



Research article

Solving ANOVA problem with restricted Type I and Type II error rates

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Abstract: The problem of solving the ordered one-way analysis of variance (ANOVA) (which consists of comparing a set of normal means) with restricted Type I and Type II error rates is considered in this paper. This case is more complicated than unordered one-way ANOVA because the detection of the monotonicity of means restrictions is necessary. To solve this issue, one of the possible formulations of the constrained Bayesian method (CBM) is applied here using the concept of directional hypotheses. The cases of known and unknown variances are examined. For unknown variances, the maximum likelihood ratio and Stein's Methods were used to overcome the problem connected with the complexity of hypotheses. The correctness of the developed methods and high quality (in comparison with existing methods) of obtained results were demonstrated by computing results of the simulated concrete scenarios. Moreover, the offered method controlled not only one Type of error, as methods do, but both Type I and Type II methods.

Keywords: ANOVA; hypothesis testing; constrained Bayesian method; Type I and Type II error rates; maximum likelihood ratio; Stein's method

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1. Introduction

The problem of comparing the means of several normally distributed random variables has attracted researchers' attention from the middle of the last century. Such problems arise in many practical applications such as when comparing treatment effects in clinical trials, or experimental studies to compare the effects of different experiments, or when comparing the failure rate of a machine with its age. In the framework of ANOVA, the problem can be formulated as follows [1]. Let us consider k normal populations $N(\mu_i, \sigma_i^2)$, $i = 1, \dots, k$. Let X_{i1}, \dots, X_{in_i} be a random sample from $N(\mu_i, \sigma_i^2)$, independent from each other. The problem is to test hypotheses

$$H_0 : \mu_1 = \dots = \mu_k \text{ vs. } H_a : \mu_i \neq \mu_j \text{ for some } i \neq j. \quad (1)$$

In some studies, researchers may believe that the true means are simply ordered ($\mu_1 \leq \dots \leq \mu_k$) before observing the data. For example, the treatment means may be a sequence of increasing dose levels of a drug. In this case, ANOVA takes the form

$$H_0 : \mu_1 = \dots = \mu_k \text{ vs. } H_a : \mu_1 \leq \dots \leq \mu_k, \quad (2)$$

with at least one strict inequality. This statement has received considerable attention in the statistical literature (see, for example, [1–5]).

The simplest case $k = 2$, i.e., the case with only two normal populations, called the Behrens–Fisher problem, is well studied. For its solution, at statement (1), a simple and accurate method is introduced in [6,7], which gives good results in both cases whether variances are equal or not. For the arbitrary case; moreover, when $k > 2$, a compact but quite good review of the methods used to solve problem (1) is given in [8], on the basis that there is a procedure that gives satisfactory results by the Type I error rate for all sample sizes, where k and parameter configurations do not exist [8,9]. To improve the situation, a parametric bootstrap (PB) approach is offered in [8,9] and it is compared with the most promoted tests: The Welch test, the James ([10]) second-order test, and the generalized F (GF) test [11,12]. The comparison was realized by Type I error rate and powers using Monte Carlo simulation, showing that the best, based on the sample sizes, values of the error variances, and the number of means being compared is a PB test. A little worse is the James second-order test. The Welch test and the GF test have fairly poor Type I errors when the value of k is medium or large and/or the sample sizes are small. In computed examples of [8], the minimum value of Monte Carlo estimates of Type I error rates is equal to 0.04 even when $k = 2$, meaning it is quite high. For other k , as a rule, it is greater.

For statement (2), a one-sided studentized range test is offered in Hayter [3]. It provides simultaneous one-sided lower confidence bounds for the ordered pairwise comparisons $\mu_i - \mu_j$, $1 \leq j < i \leq k$. Similar inference procedures are discussed in [13–17]. In some, the strong order of arranging the mathematical expectations is considered, and in others, the assumption that μ_i 's follow a simple ordering is not made. The familywise error rate (FWER) is used as an optimality criterion in these cases, the computation of which is complicated for large k . To overcome this problem, the empirical Bayesian approach of multiple testing is offered in [1]. Its essence consists of the consideration of the families of hypotheses

$$H_{i0} : \mu_{i+1} - \mu_i = 0 \text{ vs. } H_{i1} : \mu_{i+1} - \mu_i > 0, \quad i = 1, \dots, k-1, \quad (3)$$

from a Bayesian view point. To test (3), there is a new step-down multiple testing procedure, which is a step-down version of the single-step procedure [18]. It controls the familywise error rate better than a single-step procedure, which provides a simple comparison of each p -value with a single critical value. For this reason, the step-down version is more powerful than the single-step procedure.

The heart of the single-step procedure consists of the rejection of the H_{i0} hypothesis if $p_i \leq \alpha$, where p_i is H_{i0} hypothesis corresponding p -value, and $\alpha \in (0,1)$ is the significant level. In the step-down test, the differences $T_i = \mu_{i+1} - \mu_i$ and the appropriate hypotheses are in ascending order. Testing hypothesis (3) begins with the largest T -value, i.e., with $T_{(k-1)}$. If for some $T_{(k-i)}$ hypothesis $H_{(k-i)0}$ is accepted, then testing is stopped, and all the remaining hypotheses whose order number is less than $(k-i)$ are accepted; otherwise, $H_{(k-i)0}$ is rejected, and the $(i+1)$ th step is transited. In [1], the critical constants for each stage chosen so that FWER is controlled at a pre-specified level α . Because the empirical Bayesian approach is used in this method, which is based on the resampling techniques, it asymptotically controls the FWER.

An alternative method for testing hypothesis (2), similar to the parametric bootstrap method of [8] to improve the Type-I error rate, is offered in [4]. In particular, this method is an extended Hayter's test [3] for heterogeneous sample sizes and variances, and uses the idea of the PB offered in [8], based on the simulation of the sample from the null distribution with unbiased estimators of the parameters. The non-significant improvement of Type I error compared to existing methods of the Welch test (WT) [19]), the \bar{E}^2 test (which is a counterpart of the F test in the ordered case [5]) and Chen's one-stage method ([20]) for large sample sizes is shown through the modeling for the introduced novelty.

For justice, it should be said that all the considered tests have quite high Type I and Type II error rates, the values that depend on the parameters of the problem under consideration (the values of k , variances and sample sizes) and their control is impossible at a predetermined level. Therefore, to solve problems (1) and (2), we aim to develop a method that will enable us to make a decision by restricting Type I and Type II error rates to a predetermined level. To achieve this goal, the CBM is used below [21,22].

The structure of the rest of the paper is as follows. The application of CBM to solve the stated problem is given in Section 2, where formulas and criteria for testing individual and multiple Hypotheses defined by (6) are given. Algorithms for computation for known and unknown variances are developed in Section 3. Computation results for practical examples at known and unknown variances when the maximum likelihood ratio and Stein's Methods are used for overcoming the problem caused by complexity of hypotheses, are presented in Section 4. A short discussion of the obtained results is offered in Section 5, and a conclusion is presented in Section 6.

2. Method of testing multiple directional hypotheses for solving the stated problem

Without loss of generality, we can suppose that all μ_i ($i = 1, \dots, k$) are positive, because if this condition is not fulfilled, we can always achieve it by moving the origin.

Let us rewrite hypothesis (1) as follows

$$H_0 : \theta_1 = \theta_2 = \dots = \theta_{k-1} = 0 \quad \text{vs.} \quad H_a : \theta_i \neq 0, \text{ for some } i \in (1, \dots, k-1). \quad (4)$$

where $\theta_i = \mu_{i+1} - \mu_i$, $i = 1, \dots, k-1$. In this case, hypothesis (2) takes the form

$$H_0 : \theta_1 = \theta_2 = \dots = \theta_{k-1} = 0 \quad \text{vs.} \quad H_a : \theta_i \geq 0, \text{ with at least one strict inequality.} \quad (5)$$

Using the concept of directional hypotheses [22–24], similarly to (3), hypothesis (4) can be considered as a set of directional hypotheses

$$H_0^i : \theta_i = 0 \quad \text{vs.} \quad H_-^i : \theta_i < 0 \quad \text{or} \quad H_+^i : \theta_i > 0, \quad i = 1, \dots, k-1. \quad (6)$$

Statement (6) enables us to test hypotheses (1) and (2) (which are the same as hypotheses (4) and (5)) simultaneously, using techniques developed in [22,23]. In particular: After testing hypothesis (6), if for even one value of i , the left sided hypothesis is accepted among accepted alternatives, accept alternative hypothesis in (4); and if all the accepted alternative hypotheses are right sided, accept alternative hypothesis in (5); otherwise accept the null hypothesis.

