}}{z_{p*} - z_{1-\gamma}},
$$

and

$$
n = \frac{(1-\lambda)k(z_{p*} - z_{1-\gamma})^2}{(k-1)d^2}.
$$

Furthermore, we have established the following connection between the selection procedure and the hypothesis identification problem as follows:

If $\tau_{i_1}, \tau_{i_2}, \ldots, \tau_{i_j}$ ($j \leq k$) are selected, we say that these populations are homogeneous and make the hypothesis identification $H_i^j$: $\tau_{i_1} = \ldots = \tau_{i_j} \geq \max_{1 \leq l \leq k} \tau_l + \Delta \sigma.

Note that the overall identification risk connected with this problem is $\leq \gamma + (1 - P^*)$.

**Remark:** It should be pointed out that for some pairs $(\gamma, P^*)$, $\delta^0$ may not select any population. This is to be interpreted as not identifying any one of the appropriate hypotheses.
We consider some special cases to provide an idea as to the appropriate identification of one of the hypotheses. For $\gamma = 0.05, \lambda = 0.5$ and $P^* = 0.95, 0.90, 0.80$; then

(i) $k = 2$,

$H_0: \tau_1 = \tau_2, \quad H_1^1: \tau_1 \geq \tau_2 + \Delta \sigma, \quad H_2^1: \tau_2 \geq \tau_1 + \Delta \sigma.$

In this case, for specified $\Delta$-values, the smallest $d$ and $n$ needed for the optimal selection rule are given in the following table.

<table>
<thead>
<tr>
<th>$\Delta$</th>
<th>0.1</th>
<th>0.5</th>
<th>1.0</th>
<th>2.0</th>
</tr>
</thead>
<tbody>
<tr>
<td>d(0.95,0.90,0.80)</td>
<td>0.05,0.06,0.07</td>
<td>0.25,0.32,0.33</td>
<td>0.50,0.64,0.66</td>
<td>1.00,1.29,1.33</td>
</tr>
<tr>
<td>n(0.95,0.90,0.80)</td>
<td>1089,858,620</td>
<td>44,35,25</td>
<td>11,9,7</td>
<td>3,3,2</td>
</tr>
</tbody>
</table>

(ii) $k = 3$,

$H_0: \tau_1 = \tau_2 = \tau_3, \quad H_1^1: \tau_1 \geq \max(\tau_2, \tau_3) + \Delta \sigma,$

$H_2^1: \tau_2 \geq \max(\tau_1, \tau_3) + \Delta \sigma, \quad H_3^1: \tau_3 \geq \max(\tau_1, \tau_2) + \Delta \sigma,$

$H_4^1: \tau_1 = \tau_2 \geq \tau_3 + \Delta \sigma, \quad H_5^1: \tau_1 = \tau_3 \geq \tau_2 + \Delta \sigma,$

$H_6^1: \tau_2 = \tau_3 \geq \tau_1 + \Delta \sigma.$

For optimal selection rule, the minimum value of $d$ and $n$ are computed (for specified values of $\Delta$) and given in the following table.

<table>
<thead>
<tr>
<th>$\Delta$</th>
<th>0.1</th>
<th>0.5</th>
<th>1.0</th>
<th>2.0</th>
</tr>
</thead>
<tbody>
<tr>
<td>d(0.95,0.90,0.80)</td>
<td>0.05,0.06,0.07</td>
<td>0.25,0.32,0.33</td>
<td>0.50,0.64,0.66</td>
<td>1.00,1.29,1.33</td>
</tr>
<tr>
<td>n(0.95,0.90,0.80)</td>
<td>817,644,465</td>
<td>33,26,19</td>
<td>9,7,5</td>
<td>3,2,2</td>
</tr>
</tbody>
</table>

(iii) $k = 4$,

$H_0: \tau_1 = \tau_2 = \tau_3 = \tau_4, \quad H_1^1: \tau_1 \geq \max(\tau_2, \tau_3, \tau_4) + \Delta \sigma,$
\[ H_2^i: \tau_2 \geq \max(\tau_1, \tau_3, \tau_4) + \Delta \sigma, \quad H_3^i: \tau_3 \geq \max(\tau_1, \tau_2, \tau_4) + \Delta \sigma, \]
\[ H_4^i: \tau_4 \geq \max(\tau_1, \tau_2, \tau_3) + \Delta \sigma, \quad H_5^i: \tau_1 = \tau_2 \geq \max(\tau_3, \tau_4) + \Delta \sigma, \]
\[ H_6^i: \tau_1 = \tau_3 \geq \max(\tau_2, \tau_4) + \Delta \sigma, \quad H_7^i: \tau_1 = \tau_4 \geq \max(\tau_2, \tau_3) + \Delta \sigma, \]
\[ H_8^i: \tau_2 = \tau_3 \geq \max(\tau_1, \tau_4) + \Delta \sigma, \quad H_9^i: \tau_2 = \tau_4 \geq \max(\tau_1, \tau_3) + \Delta \sigma, \]
\[ H_{10}^i: \tau_3 = \tau_4 \geq \max(\tau_1, \tau_2) + \Delta \sigma, \quad H_{11}^i: \tau_1 = \tau_2 \geq \tau_3 \geq \tau_4 + \Delta \sigma, \]
\[ H_{12}^i: \tau_1 = \tau_2 = \tau_4 \geq \tau_3 + \Delta \sigma, \quad H_{13}^i: \tau_1 = \tau_3 \geq \tau_4 \geq \tau_1 + \Delta \sigma, \]
\[ H_{14}^i: \tau_2 = \tau_3 = \tau_4 \geq \tau_1 + \Delta \sigma. \]

For the optimal selection rule, the minimum value of \( d \) and \( n \) are computed (for specified values of \( \Delta \)) and given in the following table.

