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Optimization of multi-response problems with continuous functional responses by considering dispersion effects

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KEYWORDS Multiple responses optimization; Functional responses; Design of experiments; Polynomial integral. Abstract. In some processes, quality of a product should be characterized by functional relationships between response variables and a signal factor. Hence, the traditional methods cannot be used to find the optimum solution. In this paper, we propose a method, which considers two different dispersion effects in domain and between replicates variations in the functional responses. Besides, we propose an integral based measure to find the deviation from target function. A probabilistic method is applied to consider the correlation structure of functional responses. Three numerical examples and a real case from literature are studied to show the efficiency of the proposed method.

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1. Introduction

Design of experiments is one of the offline quality engineering methods to find level settings for controllable factors in order to find optimum responses. In many real production processes, there are more than one response which should be optimized. The problem will be more difficult if optimization direction of responses Many researchers tried to solve this is conflicting. problem using aggregated measures. One of the most popular measures is the desirability function. This measure was proposed by Harrington [1] at first to transform different objective values into a free scaled value between 0 and 1. He also proposed using the geometric mean of obtained desirabilities as total desirability. Thereafter, Derringer and Suich [2] proposed different desirability functions for Smaller-The-Better

*. Corresponding author. Tel.: +98 21 51212092; Fax: +98 21 51212020 E-mail address: bashiri@shahed.ac.ir (M. Bashiri). (STB), Larger-The-Better (LTB), and Nominal-The-Best (NTB) responses. There are several examples of using the desirability function in optimization problems. For example, Del Castillo et al. [3] employed modified desirability functions to optimize a wire bounding process. Jeong and Kim [4] proposed an interactive method based on the desirability function to optimize an aggregated model. Ribardo and Allen [5] incorporated mean, standard deviation, and possible shifts in the process mean in the proposed desirability function. Noorossana et al. [6] used an artificial neural network to build a prediction model and then employed a genetic algorithm to optimize a desirability function. Mostafa et al. [7] tried to minimize a desirability function, which aggregated shrinkage and warpage on the fuel filter in the injection molding process using simulated annealing. He et al. [8] proposed a desirability function by considering the worst response in the confidence region. Chen et al. [9] selected some proper weights to balance location with dispersion desirability values. Costa et al. [10] evaluated performance of some desirability-based approaches and categorized them into two less and more sophisticated approaches.

Based on the literature, we can conclude that the desirability function is used as an efficient measure in multi-response optimization problems.

In the presence of signal factor(s), we deal with a problem including functional response(s). There are different approaches in the literature that consider the signal factor in the problem. For example, Taguchi [11] proposed dynamic signal to noise ratio for such prob-Miller and Wu [12] divided signal-response lems. problems into two main categories of measurement systems and multiple target systems. They showed that Taguchi's method was appropriate for certain measurement systems, but not for multiple target systems. They proposed performance measure modeling as well as response function modeling for signalresponse problems. Tong et al. [13] applied a fuzzy TOPSIS (Technique for Order Preference by Similarity to Ideal Solution) method to solve dynamic multiresponse problems. Ozdemir and Maghsoodloo [14] developed a sensitivity measure for signal-response problems with three correlated responses. Tong et al. [15] used Data Envelopment Analysis (DEA) to analyze the relations between signal factor and responses. Fogliatto [16] proposed a desirability based method for optimizing problems with functional responses. To do this, he used the Hausdorff Distance (HD) to calculate the deviation between response value and the target value in each time point. Goethals and Cho [17] considered time as signal factor and tried to optimize the problem in certain periods of time. Maghsoodloo and Huang [18] extended quality loss function and signal to noise ratio for a problem with different types of correlated responses. Wu [19] used double-exponential desirability function to optimize multiple nonlinear dynamic quality characteristics. Tong et al. [20] used desirability function to propose an optimization procedure using dual-response-surface method. Su et al. [21] applied Artificial Neural Networks (ANN) to model the relation of controllable factors and responses, and used Scatter Search (SS) to find best setting in a problem with dynamic characteristics. Chang [22] also used ANN to build the dynamic response model and proposed modified desirability functions for optimizing phase. Chang [23] proposed a data mining approach, which used artificial neural networks, desirability functions, and simulated annealing algorithm to optimize problems with multiple dynamic responses. Chang and Chen [24] used a genetic algorithm to optimize a dynamic multi-response model, which was created using ANN. Storm et al. [25] extended classical response surface methodology for modeling of time series response data. Zhang et al. [26] proposed a goal programming approach to model multi-response optimization problems and used particle swarm optimization to search for the optimal global solution. Gauri [27] proposed a Principal Component Analysis (PCA)-based approach to optimize a multi-response problem with linear functional responses by taking into account the correlation structure of the responses. Cui and He [28] used Kriging model to establish a functional response model and applied a genetic algorithm to reach the optimum global solution. Table 1 shows a comparison between previous studies of signal-response optimization methods and the current study.

In this paper, we propose a method to optimize problems with multiple functional responses by considering two different dispersion effects. The first effect is related to variability of a functional response through the signal domain, which is named "In Domain Variation (IDV)," and the second one, which is named "Between Replicate Variation (BRV)," is related to variability among functional responses corresponding to different replicates of a treatment. Many of the signal factors, which were considered in the previous works, have a continuous nature (for example, time, pressure, etc.). However, all previous works have a discrete viewpoint to the signal factors in the calculation steps. Although increasing number of signal levels for sampling may lead to an approximately continuous viewpoint, it increases the sampling cost as well and may be economically unfeasible, especially in the destructive experiments. Moreover, in some cases, the target function has a predefined equation. In these situations, we may want to know which factor's setting leads to a function with the lowest deviation from target in the whole signal's domain and what the resulting function is, by selecting this setting. However, in the discrete approaches, we may find the setting which leads to minimum deviations by considering the selected signal levels. For better comprehension, consider the semiconductor manufacturing problem from Kang and Albin [29]. A critical device in this process is Mass Flow Controller (MFC), which controls the flow of gases in the chamber. They showed that if an MFC was in-control, the pressure in the chamber would behave as a linear function of gases flow. Thus, if we design an experiment to optimize this process, we may search for a setting which leads to a linear function with minimum deviation from the target function. Besides, in some, cases optimization may need to be done on some historical data when response values have been collected over different signal levels for different treatments and target values are available for different levels. In such cases, discrete viewpoint is not applicable because the response values of treatments cannot be compared. Another issue comes when multiple functional responses are correlative. In this situation, considering responses to be independent ignores their correlational structure. Moreover, finding an optimal setting, which is not among the experiments done, may be another issue in the mind of the analyst. In this paper, we propose

Author(s)	Year	Correlation between responses	Interaction between factors	Responses with different types	Location effect	Between replicates variation	orrelation between nctional responses	In domain variation	Continuous unctional response
Tong et al [13]	1997						F C		
Maghsoodloo and Huang [18]	2001		v	V	V				
Tong et al. [20]	2002	v	\checkmark						
Ozdemir and Maghsoodloo [14]	2004	\checkmark							
Su et al. [21]	2005		\checkmark	\checkmark	\checkmark	\checkmark			
Chang [22]	2006		\checkmark	\checkmark	\checkmark				
Chang [23]	2008		\checkmark	\checkmark	\checkmark	\checkmark			
Fogliatto [16]	2008		\checkmark	\checkmark	\checkmark				
Tong et al. [15]	2008		\checkmark	\checkmark	\checkmark	\checkmark			
Wu [19]	2009	\checkmark		\checkmark	\checkmark		\checkmark		
Chang and Chen [24]	2011		\checkmark	\checkmark	\checkmark				
Goethals and Cho [17]	2011		\checkmark	\checkmark	\checkmark	\checkmark			
Storm et al. [25]	2013		\checkmark	\checkmark	\checkmark				
Zhang et al. [26]	2014			\checkmark	\checkmark	\checkmark			
Gauri [27]	2014	\checkmark		\checkmark	\checkmark	\checkmark	\checkmark		
Cui and He [28]	2016		\checkmark		\checkmark				\checkmark
Proposed approach	-	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark

Table 1. Comparison of the previous signal-response studies with the proposed approach.

an approach to optimize multi-response problems with functional continuous response(s) by considering two types of dispersion effect. To find the best factor-level setting, a genetic algorithm is employed to search in the solution space.

The structure of the paper is as follows. In the next section, the problem statement is given. Section 3 includes the proposed methodology and the genetic algorithm applied for solving the optimization model. Three numerical examples are presented in Section 4. A real case from literature is studied in Section 5 to illustrate the application of the proposed method. Our concluding remarks as well as some future researches are given in the final section.

2. Problem statement

We deal with an experimental design with an $m \times 1$ vector \mathbf{f}_{kl} containing m continuous functional responses $(f_{1kl}, f_{2kl}, ..., f_{mkl})^T$ in replicate l of treatment k, with covariance matrix Σ_{kl} for residuals vector \mathbf{e}_{kl} ; and an $n \times 1$ vector \mathbf{y}_{kl} containing n classic responses $(y_{1kl}, y_{2kl}, ..., y_{nkl})^T$ in the form of STB, NTB, or LTB in replication l of treatment k with covariance matrix \mathbf{S}_{kl} for corresponding residuals vector $\boldsymbol{\varphi}_{kl}$. In the case of STB responses, the smallest result is desired while in the case of LTB responses, the largest result is desired. In the case of NTB responses, the result with minimum deviation from a predefined target is desired. There are q controllable factors $(x_1, x_2, ..., x_q)$, which have effect on both classic and functional type responses. The relation between classic responses and controllable factors is defined as:

$$\mathbf{y}_{kl} = \mathbf{A}\mathbf{x}_k + \boldsymbol{\varphi}_{kl},\tag{1}$$

where **A** is the coefficient matrix and \mathbf{x}_k is in the form of $(1, x_{k1}, x_{2k}, \dots, x_{qk})^T$. Moreover, there is a signal factor t, which has effect on functional responses. To address the curvature in the functional responses, we suppose that there is a polynomial relation between signal factor and functional response(s). In other words, there is a polynomial target function for each functional response. The functional responses relation can be defined as follows:

$$\mathbf{f}_{kl} = \mathbf{\Gamma} \mathbf{x}_k + \mathbf{B}_k \mathbf{t} + \mathbf{e}_{kl},\tag{2}$$

where $\mathbf{\Gamma}$ is the coefficient matrix of controllable factors, \mathbf{t} is $(1, t, t^2, ..., t^p)^T$, and \mathbf{B}_k is the matrix of polynomial coefficients in the kth treatment. It is assumed that functional responses are independent from classic ones. Moreover, the effects of possible nuisance factors are randomly distributed and, hence, are considered through the covariance matrix of the residuals. It is assumed that all responses, including functional ones, are normally distributed. However, in the case of other distributions, a transformation method may be used to satisfy the normality assumption. The goal is finding the best levels of controllable factor such that classic responses are optimized and functional response(s) have minimum deviation from their corresponding target functions.

3. Proposed methodology

The proposed method involves four main steps including regression estimation, new index calculation, aggregation, and optimization to find the best levels for the factors. The continuous nature of the signal factor is considered in the second step, while the correlation structure of functional responses is considered in the third step.

Step 1: Find the best regression function. Firstly, a regression function should be fitted to the obtained data in each treatment and should be compared with the target function. Since the number of signal factors in this paper is considered equal to one and we are going to address the curvature in the regression function, we use polynomial regression. Determining the number of discrete levels of the signal factor for sampling depends on the case. Increasing the number of discrete levels may lead to better estimation of functions, but it may also increase the cost of sampling. Hence, if we are not worried about sampling cost, we can increase the number of levels as much as possible. On the other hand, in the cases with considerable sampling cost, a bi-objective model can be defined by considering parameters estimation error and sampling costs as objective functions to facilitate finding of the optimum number of discrete levels. To find the best order of the polynomial regression, we test different orders and select the one that minimizes the Sum of Squared Errors (SSE) of the fitted model. In the case of testing different orders to avoid overfitting, we can limit the maximum order of estimated functions based on the dispersion pattern and number of data points. An overfitted function usually has several extremum (maximum or minimum) points near the data points. Hence, a function with few extremum points and small SSE can be considered as a good fit. Another alternative is to split the data into training and testing sets. However, selecting data for the test set may be another issue. A good choice to overcome this problem is using the k-fold

cross-validation method [30]. In this method, data is partitioned into k sets of equal size. Then, in each iteration, k-1 data sets are used to estimate the polynomial regression and one set is used as test set to find the SSE. This process is repeated k times with different test sets and the average of all SSEs is used to evaluate the accuracy and precision of the estimation. It should be noted that the number of data should be large enough to use this method.

Step 2: Calculating a new index for functional responses. Consider the example in Figure 1 is comprised of an NTB functional response, f, and its target function. If the researcher only considers the four discrete levels of the signal factor, t, shown in the figure, there is a negligible difference between estimated and target functions. However, it is obvious that the estimated function deviates from its target. As a proposed solution, the area enclosed between the two functions is considered as a deviation measure. Therefore, for the NTB type functional responses, cross points of fitted and target functions should be found. To do this, MATLAB software can be used to find roots of the obtained equation by subtracting the fitted function from the target one. The next step is aggregating the absolute values of integrals between roots as deviation measure. If we are only interested in the location effect and there are not any additional response variables, the treatment with the minimum Aggregated Absolute Integral Value (AAIV) can be selected as the best. By considering one functional response in the problem, the AAIV for the *l*th replicate of the kth treatment may be calculated as follows:

$$AAIV_{hkl} = \sum_{i,j} \left| \int_{i}^{j} (T_h(t) - \hat{f}_{hkl}(t)) dt \right|,$$
(3)

where i, j are two consecutive cross points of the target and estimated functions, $T_h(t)$ is target function of the *h*th functional response, and $\hat{f}_{hkl}(t)$ is estimated function of the *h*th response for the *l*th replicate in the *k*th treatment. Note that in the problems



Figure 1. A sample of estimated functional response with target function.



Figure 2. Two different functions with equal AAIVs but different shapes.

with more than one functional response, AAIV is calculated for each response separately and, then, overall index is calculated as expressed in the next step.

There are two different dispersion effects in the considered problem. The first one is IDV, which is related to the difference in deviation of estimated function from target in the signals domain. Τo consider it, the maximum difference between target and estimated functions obtained by considering all replicates may be added in the new index. This can help to distinguish sudden changes of deviation between estimated and target functions. For better illustration, two estimated functions with equal AAIV values have been depicted in Figure 2. The estimated function for treatment 1 is very close to the target function in most of its signal levels, but there is a large deviation for some levels, so it is clear that the proposed index should consider such deviations. In addition, the deviation between the target and estimated functions for treatment 2 is almost constant throughout the different signal levels, which may be preferable since both treatments have equal AAIV values. Furthermore, in some cases, positive or negative deviations may have various importance and we modify the proposed index to a Weighted Aggregated Absolute Integral Value (WAAIV) as follows:

$$WAAIV_{hkl} = w_1 \sum_{i,j} \left(\int_i^j \left(T_h(t) - \hat{f}_{hkl}(t) \right) dM \right) + w_2 \sum_{p,q} \left(\int_p^q \left(\hat{f}_{hkl}(t) - T_h(t) \right) dM \right) + w_3 \max_t \left| T_h(t) - \hat{f}_{hkl}(t) \right|,$$
(4)

where *i* and *j* are two consecutive cross points where target function of the *h*th functional response $T_h(t)$ is above the estimated function of the *h*th functional response for the *l*th replicate in *k*th treatment $\hat{f}_{hkl}(t)$



Figure 3. Three estimated functions for different replicates of a treatment.

and p, q, are two consecutive cross points where the mentioned target function is below the mentioned estimated function; and w_1 , w_2 , and w_3 are weight coefficients for the negative, positive, and in domain variations, respectively.

The second type of dispersion effect is BRV and may be considered when there are different estimated functions for each treatment because of replicates. For better comprehension, consider Figure 3. This type of dispersion effect is incorporated in the next step of the proposed method.

Some of the problems may have STB or LTBtype functional responses. For example, consider the air pollution rate through the time. It is desired that this value should be near zero throughout the whole signal domain. Thus, it may be considered as an STB functional response. In such cases, the AAIV index may be simply obtained by calculating the area under the fitted curve in each treatment. Note that we aim to minimize the AAIV for STB or NTB-type functional responses while it is desired to maximize it for LTB-type functional responses.

Step 3: Constructing the overall index for multiple responses. In this step, all calculated indices for different responses should be summarized in a total index. However, there are two possible situations here. In the first case, responses are uncorrelated and in the second one, they have a significant correlation structure. In the first case, the desirability measure is employed. To calculate the desirability of STB, LTB, and NTB-type classic responses, the proposed indices by Derringer and Suich [2] are used. In addition, the desirability for an NTB or STB functional response may be calculated based on Derringer and Suich [2] as follows:

$$dm_{h,k} = \begin{cases} \left| \frac{UM_h - M_{h,k}}{UM_h} \right|^{\alpha} & 0 < M_{h,k} < UM_h \\ 0 & M_{h,k} > UM_h \end{cases} , \quad (5)$$

where $M_{h,k}$ is the average of AAIV (or WAAIV)

values of the *h*th functional response in the *k*th treatment, UM_h is the upper specification limit for $M_{h,k}$, and α is the shape constant of the desirability function.

In the case of problems with two or more replicates in each treatment, we also have variance for AAIV (or WAAIV) values in each treatment. Hence, to incorporate the BRV as the second type of dispersion effect in the problems with uncorrelated responses, we consider the variance of the AAIV (or WAAIV) for each functional response as an STB-type response, and then calculate its desirability as follows.

$$d\nu_{h,k} = \begin{cases} \left| \frac{UV_h - V_{h,k}}{UV_h} \right|^{\alpha} & 0 < V_{h,k} < UV_h \\ 0 & V_{h,k} > UV_h \end{cases},$$
(6)

where $V_{h,k}$ is the variance of AAIV (or WAAIV) values of the *h*th functional response in the *k*th treatment, UV_h is the upper specification limit for $V_{h,k}$, and α is the shape constant of the desirability function.

