Worst-Case Tolerance Synthesis for Low-Sidelobe Sparse Linear Arrays Using a Novel Self-Adaptive Hybrid Differential Evolution Algorithm

Tao Ni*, Yong-Chang Jiao, Li Zhang, and Zi-Bin Weng

Abstract—A worst-case tolerance synthesis problem for low-sidelobe sparse linear arrays is solved by using a novel self-adaptive hybrid differential evolution (SAHDE) algorithm. First, we establish a worst-case tolerance synthesis model for low-sidelobe sparse linear arrays, in which random position errors are considered and assumed to obey the Gaussian distributions. Through the random sampling, the random model is converted to a deterministic optimization problem. Then, a novel SAHDE algorithm is presented for solving the problem. As a modification to the existing hybrid differential evolution algorithm, a simplified quadratic interpolation (SQI) operator is used to tune the control parameters self-adaptively, establishing the connections between control parameters and the fitness values. In order to determine appropriate control parameter values quickly, a selection operation is also used. Detailed implementation procedure for the SAHDE algorithm is presented, and some numerical results show its effectiveness. Finally, for the deterministic optimization problem, we present a fast way for calculating its fitness values. The SAHDE algorithm is used to obtain optimal nominal element positions. Simulated results illustrate that the worst-case peak sidelobe levels for the sparse linear arrays are improved evidently. The SAHDE algorithm is efficient for solving the worst-case tolerance synthesis problem.

1. INTRODUCTION

Over the past years, synthesis of unequally spaced arrays has been studied in depth. Compared with periodic arrays, since the array periodicity is broken, unequally spaced arrays have shown several promising characteristics and probably have some potential advantages [1–4]. Specifically, grating lobes can be restrained in this case, and the array may get lower peak sidelobe levels (PSLLs), while the sidelobes of equally spaced arrays with uniform excitation distribution are greater than about $-13.4\, dB$. Furthermore, it is possible to reduce the number of elements in one certain aperture, which helps to reduce the array cost. