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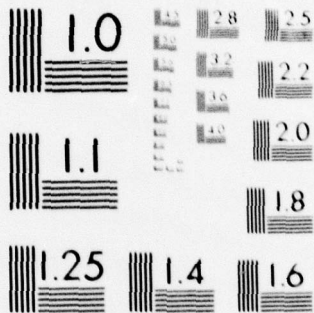
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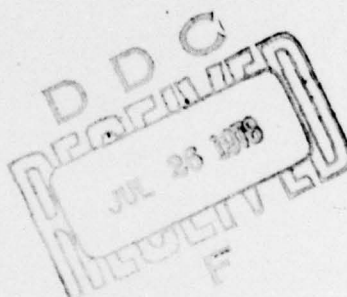
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A Note on Optimal Subset Selection Procedures*

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1. Introduction. A common problem faced by an experimenter is one of comparing several categories or populations. The classical approach to this problem has been to use a test for homogeneity, i.e., to test whether all categories (populations) are identical or not. This approach is not always realistic and it is often inadequate. The inadequacy lies in the fact that only two decisions, accept or reject the hypothesis, are available to the experimenter. The experimenter is then faced with the problem of what to do next, especially if the hypothesis is rejected. These difficulties and inadequacies may be alleviated by formulating the problem as multiple decision problems aimed at selection or ranking (ordering) of the k populations. This has led to the rapid development of selection and ranking theory during the last two decades. Many reasonable rules have been proposed. Some desirable properties of these rules have been studied. However, very little work has been done to study the optimality of selection procedures, especially in the subset selection approach. In this paper, we are interested in deriving some methods to construct optimal subset selection procedures. Some classical selection rules are constructed as special cases.

Let $\pi_1, \pi_2, \dots, \pi_k$ represent $k (\geq 2)$ populations (treatments) and let $x_{i1}, \dots, x_{in_i}$ be n_i independent random observations from π_i . The quality of

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the i th population π_i is characterized by a real-valued parameter θ_i , usually unknown. Let $\Omega = \{\underline{\theta} | \underline{\theta}' = (\theta_1, \dots, \theta_k)\}$ denote the whole parameter space.

Let $\tau_{ij} = \tau_{ij}(\underline{\theta})$ be a measure of separation between π_i and π_j . We assume that there exists a monotonically nonincreasing function h such that $\tau_{ji} = h(\tau_{ij})$.

Let $\Omega_i = \{\underline{\theta} | \tau_{ij} \geq \tau_0, j \neq i\}$, $1 \leq i \leq k$, and $\Omega_0 = \Omega - \bar{\Omega}$, where $\bar{\Omega} = \bigcup_{i=1}^k \Omega_i$. Thus

$\Omega = \bigcup_{i=0}^k \Omega_i$ is a partition of Ω . For this problem, we assume τ_0 and τ_{ii} as known with $\tau_0 > \tau_{ii}$ for all i . Let $\tau_i = \min_{j \neq i} \tau_{ij}$, $1 \leq i \leq k$. We define

$\tau^* = \max_{1 \leq i \leq k} \tau_i$. The population associated with τ^* will be called the best population.

It should be pointed out that if $\underline{\theta} \in \Omega_i$, then $\tau_i \geq \tau_j$ for all j ,

since for some j , $j \neq i$, $\tau_{ji} = h(\tau_{ij}) \leq h(\tau_0) \leq h(\tau_{ii}) = \tau_{ii} < \tau_0$. Thus if

$\underline{\theta} \in \Omega_i$, then π_i is the best population. A selection of a subset containing

the best population is called a correct selection (CS). (Note that in case

of ties any one of the best populations corresponding to τ^* is "tagged" as the best population.) To illustrate the above notation, we assume that the observa-

tions are drawn from π_i which has a normal distribution with unknown mean

θ_i ($i = 1, \dots, k$) and known variance σ^2 . We can define $\tau_{ij} = \theta_i - \theta_j$; then it

can be seen that $\tau_i = \theta_i - \theta_{[k]}$ if $\theta_i < \theta_{[k]}$ and $\tau_i = \theta_i - \theta_{[k-1]}$ if $\theta_i = \theta_{[k]}$,

where $\theta_{[1]} \leq \dots \leq \theta_{[k]}$. In this case, $\tau_{ii} = 0$ for all i . Thus the population

with the largest mean, $\theta_{[k]}$, is the best. If instead $\tau_{ij} = \theta_j - \theta_i$ then the popula-

tion with the smallest mean $\theta_{[1]}$ would be the best. In the above example, we

have $h(t) = -t$ which is a decreasing function.

Let the observed sample vector be denoted by $\underline{X}' = (X'_1, \dots, X'_k)$ where $\underline{X}_i$ has components $X_{i1}, \dots, X_{in_i}$, $i = 1, \dots, k$. Let $\delta = (\delta_1, \dots, \delta_k)$ be a selection procedure where $\delta_i(\underline{x})$ is the probability of selecting π_i ($1 \leq i \leq k$) based on the observed vector $\underline{X} = \underline{x}$ from the k populations. As measures of goodness or

optimality of a selection rule, consider two quantities (cf. Lehmann [1]) $R(\underline{\theta}, \delta)$ and $S(\underline{\theta}, \delta)$. We define $S(\underline{\theta}, \delta) = P_{\underline{\theta}}(CS|\delta)$ and $R(\underline{\theta}, \delta) = \sum_{i=1}^k R^{(i)}(\underline{\theta}, \delta_i)$, where $R^{(i)}(\underline{\theta}, \delta_i) = P(\text{Selecting } \pi_i | \delta)$. Let S be the size of the selected subset. Thus $R(\underline{\theta}, \delta) = E(S|\delta)$. For a specified γ , ($0 < \gamma < 1$), we may restrict our attention to the class $\mathcal{L}$ of all δ such that

$$(1) \quad S(\underline{\theta}, \delta) \geq \gamma \text{ for } \underline{\theta} \in \bar{\Omega}.$$

We are interested in constructing an optimal procedure δ^0 in $\mathcal{L}$ which minimizes the supremum of $R(\underline{\theta}, \delta)$ over $\bar{\Omega}$ for all $\delta \in \mathcal{L}$, i.e.,

$$(2) \quad \sup_{\underline{\theta} \in \bar{\Omega}} R(\underline{\theta}, \delta^0) = \min_{\delta \in \mathcal{L}} \sup_{\underline{\theta} \in \bar{\Omega}} R(\underline{\theta}, \delta).$$

We restrict attention to those selection procedures which depend upon the observations only through a sufficient and maximal invariant statistic Z_{ij} which are defined as follows:

$$Z_{ij} = f(X_{i1}, \dots, X_{in_i}; X_{j1}, \dots, X_{jn_j}).$$

This Z_{ij} is based on the n_i and n_j observations from π_i and π_j ($i, j = 1, 2, \dots, k$), respectively. It is well known that the distribution of Z_{ij} depends only on τ_{ij} . For any i , let the joint density of Z_{ij} , $j \neq i$, be $p_{\underline{\theta}}(z_i)$. Let $p_{\underline{\theta}}(z_i)$ be denoted by $p_0(z_i)$ when $\tau_{i1} = \dots = \tau_{ik} = \tau_{ii} = \text{constant}$ and by $p_i(z_i)$ when $\tau_{i1} = \dots = \tau_{ik} = \tau_0$, $1 \leq i \leq k$. In a given problem the function f is so chosen as to indicate the measure of separation between the populations in a reasonable way. In case of the above normal means example, a choice of Z_{ij} might be $\bar{X}_i - \bar{X}_j$, where $\bar{X}_i = \frac{1}{n_i} \sum_{\ell=1}^{n_i} X_{i\ell}$ and $\bar{X}_j = \frac{1}{n_j} \sum_{\ell=1}^{n_j} X_{j\ell}$. Let ν be a σ -finite measure on R^{k-1} .