Let us recall that for testing hypothesis (6), we have the following information: X_{i1}, \dots, X_{in_i} - a random sample from $N(\mu_i, \sigma_i^2)$, $i = 1, \dots, k$, on the basis of which we compute

$$\bar{X}_i = \frac{1}{n_i} \sum_{j=1}^{n_i} X_{ij} \quad \text{and} \quad S_i^2 = \frac{1}{n_i - 1} \sum_{j=1}^{n_i} (X_{ij} - \bar{X}_i)^2, \quad i = 1, \dots, k, \quad (7)$$

where \bar{X}_i and S_i^2 are the i th sample mean and sample variance.

To test hypotheses concerning parameter θ_i , sufficient statistics is $\bar{Y}_i = \bar{X}_{i+1} - \bar{X}_i$, where the distribution is $N(\bar{y}_i | \theta_i, \sigma_i^2/n_i + \sigma_{i+1}^2/n_{i+1})$, $i = 1, \dots, k-1$, because of independence of \bar{X}_{i+1} and \bar{X}_i . Parameter $\theta_i = 0$ at basic hypothesis H_0^i , and $\theta_i < 0$ or $\theta_i > 0$, at alternative hypotheses H_-^i or H_+^i , accordingly.

2.1. Testing a subset of individual hypotheses of (6)

For ease of presentation and understanding, let us first consider one subset of individual hypotheses of (6), and for simplicity, omit index i . Let us use one of the possible formulations of CBM, in particular, restrictions on posterior probabilities of rejected true hypotheses (Task 7) considered in [22,23]. In this formulation, the Type II error rate is minimized (that means the power is maximized), and the Type I error rate is restricted. This means, that the statement of the problem is the following [21,23]:

$$G_\delta = \max_{\{\Gamma_-, \Gamma_0, \Gamma_+\}} \{K_0 \cdot [p(H_-) \cdot P(x \in \Gamma_- | H_-) + p(H_0) \cdot P(x \in \Gamma_0 | H_0) + p(H_+) \cdot P(x \in \Gamma_+ | H_+)]\} \quad (8)$$

subject to

$$\begin{aligned}
K_1 \cdot [p(H_0) \cdot P(x \in \Gamma_- | H_0) + p(H_+) \cdot P(x \in \Gamma_- | H_+)] &\leq r_7^-, \\
K_1 \cdot [p(H_-) \cdot P(x \in \Gamma_0 | H_-) + p(H_+) \cdot P(x \in \Gamma_0 | H_+)] &\leq r_7^0, \\
K_1 \cdot [p(H_-) \cdot P(x \in \Gamma_+ | H_-) + p(H_0) \cdot P(x \in \Gamma_+ | H_0)] &\leq r_7^+.
\end{aligned} \tag{9}$$

Here, for each subset of individual hypotheses of (6), the following loss functions are used

$$L_1(H_i, \delta_j(x) = 1) = \begin{cases} 0 & \text{at } i = j, \\ K_1 & \text{at } i \neq j; \end{cases} \text{ and } L_2(H_i, \delta_j(x) = 0) = \begin{cases} K_0 & \text{at } i = j, \\ 0 & \text{at } i \neq j; \end{cases} \tag{10}$$

where K_1 is the loss of incorrectly accepted H_j when H_i is true and K_0 the loss of incorrectly rejected H_j at testing it versus H_i , $i, j \in (-, 0, +)$.

Using the undetermined Lagrange multiplier method and the concepts of posterior probabilities, to solve problems (8) and (9), we have [23]

$$\begin{aligned}
\Gamma_- &= \left\{ x : K_1 \cdot (p(H_0 | x) + p(H_+ | x)) < \frac{1}{\lambda_7^-} \cdot K_0 \cdot p(H_- | x) \right\}, \\
\Gamma_0 &= \left\{ x : K_1 \cdot (p(H_- | x) + p(H_+ | x)) < \frac{1}{\lambda_7^0} \cdot K_0 \cdot p(H_0 | x) \right\}, \\
\Gamma_+ &= \left\{ x : K_1 \cdot (p(H_- | x) + p(H_0 | x)) < \frac{1}{\lambda_7^+} \cdot K_0 \cdot p(H_+ | x) \right\},
\end{aligned} \tag{11}$$

where Lagrange multipliers λ_7^- , λ_7^0 , and λ_7^+ are determined so that in (9), the equalities hold.

The most common criteria of estimation of the quality of testing directional hypotheses are the mixed directional false discovery rate (*mdFDR*) and the false acceptance rate (*FAR*), which have the following forms [21,22]

$$mdFDR = P(\bar{y} \in \Gamma_- | H_+) + P(\bar{y} \in \Gamma_- | H_0) + P(\bar{y} \in \Gamma_+ | H_-) + P(\bar{y} \in \Gamma_+ | H_0), \tag{12}$$

and

$$FAR = P(\bar{y} \in \Gamma_0 | H_-) + P(\bar{y} \in \Gamma_0 | H_+), \tag{13}$$

respectively.

According to the theorems proved in [23], if the condition $r_7^- + r_7^+ = q \cdot K_1 \cdot P_{\min}$ is satisfied, where $0 < q < 1$, $P_{\min} = \min\{p(H_-, H_0, H_+)\}$, then the following condition $mdFDR \leq q$ is fulfilled, and if the condition $r_7^- + r_7^0 + r_7^+ = q \cdot K_1 \cdot P_{\min}$ is satisfied, the following condition $mdFDR + FAR \leq q$ is

fulfilled. Also, when λ_7^0 is chosen so that in the second condition of (9) equality takes place, the following is fulfilled:

$$FAR \leq \frac{1}{P'_{\min}} \cdot \frac{r_7^0}{K_1}, \quad (14)$$

where $P'_{\min} = \min \{p(H_-), p(H_+)\}$.

For the case of directional hypotheses, the Type I error rate (*TIER*) can be determined as follows

$$TIER = P(\bar{y} \in \Gamma_- | H_0) + P(\bar{y} \in \Gamma_+ | H_0). \quad (15)$$

Comparing (12) and (15), it is clear that $mdFDR \geq TIER$.

Let us define Type II error rates (*TIER*) as follows

$$TIER = P(\bar{y} \in \Gamma_0 | H_-) + P(\bar{y} \in \Gamma_0 | H_+). \quad (16)$$

Comparing (13) and (16), we are convincing that $FAR = TIER$. Thus, to fulfill the condition $\frac{1}{P'_{\min}} \cdot \frac{r_7^0}{K_1} = q$, the following takes place $FAR = TIER \leq q$, where $0 < q < 1$.

Theorem 1. *CBM 7 with restriction levels (9), to satisfy a condition $(r_7^- + r_7^+) / (K_1 \cdot p(H_0)) = q$, where $0 < q < 1$, ensures a decision rule with Type I error rate less or equal to q , i.e., with the condition $TIER \leq q$.*

Proof. From the first and the third conditions of the restrictions of (9), we write the following

$$\begin{aligned} P(\bar{y} \in \Gamma_- | H_0) &\leq \left[\frac{r_7^-}{K_1} - p(H_+) \cdot P(\bar{y} \in \Gamma_- | H_+) \right] \cdot \frac{1}{p(H_0)} = \\ &= \frac{r_7^-}{K_1 \cdot p(H_0)} - \frac{p(H_+)}{p(H_0)} \cdot P(\bar{y} \in \Gamma_- | H_+), \end{aligned}$$

and

$$\begin{aligned} P(\bar{y} \in \Gamma_+ | H_0) &\leq \left[\frac{r_7^+}{K_1} - p(H_-) \cdot P(\bar{y} \in \Gamma_+ | H_-) \right] \cdot \frac{1}{p(H_0)} = \\ &= \frac{r_7^+}{K_1 \cdot p(H_0)} - \frac{p(H_-)}{p(H_0)} \cdot P(\bar{y} \in \Gamma_+ | H_-). \end{aligned}$$

From here, and considering (15), we are convincing in the correctness of theorem 1.