<table>
<thead>
<tr>
<th>( \Delta )</th>
<th>0.1</th>
<th>0.5</th>
<th>1.0</th>
<th>2.0</th>
</tr>
</thead>
<tbody>
<tr>
<td>( d(0.95,0.90,0.80) )</td>
<td>0.05,0.06,0.07</td>
<td>0.25,0.32,0.33</td>
<td>0.50,0.64,0.66</td>
<td>1.00,1.29,1.33</td>
</tr>
<tr>
<td>( n(0.95,0.90,0.80) )</td>
<td>726,572,413</td>
<td>30,23,17</td>
<td>8,6,5</td>
<td>2,2,2</td>
</tr>
</tbody>
</table>

Note that \( P^* \) is the probability of correct selection for the associated subset selection rule, while the error probability \( \gamma \) is controlled at 5 percent level. The identification risk is \( 0.05 + (1-P^*) \). We can explain the cases described above as follows: for \( k = 2 \), if the selected subset contains \( \pi_1 \) only, we identify \( H_{1i}^i, i = 1,2 \); if it contains \( \pi_1 \) and \( \pi_2 \), we identify \( H_0 \). For \( k = 3 \), if the selected subset contains \( \pi_1 \) only, we identify \( H_{1i}^i, i = 1,2,3 \); if it contains \( \pi_1 \) and \( \pi_2 \), \( \pi_1 \) and \( \pi_3 \), or \( \pi_2 \) and \( \pi_3 \) only, we identify \( H_4, H_5 \) or \( H_6 \), respectively. Similar discussion applies to the case \( k = 4 \).
Now, we discuss the case where $\sigma^2$ is unknown. For any $i$, the maximum likelihood estimators of $\tau_i$ and $\sigma^2$ are:

$$\hat{\tau}_i = (C_i^t \Sigma_i^{-1} C_i)^{-1} C_i^t \Sigma_i^{-1} Y_i = \left[ \begin{array}{c} \hat{Y}_{i1} \\ \vdots \\ \hat{Y}_{ik} \end{array} \right]$$

and

$$\hat{\sigma}^2 = \frac{1}{(k-1)(n-1)} \sum_i [\Sigma_i^{-1} - \Sigma_i^{-1} C_i (C_i^t \Sigma_i^{-1} C_i)^{-1} C_i^t \Sigma_i^{-1}] Y_i.$$ 

We know that $\sigma^2$ and $\hat{\tau}_i$ are independent and the distribution $f(s)$ of $s = \frac{\hat{\sigma}^2}{\sigma^2}$ is $\chi^2_p(s)$ with $p = (k-1)(n-1)$.

As before, we define the selection rule as follows:

$$\varphi_i(\hat{\tau}_i, \hat{\sigma}) = \begin{cases} 1 & \text{if } \hat{Y}_{i1} + \ldots + \hat{Y}_{ik} \geq d_1 \hat{\sigma}, \\ 0 & \text{otherwise} \end{cases}$$

or

$$\varphi_i^0(x, \hat{\sigma}) = \begin{cases} 1 & \text{if } \bar{x}_i \geq \frac{1}{k-1} \sum_{j \neq i} \bar{x}_j + \frac{d_1}{k-1} \hat{\sigma}, \\ 0 & \text{otherwise} \end{cases}$$

Conditionally, for an observed value of $\hat{\sigma}$, we can discuss the optimality as before. However, the constant $d$ and $n$ can be determined without any difficulty by (2.8) and (2.9). Since

$$E_{\hat{\tau}_i} \varphi_i(\hat{\tau}_i, \hat{\sigma}) = \gamma, \quad \tau \in \tau_0$$

we get
(2.6) \[ \int \phi \left( - \frac{d_1 s \sqrt{n}}{\sqrt{1-\lambda)k(k-1)}} \right) f(s) ds = \gamma, \]

and

\[ \inf \mathbb{P}(CS|\theta^0) \]

(2.7) \[ = \int \phi \left[ - \frac{(d_1 s - (k-1)\Delta)^{1/2}}{\sqrt{1-\lambda)k(k-1)}} \right] f(s) ds = p^*. \]

This gives

(2.8) \[ t\left[ - \frac{d_1 \sqrt{n(n-1)}}{\sqrt{1-\lambda)k}} \right; (k-1)(n-1), 0] = \gamma, \]

and

(2.9) \[ t\left[ - \frac{d_1 \sqrt{n(n-1)}}{\sqrt{1-\lambda)k}} ; (k-1)(n-1), \frac{\Delta \sqrt{n(n-1)}}{\sqrt{1-\lambda)k}} \right] = p^*, \]

where \( t(a; b, c) \) is the percentage point of the noncentral t with \( b \) degrees of freedom and the noncentrality parameter \( c \).

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References


Selection Procedures for a Problem in Analysis of Variance

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Multiple decision rules, Completely randomized block design, Joint normal distribution, Hypothesis identification, Identification risk, Subset selection, Optimal selection procedures.

There are many situations in the analysis of variance where an experimenter would like to make comparisons among (and select the "best" set) the treatments. In this paper we study the problem where the data are based on a completely randomized block design. It is shown that the subset selection approach is a useful method to make appropriate "identification" among the hypotheses and the selected subset. We propose an optimal selection procedure which controls the error probabilities when all the parameters (treatments) are equal and which maximizes the infimum of the probability of a correct selection over some preference parameter.
space, simultaneously. Some examples are provided to illustrate the optimal subset selection rule and its interpretation in terms of the "identified" hypotheses.