Oosterhoff [2] defines a monotone likelihood ratio for a random vector $\underline{x}$ with m components as follows: Let $\underline{\theta}$ be an m -dimensional vector of parameters. A partial ordering of points in R^m is defined by $\underline{x}_1 \preceq \underline{x}_2$, $\underline{x}_i' = (x_{i1}, \dots, x_{im})$, $i = 1, 2$, meaning that $x_{1j} \leq x_{2j}$ for $j = 1, 2, \dots, k$, and the inequality is strict for at least one component. The density $f_{\underline{\theta}}(\underline{x})$ has monotone likelihood ratio (MLR) if for all $\underline{\theta}_1 \preceq \underline{\theta}_2$, $f_{\underline{\theta}_2}(\underline{x})/f_{\underline{\theta}_1}(\underline{x})$ is nondecreasing in $\underline{x}$.

Now we state and prove a theorem which provides a solution to the restricted minimax problem as stated in (1) and (2) (cf Lehmann [1]).

Theorem: Suppose that the density $p_{\underline{\theta}}(\underline{z})$ has the MLR property. If $R(\underline{\theta}, \delta^0)$ is maximized at $\tau_{ij} = \tau_{ji} = \text{constant}$, for all i, j , where δ^0 is given by

$$\delta_i^0(\underline{z}_i) = \begin{cases} 1 & \text{if } p_i(\underline{z}_i) > c p_0(\underline{z}_i), \\ \lambda_i & = \\ 0 & < \end{cases}$$

such that $c(> 0)$ and λ_i are determined by $\int \delta_i^0 p_i = \gamma$, $1 \leq i \leq k$. Then $\delta^0 = (\delta_1^0, \dots, \delta_k^0)$ minimizes $\sup_{\underline{\theta} \in \Omega} R(\underline{\theta}, \delta)$ subject to $\inf_{\underline{\theta} \in \Omega} S(\underline{\theta}, \delta) \geq \gamma$.

Proof. For any $\delta \in \mathcal{C}$,

$\underline{\theta} \in \bar{\Omega}$ implies $\underline{\theta} \in \Omega_i$ for some i , thus

$$S(\underline{\theta}, \delta) = \int \delta_i(\underline{z}_i) p_{\underline{\theta}}(\underline{z}_i) dv(\underline{z}_i) \geq \min_{1 \leq i \leq k} \inf_{\underline{\theta} \in \Omega_i} \int \delta_i(\underline{z}_i) p_{\underline{\theta}}(\underline{z}_i) dv(\underline{z}_i).$$

We have

$$\inf_{\underline{\theta} \in \bar{\Omega}} S(\underline{\theta}, \delta) = \min_{1 \leq i \leq k} \inf_{\underline{\theta} \in \Omega_i} \int \delta_i(\underline{z}_i) p_{\underline{\theta}}(\underline{z}_i) dv(\underline{z}_i).$$

Hence for any $\delta \in \mathcal{C}$, $\inf_{\underline{\theta} \in \Omega_i} \int \delta_i(\underline{z}_i) p_{\underline{\theta}}(\underline{z}_i) dv(\underline{z}_i) \geq \gamma$, $1 \leq i \leq k$, and by the assumption that $\int \delta_i^0 p_i = \gamma$, it follows that

$$\int (\delta_i - \delta_i^0)(p_i - cp_0) \leq 0$$

which implies

$$\int \delta_i^0 p_0 \leq \int \delta_i p_0.$$

By the assumption, $\delta_i^0(z_i)$ is nondecreasing in z_i , we have

$$\inf_{\underline{\theta} \in \bar{\Omega}} S(\underline{\theta}, \delta^0) = \min_{1 \leq i \leq k} \int \delta_i^0 p_i = \gamma.$$

If $R(\underline{\theta}, \delta^0)$ is maximized at $\tau_{ij} = \tau_{ii} = \text{constant}$, for all i, j , then

$$\sup_{\underline{\theta} \in \bar{\Omega}} R(\underline{\theta}, \delta) \geq \sum_{i=1}^k \int \delta_i p_0 \geq \sum_{i=1}^k \int \delta_i^0 p_0 = \sup_{\underline{\theta} \in \bar{\Omega}} R(\underline{\theta}, \delta^0),$$

which completes the proof.

Example: Let $\pi_1, \pi_2, \dots, \pi_k$ be k independent normal populations with means $\theta_1, \dots, \theta_k$ and common variance $\sigma^2 = 1$. Our interest is to select a nonempty subset of the k populations containing the best. The ordered θ_i 's are denoted by $\theta_{[1]} \leq \dots \leq \theta_{[k]}$. It is assumed that there is no prior knowledge of the correct pairing of the ordered and the unordered θ_i 's. Our goal is to select a nonempty subset of the k populations so as to include the population associated with $\theta_{[k]}$.