Because of the specific nature of the acceptance regions of CBM [21–23], it can so happen that making a simple decision on the basis of existed information becomes impossible, e.g., when the test statistic belongs to the intersection areas of the acceptance regions or does not belong to any of these

regions. In such a situation, it becomes impossible to make a simple decision with a specified confidence level on the basis of the information, and more information is required to achieve this. If acquiring more information is impossible, then the restriction levels in (9) must be changed until a simple decision can be made. When acquiring more information is possible, we appeal to the sequential experiment, i.e., to increase a sample size and apply *Procedure A*.

Procedure A

The statistics for testing a subset of individual hypotheses of (6) is $\bar{y}_i = \bar{x}_{i+1} - \bar{x}_i$, which depends on two initial samples $x_{i1}, \dots, x_{in}, \dots$ and $x_{i+1,1}, \dots, x_{i+1,n}, \dots$. Let us denote the test statistics on the basis of n observations by $\bar{y}_{i,n}$. Then the sequential procedure is as follows:

Step 1

- if $\bar{y}_{i,n}$ belongs to only region Γ_-^i , accept hypothesis H_-^i ,
- if $\bar{y}_{i,n}$ belongs to only region Γ_0^i , accept hypothesis H_0^i ,
- if $\bar{y}_{i,n}$ belongs to only region Γ_+^i , accept hypothesis H_+^i ,
- otherwise continue sampling; collect next observations and compute new test statistics $\bar{y}_{i,n+1}$;

Step 2

- if $\bar{y}_{i,n+1}$ belongs to only region Γ_-^i , accept hypothesis H_-^i ,
- if $\bar{y}_{i,n+1}$ belongs to only region Γ_0^i , accept hypothesis H_0^i ,
- if $\bar{y}_{i,n+1}$ belongs to only region Γ_+^i , accept hypothesis H_+^i ,
- otherwise, continue sampling; collect observations and compute test statistics etc.

The sampling continues until the test statistics belong to only one hypothesis acceptance region.

Note 1: Obviously, the sample size at the beginning of the sequential test is equal to any particular value, i.e., $n = n^*$ (in particular, it can be $n^* = 1$), and if the desired level of reliability of making a decision is achievable for this amount of information, this corresponds to the parallel experiment on which the testing process finishes. In case of continuing sampling, the parallel experiment generalizes to the sequential experiment naturally.

2.2. Testing of multiple hypotheses (6)

For testing multiple directional hypotheses (6), the concept of the total mixed directional false discovery rate (*tmdFDR*) is introduced in [23,25]

$$tmdFDR = \sum_{i=1}^{k-1} mdFDR_i. \quad (17)$$

To guarantee level q , in testing hypothesis (6), we have to consider $k - 1$ subsets of directional hypotheses. Then, for each of them, we use procedure A described above to provide a level of q_i for i th subset of hypotheses, so that $\sum_{i=1}^{k-1} q_i = q$ is achieved.

We act similarly to provide a level q for the total Type I error rate (*TTIER*) and for the total

Type II error rate ($TTIER$). Namely, we provide q_i , the level of the appropriate criteria for the i th subset of the individual directional hypotheses. As a result, we have

$$TTIER = \sum_{i=1}^{k-1} TIER_i, \quad (18)$$

$$TTIER = \sum_{i=1}^{k-1} TTIER_i, \quad (19)$$

where $TIER_i$ and $TTIER_i$ are the Type I error rate and the Type II error rate, respectively, of the i th subset of directional hypotheses [23,26].

The values of q_i in all three cases (for $tmdFDR$, for $TTIER$, and for $TTIER$) can be chosen to be equal, i.e., $q_i = q/(k-1)$ or different, e.g., inversely proportional to the informational distances between the tested hypotheses in the subsets of directional hypotheses [23].

To test multiple hypotheses, two family-wise error rates, type I ($FWER_I$) and type II ($FWER_{II}$), are used (see, for example, [26–28]), which are the same as $TTIER$ and $TTIER$ in the considered case, i.e., we have $TTIER = FWER_I$ and $TTIER = FWER_{II}$.

In all three considered cases, with the restriction of $tmdFDR$, $TTIER$, and $TTIER$ on the desired levels, we use the above described sequential *Procedure A*, where the sampling continues until a simple decision is made for all the subsets of multiple hypotheses (6). The stopping rules remain the same as in [23]. Theorems 3.2 and 3.3 of the work [23], proving the appropriateness of the stopping rule, are in force for the considered directional hypotheses as well.

As mentioned in [24], “Currently, in many real-life applications, we indeed encounter situations where the number of individual hypotheses in the set of multiple hypotheses (6) is very big, i.e., when data is big. In such a situation, determination of Lagrange multipliers for each subset of an individual hypothesis requires a long time for computation. Though the computation of Lagrange multipliers is completed in the preparatory stage before making a decision, still the reduction of computation time is important for many practical applications from the operational and cost considerations.” For this purpose, Theorem 4 is proved in [24] in accordance with the testing of all of subsets of individual hypotheses of (6) and can be used once with the same Lagrange multipliers, determined for a subset of individual hypothesis with lowest divergence among directional hypotheses at the level $q_i = q/(k-1)$, satisfying condition $\sum_{i=1}^{k-1} q_i = q$ provides the total mixed directional false discovery rate (17), as well as the total Type I error rate (18) and the total Type II error rate (19), restricting all of them with level q .

3. Algorithms for computation

3.1. Variances are known

Let us suppose that the variances of X_i that is σ_i^2 , $i=1, \dots, k$, are known. To realize the above developed method, the concretization of distribution laws of the statistics \bar{Y}_i at hypotheses $H_i^{(0)}$,

$H_i^{(-)}$ and $H_i^{(+)}$ is necessary, i.e., it is necessary to determine densities $p(\bar{y}_i | H_i^{(0)})$, $p(\bar{y}_i | H_i^{(-)})$, and $p(\bar{y}_i | H_i^{(+)})$.

As mentioned in the above \bar{Y}_i is normally distributed. Therefore, taking into account the intervals of definition of the mathematical expectation at different hypotheses, we can suppose that $p(\bar{y}_i | H_i^{(0)}) = N(\bar{y}_i | 0, \bar{\sigma}_i^2)$ at $H_i^{(0)}$ and $p(\bar{y}_i | H_i^{(-)})$ and $p(\bar{y}_i | H_i^{(+)})$ are the truncated normal densities $N(0, \omega_0^{-1} \bar{\sigma}_i^2)$ (ω_0 known) over $(-\infty, 0)$ and $(0, +\infty)$, respectively [23,29]. Here, $\bar{\sigma}_i^2 = (\sigma_i^2 + \sigma_{i+1}^2)/n = \mathcal{G}_i^2/n$, were $\mathcal{G}_i^2 = (\sigma_i^2 + \sigma_{i+1}^2)$. Thus, we have [24]