To calculate the overall desirability, D, at each treatment, Derringer [31] proposed a modified geometric average with weighting terms, which is employed in this paper as well.

In the second case, there are correlated responses and thus, the previous approach is not recommended because it ignores the correlation structure of responses. To consider multiple correlated responses, Chiao and Hamada [32] proposed a method in which the treatment with maximum probability of being all responses in the corresponding specification limits is selected as the best solution. The probability is calculated by considering multivariate normal distribution, which considers the covariance matrix of responses. Hence, the correlation structure is considered as well as the BRV in the calculations. The Proportion Of Conformance (POC) for correlated responses in the kth treatment can be calculated as follows:

$$POC_k = P(\mathbf{y} \in \mathbf{E} \,|\, \mathbf{x}_k),\tag{7}$$

where **y** is the vector of classic responses and **E** is the corresponding specification region. For better comprehension, consider two correlated y_1 , y_2 responses with specification region and 95% confidence interval as shown in Figure 4. The shared area, which is illustrated with dashes, shows the proportion of conformance.

On the other hand, when the functional responses correlate, we should compute the covariance matrix of AAIVs (or WAAIVs) through the replicated data. It should be noted that before calculating the POC index for AAIVs (or WAAIVs), we should check the normality assumption for them. If the AAIVs (or WAAIVs) do not follow a normal



Figure 4. Proportions of conformance for two supposed responses.

distribution, we should use a transformation method. Then, the Proportion Of Conformance (POC) for each treatment can be calculated by replacing \boldsymbol{y} with **AAIV** (or **WAAIV**) in Eq. (7). Note that the specification region for **AAIV** (or **WAAIV**) is in the form of $\begin{bmatrix} 0 & b_h \end{bmatrix}^m$, where b_h is the desired upper bound for AAIV_{hkl} (or WAAIV_{hkl}) defined by the analyst.

Step 4: Optimizing the overall index to find the best setting. By doing the previous steps, the analyzer can select the best treatment based on the overall index. However, if the analyzer wants to search for a better setting in the whole solution space, they may run this step. Before running the optimization algorithm, a relation function between controllable factors and calculated indices (POC or total desirability) should be estimated. The final step is employing an algorithm to find the best settings for the controllable Genetic Algorithm (GA) is one of the factors. popular methods to search for near-optimal solution in the problems with discrete or continuous solution spaces. In each iteration, it makes better solutions using crossover and mutation operators. There are several examples of using GA in multi-response optimization problems. For example, refer to [6,33-36]. However, it should be noted that GA is not the only alternative and the optimization algorithm may be selected by the analyst based on their preferences and the problem conditions. Thus, we recommend it to search for the optimum total index. It should be noted that tuning of GA parameters is performed using the Taguchi method. In this method, through a factorial design, the mean of calculated responses in each try for different parameter levels is computed and the best level for each parameter is selected.

4. Numerical examples

4.1. Example 1: Performance evaluation

As the first example, suppose a 2^2 factorial design with a signal factor t limited by [-5, 5] and a functional



Figure 5. Target function versus fitted functions for different treatments in Example 1.

 Table 2. Experimental design of Example 1

Treatment	x_1	x_2	t=-5	t = -3	t = -1	t = 1	t=3	t=5	SSE
1	1	1	-3168.01	-305.97	-42.02	-342.23	-202.08	2711.04	4.56×10^{-25}
2	1	-1	-2927.91	-216.87	-52.45	-22.69	57.38	2871.99	4.31×10^{-25}
3	-1	1	-2527.56	234.19	109.01	197.67	488.40	2982.20	7.51×10^{-25}
4	-1	-1	186.15	-162.13	4.32	2.83	-2826.86	2876.91	7.76×10^{-23}

response y. It is supposed that the target function for y is $t^5-2t^3+t^2-4t+1.5$. To evaluate the performance of the proposed method, suppose that we know the exact functional response corresponding to each treatment as shown in Eqs. (8)-(11), respectively:

$$0.6149 t^{5} + 0.3904 t^{4} + 14.7584 t^{3} - 11.6588 t^{2} - 165.2982t - 181.4266, \qquad (8)$$

 $1.2314 t^5 + 0.3522 t^4 - 8.4762 t^3$

 $-8.7811 t^2 - 22.2397 t - 29.3061, (9)$

 $1.3374 t^5 - 1.4330 t^4 + 13.6787 t^3$

 $+ 40.3379 t^2 - 57.1314 t - 113.9450, \quad (10)$

$$4.1653 t^{5} + 15.6840 t^{4} - 97.0485 t^{3} - 344.1352 t^{2} - 91.9832 t - 331.6912.$$
(11)

Figure 5 shows the target and treatment functions. For the first evaluation, we consider (-5, -3, -1, 1, 3, 5) as initial signal levels to generate response values and add a standard normal residual to them. The resulting response values under different signal levels are reported in Table 2.