Let $\bar{x}_i$, $1 \leq i \leq k$, denote the sample means of independent samples of size n from these populations. Let the joint likelihood function of $\bar{x}_i$, $i=1, 2, \dots, k$, be

$$g_{\underline{\theta}}(\underline{x}) = \prod_{j=1}^k g_{\theta_j}(\bar{x}_j),$$

where $g_{\theta_i}(\bar{x}_i) = \frac{\sqrt{n}}{\sqrt{2\pi}} e^{-\frac{n}{2}(\bar{x}_i - \theta_i)^2}$, $1 \leq i \leq k$. Let

$\tau_{ij} = \tau_{ij}(\underline{\theta}) = \theta_i - \theta_j$, $1 \leq j \leq k$, $j \neq i$, $\tau_{ii} = 0$, $\tau_0 = \Delta > 0$ and

$z_{ij} = \bar{x}_i - \bar{x}_j$, $j \neq i$. Let $\underline{z}_i' = (z_{i1}, \dots, z_{ik})$ and $\underline{\tau}_i' = (\tau_{i1}, \dots, \tau_{ik})$.

$$p_{\underline{\theta}}(\underline{z}_i) = (2\pi)^{\frac{k-1}{2}} |\Sigma|^{-1/2} \exp(-(\underline{z}_i - \underline{z}_i)' \Sigma^{-1} (\underline{z}_i - \underline{z}_i)),$$

where $\Sigma_{(k-1) \times (k-1)} = \frac{1}{n} \begin{pmatrix} 2 & & 1 \\ & \ddots & \\ 1 & & 2 \end{pmatrix}$ is the positive definite covariance matrix of Z_{ij} 's. We know that

$$\frac{p_i(\underline{z}_i)}{p_0(\underline{z}_i)} = \exp(\underline{z}_i' \Sigma^{-1} \underline{\Delta} + \underline{\Delta}' \Sigma^{-1} \underline{z}_i - \underline{\Delta}' \Sigma^{-1} \underline{\Delta})$$

is nondecreasing in z_{ij} , $j \neq i$, where $\underline{\Delta}' = (\dots, \Delta)$. And

$$\frac{p_i(\underline{z}_i)}{p_0(\underline{z}_i)} > c$$

is equivalent to

$$(3) \quad \bar{x}_i > \frac{1}{k-1} \sum_{j \neq i} \bar{x}_j + d.$$

For any i , let $\mu_i = \sum_{j \neq i} (\theta_j - \theta_i)$, we have

$$\mu_i = \mu - k\theta_i$$

where $\mu = \sum_{j=1}^k \theta_j$. Hence, $\mu_1 = \mu_2 = \dots = \mu_k$ if and only if $\theta_1 = \theta_2 = \dots = \theta_k$. We order μ_i 's as $\mu_{[1]} \leq \dots \leq \mu_{[k]}$. Since

$$\begin{aligned} R(\underline{\theta}, \delta) &= \sum_{i=1}^k P(\bar{X}_i > \frac{1}{k-1} \sum_{j \neq i} \bar{X}_j + d) \\ &= \sum_{i=1}^k \int \phi(\sqrt{k-1}(y - \frac{1}{k-1} \mu_i - d)) d\phi(y) \\ &= f(\mu_1, \dots, \mu_k) \\ &\leq f(\mu_{[1]}, \dots, \mu_{[1]}) \\ &= f(\theta_{[k]}, \dots, \theta_{[k]}) \end{aligned}$$

$$= f(0, \dots, 0)$$

$$= R(\underline{0}, \delta),$$

hence, it follows that the supremum of $R(\underline{\theta}, \delta^0)$ over Ω occurs at $\theta_1 = \dots = \theta_k$, i.e., $r_{ij} = r_{ii} = 0$. Thus the result of the theorem can be applied.

Note that the above procedure (3) is a rule of the type proposed by Seal [3] to select a subset containing the population associated with the largest θ_i 's.

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