$$\begin{aligned}
 p(\bar{y}_i | H_i^{(0)}) &= \frac{\sqrt{n}}{\sqrt{2\pi} \cdot \mathcal{G}_i} \cdot \exp\left\{-\frac{n\bar{y}_i^2}{2\mathcal{G}_i^2}\right\}, \\
 p(\bar{y}_i | H_i^{(-)}) &= \int_{-\infty}^0 \frac{\sqrt{n}}{\sqrt{2\pi} \cdot \mathcal{G}_i} \exp\left\{-\frac{n(\bar{y}_i - \theta_i)^2}{2\mathcal{G}_i^2}\right\} \cdot \frac{2\sqrt{\omega_0}}{\sqrt{2\pi} \cdot \mathcal{G}_i} \cdot \exp\left\{-\frac{\omega_0^2 \theta_i^2}{2\mathcal{G}_i^2}\right\} d\theta_i = \\
 &= \frac{2\sqrt{\omega_0}}{\sqrt{n + \omega_0}} \cdot \frac{\sqrt{n}}{\sqrt{2\pi} \cdot \mathcal{G}_i} \cdot (1 - \Phi(u_i)) \exp\left\{-\frac{\omega_0 u_i^2}{2n}\right\}, \\
 p(\bar{y}_i | H_i^{(+)}) &= \int_0^{+\infty} \frac{\sqrt{n}}{\sqrt{2\pi} \cdot \mathcal{G}_i} \exp\left\{-\frac{n(\bar{y}_i - \theta_i)^2}{2\mathcal{G}_i^2}\right\} \cdot \frac{2\sqrt{\omega_0}}{\sqrt{2\pi} \cdot \mathcal{G}_i} \cdot \exp\left\{-\frac{\omega_0^2 \theta_i^2}{2\mathcal{G}_i^2}\right\} d\theta_i = \\
 &= \frac{2\sqrt{\omega_0}}{\sqrt{n + \omega_0}} \cdot \frac{\sqrt{n}}{\sqrt{2\pi} \cdot \mathcal{G}_i} \cdot \Phi(u_i) \exp\left\{-\frac{\omega_0 u_i^2}{2n}\right\}. \tag{20}
 \end{aligned}$$

For these distributions, hypotheses acceptance regions (11) for observations \bar{y}_i are the following [23]

$$\begin{aligned}
 \Gamma_- &= \left\{ \bar{y}_i : \frac{p(H_0) \cdot \sqrt{n + \omega_0} \cdot \exp\{-u_i^2/2\} + p(H_+) \cdot 2 \cdot \sqrt{\omega_0} \cdot \Phi(u_i)}{p(H_-) \cdot 2 \cdot \sqrt{\omega_0} \cdot (1 - \Phi(u_i))} < \frac{1}{\lambda_7^-} \cdot \frac{K_0}{K_1} \right\}, \\
 \Gamma_0 &= \left\{ \bar{y}_i : \frac{p(H_-) \cdot 2 \cdot \sqrt{\omega_0} \cdot (1 - \Phi(u_i)) + p(H_+) \cdot 2 \cdot \sqrt{\omega_0} \cdot \Phi(u_i)}{p(H_0) \cdot \sqrt{n + \omega_0} \cdot \exp\{-u_i^2/2\}} < \frac{1}{\lambda_7^0} \cdot \frac{K_0}{K_1} \right\}, \\
 \Gamma_+ &= \left\{ \bar{y}_i : \frac{p(H_-) \cdot 2 \cdot \sqrt{\omega_0} \cdot (1 - \Phi(u_i)) + p(H_0) \cdot \sqrt{n + \omega_0} \cdot \exp\{-u_i^2/2\}}{p(H_+) \cdot 2 \cdot \sqrt{\omega_0} \cdot \Phi(u_i)} < \frac{1}{\lambda_7^+} \cdot \frac{K_0}{K_1} \right\}. \tag{21}
 \end{aligned}$$

Here, n is the quantity of \bar{y}_i on the basis of which hypothesis (6) is tested using sequential Procedure A, $u_i = n \cdot \bar{y}_i / (\mathcal{G}_i \sqrt{n + \omega_0})$, and $\Phi(\cdot)$ is the standard normal distribution function.

The Lagrange multipliers are determined so that in condition (9), the equalities hold. For the solution of the relevant equations, the suitable probability integrals are computed by the Monte-Carlo method (see, for example, [21,23]).

3.2. Variances are not known

Let us suppose that the variances of X_i that is σ_i^2 , $i = 1, \dots, k$, are not known. In this case, we can consider two methods based on the maximum likelihood ratio and Stein's method [30–34]. The essence of these methods consist of the following. In the maximum likelihood ratio approach, the density with unknown parameters' values is replaced with the maximum of the density over all possible values of these parameters at the truth of testing hypotheses, computed by existing samples. Stein's method integrates the density over unknown parameters using special measures to obtain the density of the maximum invariant statistic, which can then be used to analyze the problems; for example, to find the uniformly most powerful invariant test [32].

The method based on the maximum likelihood ratio can be applied when the numbers of observations n_i , $i = 1, \dots, k$, are large, which enable us to divide every subset of the observations into two parts $x_{i,1}, \dots, x_{i,m_i}$ and $x_{i,m_i+1}, \dots, x_{i,n_i}$, where $m_i < n_i$. Using m_i observations, $\bar{S}_i^2 = S_i^2 + S_{i+1}^2$, $i = 1, \dots, k-1$, estimators of the variance $\bar{\sigma}_i^2$ of the statistics $\bar{Y}_i = \bar{X}_{i+1} - \bar{X}_i$ are computed and are used in formulae (20) and (21) instead of $\bar{\sigma}_i^2$. Statistics u_i are computed using observations $x_{i,m_i+1}, \dots, x_{i,n_i}$, $i = 1, \dots, k$, to make decisions based on hypotheses acceptance regions (21) when testing hypothesis (6).

When applying Stein's method, to average the influence of $\bar{\sigma}_i^2$, it is necessary to use a priori distribution of \bar{S}_i^2 , which is difficult to determine functionally with known parameters at the considering case. Therefore, we conduct the following: Construct a confidence interval of $\bar{S}_i^2 = S_i^2 + S_{i+1}^2$ with the given confidence probability $1 - \alpha$ ($0 < \alpha < 1$) and assume that \bar{S}_i^2 takes values from this interval according to a uniform distribution.

Finding confidence intervals of \bar{S}_i^2 , $i = 1, \dots, k-1$ is not difficult. It looks like:

$$\bar{S}_{i,L}^2 < \bar{\sigma}_i^2 < \bar{S}_{i,U}^2, \quad (22)$$

where

$$\begin{aligned} \bar{S}_{i,L}^2 &= \frac{(m_i - 1)S_i^2}{\chi_{m_i-1, 1-\alpha}^2} + \frac{(m_{i+1} - 1)S_{i+1}^2}{\chi_{m_{i+1}-1, 1-\alpha}^2}, \\ \bar{S}_{i,U}^2 &= \frac{(m_i - 1)S_i^2}{\chi_{m_i-1, \alpha}^2} + \frac{(m_{i+1} - 1)S_{i+1}^2}{\chi_{m_{i+1}-1, \alpha}^2} \end{aligned} \quad (23)$$

where $\chi_{m_i-1, \alpha}^2$ is a quantile of the α -th order of the chi-square distribution with degree of freedom $m_i - 1$.

To overcome the uncertainty in the densities (20) caused by not knowing the value of the variance, we are averaging the effect of the variance on these densities by integrating them over the confidence interval of the variance with uniform density. In this case, conditional distribution densities (20) take the form

$$\begin{aligned}
p(\bar{y}_i | H_i^0) &= k_i \int_{\bar{S}_{i,L}^2}^{\bar{S}_{i,u}^2} \frac{1}{x} \cdot \exp\left\{-\frac{a_i}{x^2}\right\} dx, \\
p(\bar{y}_i | H_i^-) &= d_i \int_{-\infty}^0 \int_{\bar{S}_{i,L}^2}^{\bar{S}_{i,u}^2} \frac{1}{x^2} \exp\left\{-\frac{a_i - b_i z + c_i z^2}{x^2}\right\} dz dx, \\
p(\bar{y}_i | H_i^+) &= d_i \int_0^{+\infty} \int_{\bar{S}_{i,L}^2}^{\bar{S}_{i,u}^2} \frac{1}{x^2} \exp\left\{-\frac{a_i - b_i z + c_i z^2}{x^2}\right\} dz dx,
\end{aligned} \tag{24}$$

where $k_i = \frac{\sqrt{n_i}}{\sqrt{2\pi(\bar{S}_{i,u}^2 - \bar{S}_{i,L}^2)}}$, $d_i = \frac{\sqrt{n_i \cdot \omega_0}}{\pi(\bar{S}_{i,u}^2 - \bar{S}_{i,L}^2)}$, $a_i = \frac{n_i \bar{y}_i^2}{2}$, $b_i = n_i \bar{y}_i$, $c_i = \frac{n_i + \omega_0}{2}$.