To evaluate the efficiency of the proposed measure, we compare it with HD proposed by Fogliatto [16]. The HD between two vectors \mathbf{y}_1 and \mathbf{y}_2 is calculated as follows:

$$h(\mathbf{y}_{1}, \mathbf{y}_{2}) = \max\left\{ \max_{y_{1} \in \mathbf{y}_{1}} \left(\min_{y_{2} \in \mathbf{y}_{2}} \sqrt{(y_{1} - y_{2})^{2}} \right), \\ \max_{y_{2} \in \mathbf{y}_{2}} \left(\min_{y_{1} \in \mathbf{y}_{1}} \sqrt{(y_{1} - y_{2})^{2}} \right) \right\},$$
(12)

As can be seen in Eq. (12), HD uses two points, which have minimum deviation from each other, to calculate the distance between the two vectors and ignores the order of points. For example, by considering $y_1 = (2, 10, 5)$ and $y_2 = (10, 5, 2)$, the result is $h(\mathbf{y}_1, \mathbf{y}_2) = 0$; however, it is obvious that they are different. To calculate the AAIV for each treatment, we estimate a function for it based on the results reported in Table 2. The goodness of fitness is evaluated through the sum of squared differences between each observation and its estimated value (SSE), which is reported in Table 2 as well. Table 3 shows the AAIV and HD for each treatment. It can be seen that using the HD (based on [16]) as comparing measure leads to selecting treatment 4 as the best one. However, it is obvious from Figure 5 that it is not the best choice. This shows that the HD is not an efficient measure in such problems. Another conflict of results can be seen between treatments 1 and 3. Based on the AAIV measure, treatment 1 dominates 3, but based on the HD measure, the result is the opposite. This is another problem with employing HD and occurs due

Table 3. AAIV and HD results for each treatment in Example 1 by considering [-5, 5] as signal domain.

Treatment	x_1	x_2	AAIV	HD
1	1	1	2389.98	339.51
2	1	-1	681.02	130.11
3	-1	1	2615.77	300.93
4	-1	-1	12180.79	5.32



Figure 6. AAIV changes due to change in signal levels.



Figure 7. HD changes due to change in signal levels.

to its discrete nature. To compare the variations of AAIV and HD results, we change each signal level in a specified interval [-0.5, 0.5]. Hence, the signal levels is changed from (-5.5, -3.5, -1.5, 0.5, 2.5, 4.5) to (-4.5, -2.5, -0.5, 1.5, 3.5, 5.5) in 10 iterations. As shown in Figure 6, by changing the signal levels, AAIVs are approximately robust against the added values to the initial signal levels; however, the changes in HD according to the signal level shifts are considerable as shown in Figure 7. This shows that HD is related to signal levels and the best treatment may be changed by selecting different signal levels, but AAIV is more robust. Nonetheless, HD may be a good choice in the cases that we deal with monotonic functions for the treatments. In such problems, response values in different signal levels may be far from each other and, hence, ignoring the order due to the use of HD cannot affect the selected solution. For better illustration, consider [4, 5] as limit of signal domain. As can be seen in Figure 5, all of the functions are monotonic in this range. To compute the HD and AAIV in this

Table 4. AAIV and HD results for each treatment in Example 1 by considering [4, 5] as signal domain.

Treatment	x_1	x_2	AAIV	HD
1	1	1	195.55	251.74
2	1	-1	115.76	169.40
3	-1	1	156.65	218.50
4	$^{-1}$	-1	2350.59	3633.97

Table 5. Supposed experimental results for each signallevel in Example 2.

Treatment	x_1	x_2	t = 0	t = 1	t=2	t = 3
1	1	1	4.4	7.4	11.0	6.3
2	1	-1	5.5	8.3	10.0	7.4
3	-1	1	7.3	6.4	8.0	3.3
4	-1	-1	1.5	5.0	6.5	3.4



Figure 8. Fitted functional responses in Example 2.

range, we consider (4, 4.2, 4.4, 4.6, 4.8, 5) as signal levels and generate response values based on Eqs. (9)-(12) by adding standard normal residuals to them. It can be seen in Table 4 that HD results confirm the AAIV results. It should be noted that although the proposed method considers the continuous nature of the responses, it may incorporate an estimation error in the calculations. Therefore, it is very important to find an estimated function with acceptably small error.

4.2. Example 2: In-domain dispersion effect

Suppose that there is a problem with one functional response and two controllable factors. The functional response is affected by a signal factor (t) with four levels (0, 1, 2, 3). The target values are supposed to be (4.5, 7.3, 9, 6.4) and the experimental results are supposed as reported in Table 5. Suppose that the analyst is interested in ranking treatments by considering deviation from target as well as IDV. By considering polynomial order equal to 6, the SSE values are approximately zero and each function has only one maximum point. Hence, we are not worried about overfitting. Figure 8 illustrates the fitted functions for target and treatments. As can be seen, the fitted responses corresponding to treatments 1 and 2 have

small deviations from the target in comparison with treatments 3 and 4.

To better illustrate the efficiency of the proposed measure, consider another approach from the literature with discrete viewpoint to the signal levels. Wu [19] proposed an approach for optimizing multiple functional responses using double-exponential desirability function. Based on his proposition, the desirability for the kth treatment by considering NTB type uncorrelated responses is calculated as follows:

$$WD_{k} = \left[\prod_{h=1}^{m} \left[\prod_{u=1}^{\nu} \exp\left(-c_{h} \left|y_{hku} - g_{hu}\right|^{\alpha_{h}}\right)\right]^{\frac{1}{\nu}}\right]^{\frac{1}{m}}, (13)$$

where y_{hku} is the value of the *h*th response for the *u*th signal level in treatment k, g_{hu} is target value of the *h*th response for the *u*th signal level, ν is number of signal levels, *m* is number of functional responses, α_h is shape constant of the *h*th response, and c_h is scale constant of the *h*th response.

Using the predefined equations, WD (by considering $\alpha_h = c_h = 1$), AAIV, and WAAIV (by considering [0.2 0.2 0.6] as weight vector) are calculated and the results are summarized in Table 6. Based on the WD results, treatment 1 is selected as the best by a wide difference from the second treatment. This is

Table 6. The results of different measures in Example 2.

Treatment	WD	AAIV	WAAIV
1	0.56	3.25	2.39
2	0.37	3.00	1.20
3	0.12	5.20	3.96
4	0.07	7.89	3.67

because of ignoring the IDV. By employing the AAIV, treatment 2 may be selected as the best while it has a negligible difference from treatment 1. However, by employing the WAAIV, which can consider the comments of the analyst as weights, it can be seen that treatment 2 is very better than treatment 1.

4.3. Example 3: Correlated functional responses

Expand Example 1 to include two controllable factors, x_1 and x_2 , a signal factor, t, and two correlated functional responses, y_1 and y_2 . Target profiles for functional responses are $T_1 = t^5 + 2t^4 + 3t^3 - 2t^2 + t + 2.5$ and $T_2 = t^5 - 2t^3 + t^2 - 4t + 1.5$, respectively, which are depicted in Figure 9. The covariance matrix of the response functions is $\begin{bmatrix} 10 & 11.4\\ 11.4 & 18 \end{bmatrix}$. By replicating the experiment in each signal level of each treatment for 10 times, mean **m** and covariance matrix **s** of the functional responses may be estimated based on the observed values of the responses. Table 7 shows the experimental design, including mean and covariance



Figure 9. Target profiles for functional responses in Example 3.

Table 7. Experimental design of Example 3.

Treatment	t = -5	t = -3	t = -1	t = 1	t = 3	t = 5
1	$\mathbf{M}' = \begin{bmatrix} -2301.18\\ -2826.89 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} -180.81\\ -166.27 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} -3.49\\ -650 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} 7.22\\ -2.64 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} -475.95\\ -191.01 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} -4707.80\\ -2881.95 \end{bmatrix}$
	$\mathbf{S} = \begin{bmatrix} 3.57 & 5.81\\ 5.81 & 9.85 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 7.90 & 8.38\\ 8.38 & 11.47 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 7.17 & 9.44\\ 9.44 & 13.30 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 9.85 & 14.92\\ 14.92 & 24.29 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 6.03 & 7.40 \\ 7.40 & 9.28 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 8.18 & 10.27\\ 10.27 & 14.02 \end{bmatrix}$
2	$\mathbf{M}' = \begin{bmatrix} -2304.83\\ -2832.15 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} -180.97\\ -167.51 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} -2.31\\ 7.71 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} 8.84\\ -1.23 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} 471.24\\184.15 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} 4710.44\\2885.08 \end{bmatrix}$
2	$\mathbf{S} = \begin{bmatrix} 1.83 & 2.64\\ 2.64 & 4.64 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 5.49 & 6.98\\ 6.98 & 9.88 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 15.68 & 19.25\\ 19.25 & 24.77 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 6.81 & 7.81 \\ 7.81 & 10.12 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 17.07 & 21.63\\ 21.63 & 28.13 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 13.23 & 16.41 \\ 16.41 & 21.56 \end{bmatrix}$
9	$\mathbf{M}' = \begin{bmatrix} -2302.74\\ -2829.05 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} -180.66\\ -166.38 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} -2.24\\ 8.71 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} 7.78\\-1.73 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} 474.01\\188.15 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} 4705.18\\ 2878.56 \end{bmatrix}$
3	$\mathbf{S} = \begin{bmatrix} 18.01 & 24.57\\ 24.57 & 35.46 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 12.97 & 17.42\\ 17.42 & 24.00 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 8.95 & 12.69\\ 12.69 & 19.47 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 6.05 & 7.86\\ 7.86 & 11.60 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 6.72 & 8.83 \\ 8.83 & 12.20 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 4.61 & 6.85 \\ 6.85 & 10.64 \end{bmatrix}$
	$\mathbf{M}' = \begin{bmatrix} -2302.89 \\ -2828.66 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} -181.45 \\ -168.18 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} -2.00 \\ 7.66 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} 6.69\\ -4.18 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} 471.89\\ 185.19 \end{bmatrix}$	$\mathbf{M}' = \begin{bmatrix} 4708.43 \\ 2883.12 \end{bmatrix}$
4	$\mathbf{S} = \begin{bmatrix} 6.47 & 7.32 \\ 7.32 & 8.73 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 13.21 & 16.81\\ 16.81 & 21.84 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 6.29 & 8.84 \\ 8.84 & 13.28 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 4.92 & 5.66\\ 5.66 & 7.45 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 14.01 & 19.17\\ 19.17 & 27.00 \end{bmatrix}$	$\mathbf{S} = \begin{bmatrix} 6.24 & 8.38\\ 8.38 & 11.93 \end{bmatrix}$

Treatment	x_1	x_2	MAAIV	SAAIV	POC_k	D
1	1	1	(24.70 33.81)	$\begin{bmatrix} 166.76 & 188.29 \\ 188.29 & 219.48 \end{bmatrix}$	0.3872	0.5566
2	1	-1	$(29.67 \ 39.05)$	$\begin{bmatrix} 37.66 & 36.28 \\ 36.28 & 49.12 \end{bmatrix}$	0.0978	0.6194
3	-1	1	(25.48 35.28)	$\begin{bmatrix} 76.79 & 91.72 \\ 91.72 & 126.41 \end{bmatrix}$	0.3186	0.6222
4	-1	-1	$(26.14 \ 36.46)$	$\begin{bmatrix} 112.61 & 161.82 \\ 161.82 & 259.25 \end{bmatrix}$	0.3381	0.5361

Table 8. AAIVs and POC results in Example 3.

Table 9. Polynomial regression parameters corresponding to each treatment in Example 4.

${\it Treatment}$	x_1	x_2	x_3	B
1	-1	-1	-1	[-5.74,-50.91,784.57,1201,10724.97,-51981.87,-22339.75,-128407.13]
2	-1	0	0	[2,-45.91,300,-3000,11724.97,-10000,21000,-120000]
3	-1	1	1	[9.74,-40.91,-184.57,-7201,12724.97,31981.87,64339.75,-111592.87]
4	0	-1	0	$[-5,\ 5,\ -120,\ 3501,\ -2000,\ 1000,\ -48339.75,\ 110000]$
5	0	0	1	$[2.74,\ 10,\ -604.57,\ -700,\ -1000,\ 42981.87, -5000,\ 118407.13]$
6	0	1	-1	[2.26,-15,724.57,-2801,3000,-43981.87,53339.75,-228407.13]
7	1	-1	1	$[-4.26,\ 60.91,\ -1024.57,\ 5801,\ -14724.97,\ 53981.87,\ -74339.75,\ 348407.13]$
8	1	0	-1	$[-4.74,\ 35.91,\ 304.57,\ 3700,\ -10724.97,\ -32981.87,\ -16000,\ 1592.87]$

matrices for the functional responses in each signal level.