Let us denote:

$$\begin{aligned}
p^0(\bar{y}_i | H_i^0) &= \int_{\bar{S}_{i,L}^2}^{\bar{S}_{i,u}^2} \frac{1}{x} \cdot \exp\left\{-\frac{a_i}{x^2}\right\} dx, \\
p^-(\bar{y}_i | H_i^-) &= \int_{-\infty}^0 \int_{\bar{S}_{i,L}^2}^{\bar{S}_{i,u}^2} \frac{1}{x^2} \exp\left\{-\frac{a_i - b_i z + c_i z^2}{x^2}\right\} dz dx, \\
p^+(\bar{y}_i | H_i^+) &= \int_0^{+\infty} \int_{\bar{S}_{i,L}^2}^{\bar{S}_{i,u}^2} \frac{1}{x^2} \exp\left\{-\frac{a_i - b_i z + c_i z^2}{x^2}\right\} dz dx,
\end{aligned} \tag{25}$$

Hypotheses acceptance regions (11) take the following form using these denotations:

$$\begin{aligned}
\Gamma_0 &= \left\{ \bar{y}_i : \frac{p(H_-^i) \cdot p^-(\bar{y}_i | H_-^i) + p(H_+^i) \cdot p^+(\bar{y}_i | H_+^i)}{p(H_0^i) \cdot p^0(\bar{y}_i | H_0^i)} \cdot \sqrt{\frac{2\omega_0}{\pi}} < \frac{1}{\lambda_7^0} \cdot \frac{K_0}{K_1} \right\}, \\
\Gamma_- &= \left\{ \bar{y}_i : \frac{\sqrt{\frac{\pi}{2\omega_0}} \cdot p(H_0^i) \cdot p^0(\bar{y}_i | H_0^i) + p(H_+^i) \cdot p^+(\bar{y}_i | H_+^i)}{p(H_-^i) \cdot p^-(\bar{y}_i | H_-^i)} < \frac{1}{\lambda_7^-} \cdot \frac{K_0}{K_1} \right\}, \\
\Gamma_+ &= \left\{ \bar{y}_i : \frac{p(H_-^i) \cdot p^-(\bar{y}_i | H_-^i) + \sqrt{\frac{\pi}{2\omega_0}} \cdot p(H_0^i) \cdot p^0(\bar{y}_i | H_0^i)}{p(H_+^i) \cdot p^+(\bar{y}_i | H_+^i)} < \frac{1}{\lambda_7^+} \cdot \frac{K_0}{K_1} \right\},
\end{aligned} \tag{26}$$

where the Lagrange multipliers are determined, so that in conditions (9), the equalities hold.

4. Computation results

To compare the offered method with the known ones, we present computation outcomes of the examples considered in ([4], p. 26, Tables 2 and 6) in Tables below where the results are obtained by the method offered above (see Item 3.1) and the Hayter PB method. As a priori probabilities, we take $p(H_-) = p(H_0) = p(H_+) = 1/3$, coefficient $\omega_0 = 4$ for the results given in Tables 1–5, and $\omega_0 = 13$ for Stein's method for true alternative hypotheses the results of which are given in Table 6. To keep $mdFDR$, $TIER$, and $TTIER$ at $q = 0.05$, to test individual hypotheses of (6), we have to choose the following restriction levels in (9) for each subset of the hypotheses: $r_7^- = r_7^0 = r_7^+ = 0.0083(3)$. To test multiple hypotheses with the $k-1$ subset of individual hypotheses for ensuring the same restriction levels of the criteria $mdFDR$, $TIER$, and $TTIER$, we have to choose the following restriction levels: $r_7^- = r_7^0 = r_7^+ = 0.0083(3)/(k-1)$. To test of all subsets of individual hypotheses, we use one and the same Lagrange multipliers, which are determined for a subset of individual hypotheses with the lowest divergence among directional hypotheses [24]. Because of $k-1=3$, the restriction levels in (9) are equal to $0.0027(7)$. For the solution of equations (9), the necessary probability integrals are computed by the Monte Carlo method with a sample size equal to 5,000 (see, for example, [21,35]). The obtained values of the Lagrange multipliers are: $\lambda_7^- = 5.625$, $\lambda_7^0 = 0.965194702148438$, and $\lambda_7^+ = 2.34375$. The values of $TTIER$, $TTIER$, and $mdFDR$ are computed by the Monte Carlo simulation of random sequences with 5,000 observations. The codes for all necessary computations are written in MATLAB R2021b.

When using CBM, principle complexity does not arise when the problem dimension increases [21,22]. The computation time required to make a decision in CBM is equal to the time required for the classical Bayes method. The time required to determine the Lagrange multipliers certainly increases, but it is realized at the preparatory stage and does not directly affect the decision-making time.

4.1. Variances are known

Computational results, when variances are known, are given in Table 1 (null hypotheses are true) and Table 2 (alternative hypotheses are true) for different n_i and σ_i , $i = 1, 2, 3, 4$. From here, it is seen that for all computed combinations, $TTIER = mdFDR$ and they all are significantly less than the type I error rate of the Hayter PB method given in [4]. The powers for all computed combinations are equal to 1 and significantly surpass the appropriate values of the Hayter PB method. The offered method perfectly distinguishes one- and two-sided alternatives whose possibility is not discussed in ([4] p. 26, Table 2).

Table 1. Comparison of type-I errors at $k = 4$ (variances are known, null hypotheses are true).

σ	n	CBM at $q=0.05$		AN	α HPB Nominal $\alpha=0.05$	AH
		<i>TTIER</i>	<i>tmdFL</i>			
[1,2,2,3]	[12,12,12,12]	0.0190	0.0190	12.04 at H_0 11.89 at H_0 11.93 at H_0	0.0508	$H_0(5)$
[2,2,2,2]	[12,12,12,12]	0.0082	0.0082	11.93 at H_0 12.07 at H_0 11.96 at H_0	0.0528	$H_0(5)$
[1,2,2,3]	[6,8,10,12]	0.0081	0.0081	11.88 at H_0 11.93 at H_0 12.00 at H_0	0.0444	$H_0(5)$
[2,2,2,2]	[6,8,10,12]	0.0136	0.0136	11.98 at H_0 11.87 at H_0 11.91 at H_0	0.0476	$H_0(5)$
[3,2,2,1]	[6,8,10,12]	0.0109	0.0109	12.10 at H_0 11.96 at H_0 11.95 at H_0	0.0484	$H_0(5)$
[1,2,2,3]	[6,6,6,6]	0.0082	0.0082	11.94 at H_0 11.97 at H_0 11.96 at H_0	0.0516	$H_0(5)$
[2,2,2,2]	[6,6,6,6]	0.0054	0.0054	12.07 at H_0 12.00 at H_0 11.97 at H_0	0.0484	$H_0(5)$
[1,2,2,3]	[6,3,3,6]	0.0135	0.0135	11.86 at H_0 11.94 at H_0 11.91 at H_0	0.0460	$H_0(5)$
[2,2,2,2]	[6,3,3,6]	0.0136	0.0136	11.96 at H_0 11.90 at H_0 12.07 at H_0	0.0428	$H_0(5)$
[1,2,2,3]	[3,3,3,3]	0.0109	0.0109	11.97 at H_0 12.02 at H_0 11.91 at H_0	0.0424	$H_0(5)$
[2,2,2,2]	[3,3,3,3]	0.0217	0.0217	11.92 at H_0 11.88 at H_0 11.99 at H_0	0.0344	$H_0(5)$
[1,2,2,3]	[2,3,2,3]	0.0082	0.0082	11.96 at H_0 12.01 at H_0 12.00 at H_0	0.0368	$H_0(5)$

Continued on next page

σ	n	CBM at $q=0.05$		AN	α HPB	AH
		<i>TTIER</i>	<i>tmdFL</i>		Nominal $\alpha=0.05$	
[2,2,2,2]	[2,3,2,3]	0.0136	0.0136	11.88 at H_0 12.00 at H_0 11.97 at H_0	0.0336	$H_0(5)$
[3,2,2,1]	[2,3,2,3]	0.0136	0.0136	11.89 at H_0 11.95 at H_0 11.98 at H_0	0.0424	$H_0(5)$

Note: AN: Average number of observations necessary for making a decision; α HPB: Type-I Error Rate for Hayter PB Test; $\alpha \equiv TTIER$: Type I error rate for CBM; AH: Accepted hypotheses; $H_0(5)$: basic hypothesis of (5) is accepted.

Table 2. Powers comparison table at $k = 4$ (variances are known, alternative hypotheses are true).

n	σ	μ	CBM at $q=0.05$				AN	Hayter PB Nomin al $\alpha=0.05$ Power	AH
			<i>TTIER</i>	Power	<i>tmdFL</i>	<i>TTIER</i>			
[6,8,10,12]	[2,2,2,2]	[0,0,0,2]	0.0107	1	0.0107	0	11.99 at H_0 11.88 at H_0 1.64 at H_+	0.6388	$H_a(5)$
		[0,0,0.7,1.3]	0.0052	1	0.0052	0	11.99 at H_0 4.955 at H_+ 5.4 at H_+	0.2976	$H_a(5)$
		[0,1,2,3]	0	1	0	0	3.92 at H_+ 3.41 at H_+ 3.01 at H_+	0.7700	$H_a(5)$
		[-2,0,0,0]	0.0188	1	0.0188	0	2.028 at H_+ 11.88 at H_0 11.91 at H_0	0.4864	$H_a(5)$
		[-1.3,-0.7,0,0]	0.0078	1	0.0078	0	7.35 at H_+ 4.95 at H_+ 11.92 at H_0	0.2600	$H_a(5)$
		[-3,-2,-1,0]	0	1	0	0	3.957 at H_+ 3.45 at H_+ 3.05 at H_+	0.7700	$H_a(5)$
	[1,2,2,3]	[0,0,0,2]	0.0107	1	0.0107	0	11.93 at H_0 11.98 at H_0 2.029 at H_+	0.8020	$H_a(5)$

Continued on next page

n	σ	μ	CBM at $q=0.05$				AN	Hayter PB Nominal $\alpha=0.05$ Power	AH
			$TTIER$	Power	$tmdFL$	$TTIHR$			
		[0,0,0.7,1.3]	0.0052	1	0.0052	0	11.92 at H_0 4.93 at H_+ 7.15 at H_+	0.3384	$H_a(5)$
		[0,1,2,3]	0	1	0	0	2.92 at H_+ 3.37 at H_+ 3.84 at H_+	0.7356	$H_a(5)$
		[-2,0,0,0]	0.0080	1	0.0080	0	1.63 at H_+ 11.99 at H_0 11.95 at H_0	0.2836	$H_a(5)$
		[-1.3,- 0.7,0,0]	0	1	0	0	4.98 at H_+ 5.10 at H_+ 12.03 at H_0	0.2080	$H_a(5)$
		[-3,-2,-1,0]	0	1	0	0	2.98 at H_+ 3.39 at H_+ 3.85 at H_+	0.7356	$H_a(5)$
[3,2,2,1]		[0,0,0,2]	0.0107	1	0.0107	0	11.94 at H_0 11.91 at H_0 1.40 at H_+	0.7574	$H_a(5)$
		[0,0,0.7,1.3]	0.0026	1	0.0026	0	12.07 at H_0 5.07 at H_+ 4.23 at H_+	0.2210	$H_a(5)$
		[0,1,2,3]	0	1	0	0	5.29 at H_+ 3.42 at H_+ 2.52 at H_+	0.6724	$H_a(5)$
		[-2,0,0,0]	0.0027	1	0.0027	0	2.53 at H_+ 12.04 at H_0 12.03 at H_0	0.1886	$H_a(5)$
		[-1.3,- 0.7,0,0]	0	1	0	0	11.23 at H_+ 5.05 at H_+ 11.96 at H_0	0.1108	$H_a(5)$
		[-3,-2,-1,0]	0	1	0	0	5.35 at H_+ 3.43 at H_+ 2.48 at H_+	0.6724	$H_a(5)$

Continued on next page

number of observations used to make a decision.

Table 3. Comparison of Type-I errors, $k = 4$ (unknown variances are estimated by maximum likelihood method, null hypotheses are true).

σ	n	CBM at $q=0.05$		AN	α HPB Nominal $\alpha=0.05$	AH
		<i>TTIER</i>	<i>tmdFL</i>			
[1,2,2,3]	[12,12,12,12]	0.0190	0.0190	11.90 at H_0 11.96 at H_0 11.93 at H_0	0.0508	$H_0(5)$
[2,2,2,2]	[12,12,12,12]	0.0055	0.0055	12.03 at H_0 12.03 at H_0 12 at H_0	0.0528	$H_0(5)$
[1,2,2,3]	[6,8,10,12]	0.0054	0.0054	11.99 at H_0 11.99 at H_0 11.99 at H_0	0.0444	$H_0(5)$
[2,2,2,2]	[6,8,10,12]	0.0163	0.0163	11.91 at H_0 11.90 at H_0 11.95 at H_0	0.0476	$H_0(5)$
[3,2,2,1]	[6,8,10,12]	0.0191	0.0191	12.01 at H_0 11.93 at H_0 11.95 at H_0	0.0484	$H_0(5)$
[1,2,2,3]	[6,6,6,6]	0.0163	0.0163	11.99 at H_0 11.87 at H_0 11.95 at H_0	0.0516	$H_0(5)$
[2,2,2,2]	[6,6,6,6]	0.0217	0.0217	11.79 at H_0 12.05 at H_0 11.89 at H_0	0.0484	$H_0(5)$
[1,2,2,3]	[6,3,3,6]	0.0082	0.0082	11.99 at H_0 12.03 at H_0 11.95 at H_0	0.0460	$H_0(5)$
[2,2,2,2]	[6,3,3,6]	0.0245	0.0245	12 at H_0 11.85 at H_0 11.88 at H_0	0.0428	$H_0(5)$
[1,2,2,3]	[3,3,3,3]	0.0190	0.0190	12.02 at H_0 11.85 at H_0 12.02 at H_0	0.0424	$H_0(5)$
[2,2,2,2]	[3,3,3,3]	0.0164	0.0164	11.94 at H_0 12.04 at H_0 11.98 at H_0	0.0344	$H_0(5)$

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σ	n	CBM at $q=0.05$		AN	α HPB	AH
		$TTIER$	$tmdFL$		Nominal $\alpha=0.05$	
[1,2,2,3]	[2,3,2,3]	0.0163	0.0163	11.93 at H_0 11.88 at H_0 11.89 at H_0	0.0368	$H_0(5)$
[2,2,2,2]	[2,3,2,3]	0.0163	0.0163	11.96 at H_0 11.95 at H_0 11.96 at H_0	0.0336	$H_0(5)$
[3,2,2,1]	[2,3,2,3]	0.0164	0.0164	11.94 at H_0 11.95 at H_0 11.97 at H_0	0.0424	$H_0(5)$

Note: AN: Average number of observations necessary for making a decision; α HPB: Type-I Error Rate for Hayter PB Test; $\alpha \equiv TTIER$: Type I error rate for CBM; AH: Accepted hypotheses; $H_0(5)$: basic hypothesis of (5) is accepted.