Now, we estimate the functional responses for each replicate and, then, compute the mean and covariance matrices of the AAIVs for each treatment. To calculate POC, we test the AAIVs for normality and consider $[0 \ 30]^2$ as their desired specification region. Now, we ignore the covariance structure and calculate the desirabilities to compare the results. To do this for each response, h, a desirability corresponding to mean with $UM_h = 60$ and a desirability corresponding to variance with $UV_h = 500$ are calculated and the total desirability is found by considering 0.25 as weight of each desirability. Results are summarized in Table 8; as shown, the first treatment presents the best POC_k value; however, the third treatment presents the best D value. This shows that ignoring covariance structure in the problems with correlative responses may result in a wrong choice.

4.4. Example 4: Dissolved dose of tablet

Suppose the problem of finding the best composition of a tablet comprised of three raw materials x1, x2, and x3, such that the relation between time t and dissolution of tablet y (based on milligrams) follows

a target function, T, given in Eq. (14):

$$T = -1.13t^{7} + 29.53t^{6} - 327.45t^{5} + 1999.12t^{4} - 7270.48t^{3} + 15805.83t^{2} - 19130.77t + 110033.64.$$
(14)

We consider an orthogonal array with 8 treatments as experimental design. The polynomial regression parameters, **b**, for each treatment are given in Table 9.

It may be dangerous if the dissolving speed of the tablet is higher than the target or if the dissolution in a time point has a large difference from its target. Hence, we should assign a larger weight to integral values when the target function is below the treatment function. Also, the maximum difference should be considered as well, because increasing the speed of dissolution may be dangerous. Since there is only one functional response in this problem, we do not need to construct the overall index and optimization may be done on the WAAIV. By considering [0.1, 18, 3] as the weight vector and [0, 10] as range, WAAIVs are calculated with the results given in Table 10. The estimate of WAAIV is obtained using ordinary least squares method after removing the terms with large p-values (p-values are provided

treatm

${\it Treatment}$	x_1	x_2	x_3	WAAIV
1	-1	-1	-1	797098.7
2	-1	0	0	443368.4
3	-1	1	1	700695.5
4	0	-1	0	427297.3
5	0	0	1	5783643.0
6	0	1	-1	1733688.0
7	1	-1	1	14908102.0
8	1	0	-1	1096071.0

Table 10. WAAIV for each treatment in Example 4.

ō	1	0	-1	1090071.0

Treatment	x_1	x_2	x_3	y_1	y_2	y_3
1	-1	-1	-1	5.1	8.3	0.78
2	-1	1	-1	5.4	8.24	0.72
3	1	-1	-1	6.1	8.49	0.7
4	1	1	-1	6.2	8.48	0.77
5	0	0	-1	5.6	8.17	0.78
6	-1.41	0	1	5.8	9	0.65
7	1.41	0	1	7.3	9.56	0.67
8	0	-1.41	1	7.6	9.28	0.69

Table 11. Experimental design of the problem.

in parentheses), as follows:

0

0

9

10

 $\widehat{WAAIV} = 3679816(0.006) + 3548565_{x_1}(0.016)$

1.41

0

1

1

7.9

5.1

9.62

8.95

0.65

0.62

 $+2960930_{x_3}(0.021)$

 $+3477109_{x_1 x_3}(0.024), R^2 = 91.40\%.$ (15)

Finally, by employing GA with population size of 1000, and crossover and mutation rates of 0.7 and 0.4, after 60 iterations, we achieve [-0.9605, 0.2288, 0.7172] as the best factors levels array with WAAIV equal to 45.7711. A confirmation experiment may be performed to check the validity of the GA result.

5. A real case

Consider the real case given by Fogliatto [16] from the pet food production industry about the development of an alternative formula for a well-known brand of dog biscuits. The percentages of three ingredients are rewritten as two mixture variables and, thus, by considering the biscuit thickness, there are three controllable factors. Levels are defined based on a Central Composite Design (CCD) including ten treatments illustrated in Table 11.

There are three real responses and a simulated functional response in this problem. Response types

ent.		
Treatment	Best order	SSE
Target profile	8	0.16968
1	8	1.21
2	8	0.11237
3	8	0.16784
4	8	0.13582
5	8	0.13898
6	8	0.22952
7	8	0.18642
8	8	0.18127
9	8	0.23478
10	8	0.20691

Table 12. Best fitting order for each target and

and specifications are as follows: y_1 is nominal-thebest ($U_1 = 8$; $m_1 = 7.5$; $L_1 = 5$), y_2 is nominalthe-best ($U_2 = 10$; $m_2 = 9.5$; $L_2 = 8$), and y_3 is larger-the-better ($U_3 = 0.8$; $L_3 = 0.6$), where m_i is the target value and U_i and L_i are the upper and lower specification limits of the *i*th response, respectively. Note that the values of functional response in signal levels are driven from Table AI of Fogliatto [16].

As expressed in the previous section, the next step is finding the best fit with the best order for each treatment. To avoid overfitting, we limit the maximum order of functions between 5 and 9. Table 12 shows the best order with corresponding SSE for each treatment.

Figure 10 shows the fitted functions for target and the other treatments. The shapes of fitted functions are smooth enough and there are not several maximum or minimum points for each function. Hence, overfitting is unlikely. It can be seen that fitted profile of the treatment 3 can visually be selected as the nearest profile to the target.

The next step is calculating the absolute integral value between target profile and each of the treatments from one cross point to the next one. To do this, cross points between target and each of profiles should be found. After that, aggregating absolute integral values and selecting the treatment with minimum AAIV are desirable. Table 13 shows the cross points and AAIV for each of the treatments. It should be mentioned that lower limit of the integral range is equal to zero and upper limit is assumed to be equal to the number of values in functional response for each treatment.

Now, the desirability of each response and AAIV should be calculated for each treatment. Then, we should find the estimating equation for desirabilities. By considering $\alpha = 1$, and $UM_h = 25$, desirabilities (d_i) can be calculated as shown in Table 14. Note that total desirability (D_i) has been calculated by considering $w_i = 0.25$, which is same as the first scenario in Fogliatto [16].