Table 4. Power comparison table, $k = 4$ (unknown variances are estimated by the maximum likelihood method, alternative hypotheses are true).

n	σ	μ	CBM at $q=0.05$				AN	HayterPB	AH
			$TTIER$	Power	Power	$TTIER$		Nominal $\alpha=0.05$ Power	
[6,8,10,12]	[2,2,2,2]	[0,0,0,2]	0.008	1	0.008	0	11.95 at H_0 11.91 at H_0 2.16 at H_+	0.6388	$H_a(5)$
		[0,0,0.7,1.3]	0.01	1	0.01	0	11.94 at H_0 7.79 at H_+ 7.70 at H_+	0.2976	$H_a(5)$
		[0,1,2,3]	0	1	0	0	4.28 at H_+ 2.89 at H_+ 2.13 at H_+	0.7700	$H_a(5)$
		[-2,0,0,0]	0.0054	1	0.0054	0	4.28 at H_+ 11.98 at H_0 11.96 at H_0	0.4864	$H_a(5)$
		[-1.3,-0.7,0,0]	0	0.9948	0	0.0052	6.10 at H_+ 6.01 at H_+ 11.94 at H_0	0.2600	$H_a(5)$
		[-3,-2,-1,0]	0	1	0	0	2.74 at H_+ 3.73 at H_+ 3.43 at H_+	0.7700	$H_a(5)$

Continued on next page

n	σ	μ	CBM at $q=0.05$				AN	HayterPB Nominal $\alpha=0.05$ Power	AH
			$TTIER$	Power	Power	$TTIER$			
	[1,2,2,3]	[0,0,0,2]	0.0054	1	0.0054	0	11.89 at H_0 11.97 at H_0 8.63 at H_+	0.8020	$H_a(5)$
		[0,0,0.7,1.3]	0.0105	1	0.0105	0	11.87 at H_0 5.98 at H_+ 8.78 at H_+	0.3384	$H_a(5)$
		[0,1,2,3]	0	1	0	0	1.93 at H_+ 2.55 at H_+ 2.92 at H_+	0.7356	$H_a(5)$
		[-2,0,0,0]	0.0054	1	0.0054	0	3.77 at H_+ 11.97 at H_0 12.01 at H_0	0.2836	$H_a(5)$
		[-1.3,-0.7,0,0]	0.0078	1	0.0078	0	4.91 at H_+ 3.69 at H_+ 11.95 at H_0	0.2080	$H_a(5)$
		[-3,-2,-1,0]	0	1	0	0	2.56 at H_+ 2.74 at H_+ 4.37 at H_+	0.7356	$H_a(5)$
	[3,2,2,1]	[0,0,0,2]	0	0.9919	0	0.0081	11.99 at H_0 12.05 at H_0 1.12 at H_+	0.7574	$H_a(5)$
		[0,0,0.7,1.3]	0.0026	1	0.0026	0	11.97 at H_0 5.16 at H_+ 3.06 at H_+	0.2210	$H_a(5)$
		[0,1,2,3]	0	1	0	0	4.70 at H_+ 2.17 at H_+ 2.39 at H_+	0.6724	$H_a(5)$
		[-2,0,0,0]	0.0081	1	0.0081	0	2.71 at H_+ 11.98 at H_0 11.97 at H_0	0.1886	$H_a(5)$
		[-1.3,-0.7,0,0]	0	1	0	0	9.09 at H_+ 7.84 at H_+ 12.01 at H_0	0.1108	$H_a(5)$
		[-3,-2,-1,0]	0	1	0	0	4.33 at H_+ 2.04 at H_+ 1.84 at H_+	0.6724	$H_a(5)$

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n	σ	μ	CBM at $q=0.05$				AN	HayterPB AH	
			$TTIER$	Power	Power	$TTIER$		Nominal $\alpha=0.05$	Power
[6,8,10,12]	[2,2,2,2]	[0,0,0,-2]	0.0054	1	0.0054	0	11.99 at H_0 12.067 at H_0 2.097 at H_-	-	$H_a(5)$
		[0,0,-0.7,1.3]	0.0027	0.9973	0.0027	0.0027	12.04 at H_0 11.38 at H_- 1.98 at H_+	-	$H_a(4)$
		[0,-1,2,3]	0	1	0	0	4.06 at H_- 1.12 at H_+ 2.49 at H_+	-	$H_a(4)$
		[-2,0,1,0]	0	1	0	0	1.96 at H_+ 4.47 at H_+ 5.65 at H_-	-	$H_a(4)$
		[-1.3, 0.7, 0,0]	0.0026	1	0	0.0026	2.29 at H_+ 8.74 at H_- 11.97 at H_0	-	$H_a(4)$
		[-3,2,-1,0]	0	1	0	0	1.003 at H_+ 1.90 at H_- 3.81 at H_+	-	$H_a(4)$

Note: AN: Average number of observations necessary for making a decision; α HPB: Type-I Error Rate for Hayter PB Test; $\alpha \equiv TTIER$: Type I error rate for CBM; AH: Accepted hypotheses; $H_a(5)$: Alternative hypothesis of (5) is accepted; $H_a(4)$: Alternative hypothesis of (6) is accepted.

4.2.2. Computational results obtained by Stein's method

Computational results obtained by the Stein's method when the uniform distributions are used as a priori distributions of the unknown variances, are given in Table 5 (null hypotheses are true) and Table 6 (alternative hypotheses are true). The obtained results are very reliable. The probabilities of incorrect decisions are equal to zero, which are provided by the big number of observations necessary for making decisions when alternative hypotheses are true.

Table 5. Comparison of type-I errors, $k = 4$ (Stein's method, null hypotheses are true).

σ	n	CBM at $q=0.05$		AN	α HPB Nominal $\alpha=0.05$	AH
		<i>TTIER</i>	<i>tmdFL</i>			
[1,2,2,3]	[12,12,12,12]	0	0	2 at H_0 2 at H_0 4.18 at H_0	0.0508	$H_0(5)$
[2,2,2,2]	[12,12,12,12]	0	0	2 at H_0 2 at H_0 1 at H_0	0.0528	$H_0(5)$
[1,2,2,3]	[6,8,10,12]	0	0	2 at H_0 1 at H_0 6.05 at H_0	0.0444	$H_0(5)$
[2,2,2,2]	[6,8,10,12]	0	0	1.03 at H_0 1.54 at H_0 1 at H_0	0.0476	$H_0(5)$
[3,2,2,1]	[6,8,10,12]	0	0	1 at H_0 2 at H_0 1 at H_0	0.0484	$H_0(5)$
[1,2,2,3]	[6,6,6,6]	0	0	2 at H_0 2 at H_0 2.25 at H_0	0.0516	$H_0(5)$
[2,2,2,2]	[6,6,6,6]	0	0	2 at H_0 2 at H_0 1 at H_0	0.0484	$H_0(5)$
[1,2,2,3]	[6,3,3,6]	0	0	1 at H_0 1 at H_0 1 at H_0	0.0460	$H_0(5)$
[2,2,2,2]	[6,3,3,6]	0	0	1 at H_0 1 at H_0 1 at H_0	0.0428	$H_0(5)$
[1,2,2,3]	[3,3,3,3]	0	0	1 at H_0 1 at H_0 2 at H_0	0.0424	$H_0(5)$
[2,2,2,2]	[3,3,3,3]	0	0	1 at H_0 1 at H_0 1 at H_0	0.0344	$H_0(5)$
[1,2,2,3]	[2,3,2,3]	0	0	1 at H_0 1 at H_0 13.25 at H_0	0.0368	$H_0(5)$

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σ	n	CBM at $q=0.05$		AN	α HPB Nominal $\alpha=0.05$	AH
		$TTIER$	$tmdFL$			
[2,2,2,2]	[2,3,2,3]	0	0	1 at H_0 1 at H_0 1 at H_0	0.0336	$H_0(5)$
[3,2,2,1]	[2,3,2,3]	0	0	1 at H_0 1 at H_0 13.81 at H_0	0.0424	$H_0(5)$

Note: AN: Average number of observations necessary for making a decision; α HPB: Type-I Error Rate for Hayter PB Test; $\alpha \equiv TTIER$: Type I error rate for CBM; AH: Accepted hypotheses; $H_0(5)$: basic hypothesis of (5) is accepted.

Table 6. Power comparison table, $k = 4$ (Stein's method, alternative hypotheses are true).

n	σ	μ	CBM at $q=0.05$			AN	HayterPB Nominal $\alpha=0.05$ Power	AH
			Power	$TTIER$, $tmdFL$	$TTIER$			
[6,8,10,12]	[2,2,2,2]	[0,0,0,2]	1	0	0	390.42 at H_0 389.50 at H_0 169.46 at H_+	0.6388	$H_a(5)$
		[0,0,0.7,1.3]	1	0	0	390 at H_0 621.5 at H_+ 3001 at H_+	0.2976	$H_a(5)$
		[0,1,2,3]	1	0	0	350.90 at H_+ 242.40 at H_+ 480.70 at H_+	0.7700	$H_a(5)$
		[-2,0,0,0]	1	0	0	79.62 at H_+ 390 at H_0 121.54 at H_0	0.4864	$H_a(5)$
		[-1.3,-0.7,0,0]	1	0	0	137.50 at H_+ 65.13 at H_+ 650.86 at H_0	0.2600	$H_a(5)$
		[-3,-2,-1,0]	1	0	0	1085.50 at H_+ 146.60 at H_+ 1014.75 at H_+	0.7700	$H_a(5)$

Continued on next page

n	σ	μ	CBM at $q=0.05$			AN	HayterPB Nominal $\alpha=0.05$ Power	AH
			Power	$TTIER,$ $tmdFL$	$TTIIR$			
	[1,2,2,3]	[0,0,0,2]	1	0	0	390.33 at H_0 390.42 at H_0 100.38 at H_+	0.8020	$H_a(5)$
		[0,0,0.7,1.3]	1	0	0	390.25 at H_0 1074.33 at H_+ 1500.67 at H_+	0.3384	$H_a(5)$
		[0,1,2,3]	1	0	0	140.80 at H_+ 205.80 at H_+ 522.33 at H_+	0.7356	$H_a(5)$
		[-2,0,0,0]	1	0	0	27.44 at H_+ 390.22 at H_+ 554.44 at H_0	0.2836	$H_a(5)$
		[-1.3,-0.7,0,0]	1	0	0	353.50 at H_+ 3563 at H_+ 2156 at H_0	0.2080	$H_a(5)$
		[-3,-2,-1,0]	1	0	0	573.50 at H_+ 297.25 at H_+ 580 at H_+	0.7356	$H_a(5)$
	[3,2,2,1]	[0,0,0,2]	1	0	0	390.58 at H_0 389.83 at H_0 122.69 at H_+	0.7574	$H_a(5)$
		[0,0,0.7,1.3]	1	0	0	391.25 at H_0 275.69 at H_+ 359.75 at H_+	0.2210	$H_a(5)$
		[0,1,2,3]	1	0	0	2070.50 at H_+ 671.33 at H_+ 600.67 at H_+	0.6724	$H_a(5)$
		[-2,0,0,0]	1	0	0	74.15 at H_+ 390.33 at H_0 75.38 at H_0	0.1886	$H_a(5)$
		[-1.3,-0.7,0,0]	1	0	0	1567.67 at H_+ 174.25 at H_+ 8.50 at H_0	0.1108	$H_a(5)$

Continued on next page

n	σ	μ	CBM at $q=0.05$			AN	HayterPB Nominal $\alpha=0.05$ Power	AH
			Power	$TTIER$, $tmdFL$	$TTIER$			
		[-3,-2,-1,0]	1	0	0	3582 at H_+ 790.50 at H_+ 1384 at H_+	0.6724	$H_a(5)$
[6,8,10,12]	[2,2,2,2]	[0,0,0,-2]	1	0	0	390.5 at H_0 391.67 at H_0 127.92 at H_-	-	$H_a(4)$
[6,8,10,12]	[2,2,2,2]	[0,0,-0.7,1.3]	1	0	0	389.2 at H_0 959.4 at H_- 76.17 at H_+	-	$H_a(4)$
		[0,-1,2,3]	1	0	0	2801 at H_- 26.5 at H_+ 287.5 at H_+	-	$H_a(4)$
		[-2,0,1,0]	1	0	0	46 at H_+ 72.33 at H_+ 868.2 at H_-	-	$H_a(4)$
		[-1.3, 0.7, 0,0]	1	0	0	213 at H_+ 4592 at H_- 65 at H_0	-	$H_a(4)$
		[-3,2,-1,0]	1	0	0	32 at H_+ 318.27 at H_- 452 at H_0	-	$H_a(4)$

Note: AN: Average number of observations necessary for making a decision; α HPB: Type-I Error Rate for Hayter PB Test; $\alpha \equiv TTIER$: Type I error rate for CBM; AH: Accepted hypotheses; $H_a(5)$: Alternative hypothesis of (5) is accepted; $H_a(4)$: Alternative hypothesis of (6) is accepted.

5. Discussion

The offered method of solving the ANOVA problem, for known and unknown variances of the results, with restricted Type I and Type II error rates based on CBM gives very reliable results that surpass the methods of today. It is a sequential method that requires a set of observations to make a decision, the number of which is especially large in Stein's method in the case of true alternative hypotheses. Nevertheless, due to the large practical application of the ANOVA problem and the fact that, in many cases, the number of data is quite large, and the requirements for the accuracy of the decisions are high, we consider the results to be important and to deserve high attention.

It is worth noting that the CBM method using the maximum likelihood method requires more observations than the same CBM method using Stein's method to test the null hypothesis; this situation is reversed when testing alternative hypotheses. In this case, using Stein's method requires more

observations than using the maximum likelihood method. The reason lies in the use of different values of ω_0 . It should be noted that the increase of ω_0 causes the increase in the quantity of observations necessary to make a decision and, accordingly, the increase of the accuracy of made decisions [24,33].

The computing results obtained by CBM and the Hayter PB methods are given in Tables 1–6. In order to compare the results obtained by these methods, $q = 0.05$ is taken in CBM, since in Hayter PB methods, the calculations have been realized for a Type I error rate of $\alpha = 0.05$ (see Tables 2–4, 6 in [4]). Since each scenario has four normal population correspondences, i.e., $k = 4$, we have to test three hypotheses each time (see (6)). In each case under consideration, the true hypotheses are determined by the values of $\theta_i = \mu_{i+1} - \mu_i$, $i = 1, \dots, k - 1$. Therefore, in each scenario, three hypotheses (from the set of hypotheses H_0 , H_- and H_+) are accepted, and depending on θ_i , $i = 1, \dots, k - 1$, the values of the criteria $TTIER$, $TIIER$, $tmdFDR$, and $Power = 1 - TIIER$ are computed. The values criteria $TTIER$ and $tmdFDR$ are given when null hypotheses are true in all three cases (see Tables 1, 3, and 5), and the values all four considered criteria $TTIER$, $TIIER$, $tmdFDR$, and $Power = 1 - TIIER$ are given when the alternative hypothesis is true even for one of the three cases. In the columns n , σ , and μ , the values of sample sizes for considered normal populations, standard deviations, and mathematical expectations, respectively, are given for each considered case. The second to last the final column contains the results obtained by Hayter PB methods from [4], and the last column contains each scenario in the CBM accepted hypotheses defined by (5) or (6). The column AN (average number) contains the average number of observations necessary to make a decision in CBM (averaged over the results of 5,000 experiments).

From the results of these Tables, the superiority of CBM over Hayter PB methods is evident. Moreover, unlike the Hayter PB method, CBM enables us to distinguish (4) and (5), which are the same as (1) and (2).

6. Conclusions

The problem of testing hypotheses concerning the means of a set of normally distributed random variables at the restriction of both Type I and Type II error rates are considered in this paper. To solve the stated problem, one of the possible statements of CBM is used. In order to overcome the difficulties related to the complexity of testing hypotheses caused by not knowing the values of the variances of random variables, the maximum likelihood and Stein's methods are used. The ability to limit Type I and Type II error rates and other relevant criteria to the required level is shown for the developed algorithms. The latter is confirmed by the calculation results of practical examples by simulating different scenarios.

Author contributions

Kartlos J. Kachiashvili: Conceptualization, Formal analysis, Investigation, Methodology, Project administration, Resources, Software, Supervision, Validation, Visualization, Writing—original draft, Writing—review & editing; Ashis SenGupta: Conceptualization, Formal analysis, Investigation, Resources, Validation, Writing—original draft, Writing—review & editing; Joseph K. Kachiashvili: Data curation, Formal analysis, Investigation, Resources, Software, Visualization, Writing—original draft, Writing—review & editing.

Use of Generative-AI tools declaration

The authors declare they have not used Artificial Intelligence (AI) tools in the creation of this article.

Conflict of interest

All Authors declare that they have no financial interests.

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