As shown in Table 14, treatment 8 can be selected



Figure 10. Fitted polynomial profiles for target and treatments.

Table 13. Cr	oss points	and AA	IV for	each	treatment.
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Treatment	Cross points					AAIV	
1	1.3011	3.4710	7.7301	20.3327	28.0974	31.5621	3.6308
2	1.2301	38.9803	40.8241	45.0043			12.7561
3	1.3257	34.2576	40.1817	44.7791			0.7264
4	1.2358	37.5111	40.4301	44.8857			7.3987
5	1.2368	37.4034	40.3069	44.8664			6.7312
6	1.2069	40.0524	44.6308				25.8691
7	1.1436	40.1416	44.7220				3.0190
8	1.0653	40.1753	44.7377				2.0058
9	1.2044	40.0414	44.6190				20.0280
10	1.1930	40.0974	44.6747				1080.1195

Table 14. Desirability values for each treatment.

Treatment	d_1	d_2	d_3	d_{AAIV}	D
1	0.0400	0.2000	0.9000	0.8185	0.2771
2	0.1600	0.1600	0.6000	0.3622	0.2731
3	0.4400	0.3267	0.5000	0.9637	0.5130
4	0.4800	0.3200	0.8500	0.6301	0.5355
5	0.2400	0.1133	0.9000	0.6634	0.3570
6	0.3200	0.6667	0.2500	0	0
7	0.9200	0.8800	0.3500	0.8491	0.7004
8	0.8000	0.8533	0.4500	0.8997	0.7251
9	0.2000	0.7600	0.2500	0	0
10	0.0400	0.6333	0.1000	0	0

as the best one. To search for better treatments, we need to find fitting models for desirabilities. Note that normality assumption cannot be rejected for any of the desirabilities. After fitting different regression models and removing terms with large p-values (p-values are provided in parentheses), we find the following equations with acceptable coefficients of determination, R^2 :

$$\hat{d}_1 = 0.26556 (0.042) + 0.19633_{x_1} (0.046),$$

 $R^2 = 81.42\%,$ (16)

$$d_{2} = 0.373323 (0.015) + 0.073657_{x_{1}} (0.037) + 0.267682_{x_{3}} (0.009) + 0.0656_{x_{1}^{2}} (0.055) + 0.082349_{x_{2}^{2}} (0.044), \quad R^{2} = 99.98\%,$$
(17)

$$\hat{d}_3 = 0.515 (0.000) - 0.235_{x_3} (0.003)$$

+ $0.1625_{x_1x_2} (0.066), \quad R^2 = 87.42\%,$ (18)



Figure 11. Changes in total desirability and controllable factors versus iteration numbers.

 $\hat{d}_{AAIV} = 0.51867(0.000) + 0.20219_{x_1}(0.035)$

$$-0.25808_{x_2} (0.013) - 0.16891_{x_3} (0.044),$$

$$R^2 = 86.67\%.$$
 (19)

To prevent unreal results, we set 1 for $d_i > 1$ and 0 for $d_i < 0$. Now, we can search for better treatments. To do this, we employ GA with population size of 1000, and crossover and mutation rates of 0.7 and 0.4, respectively. After 60 iterations, we find a treatment with $x_1 = 1.41$, $x_2 = 1.41$, $x_3 = -0.69$, and $\hat{D}=0.6487$ as the best result. Note that although based on Table 14, the total desirability of treatment 8 is 0.7251, we find D=0.6487 for it. Hence, it is necessary to do a real experiment with the setting of $x_1 = 1.41, x_2 = 1.41$, and $x_3 = -0.69$ to find the exact total desirability value and compare it with treatment 8. Figure 11 shows the changes in total desirability and controllable factors in each treatment versus iteration numbers. As can be seen, increasing x_1 and x_2 will increase total desirability.

It should be noted that in [16], treatment 8 is selected as the best by considering equal weights for calculating total desirability. By solely considering functional response, Fogiatto [16] found treatment 4 using maximum and summation operators and treatment 5 using average operator as the best. However, as can be seen in Table 13, we find treatment 3 as the best by solely considering functional response. Hence, it can be concluded that finding treatment 8 as best by both methods is because of other responses (i.e. y_1, y_2 , and y_3).

6. Conclusion and future researches

We proposed a novel method for optimizing multiresponse problems with continuous functional response(s) and considered two different dispersion ef-The first dispersion effect was "In Domain fects. Variation (IDV)," which was related to the difference in deviation of estimated function, and the second one was "Between Replicates Variation (BRV)," which might be considered when there were different estimated functions for each treatment because of replicates. The first important step of the proposed method was finding an acceptable polynomial fitting model for functional response in each treatment from the observed values. Thus, if we could not find fitted functions with acceptable estimating errors, the proposed method might not be applicable. We found the cross points between target function(s) and estimated function(s) in each treatment. Thereafter, we proposed the absolute integral value as the measure. In some special problems, it was important to distinguish between positive and negative deviations from the target function. Moreover, IDV might be considered in this step. For analyzing such cases, WAAIV was proposed. To aggregate uncorrelated AAIVs (or WAAIVs) in one measure, we used desirability function. In the problems with correlated functional responses, we employed Proportion Of Conformance (POC) proposed by Chiao and Hamada [32] to transform correlated AAIVs (or WAAIVs) into one measure. The BRV might be considered in this step. Finally, we applied GA to find the best levels for factors. Some illustrative examples showed that our proposed continuous method was efficient in the problems with functional responses. The application of the proposed method was illustrated through a real case in pet food production extracted from [16]. In this paper, we supposed that the effect of nuisance factor was randomly distributed through the randomization. Considering these factors in the proposed method can be a direction for future research. Moreover, we supposed that there was a polynomial relation between functional response and the signal factor. A future study may be done to focus on other types of estimated functions and use spline regression to model the functional responses. In addition, investigating the multi-response problems with correlational structure between classic and functional responses can be a fruitful area of research.

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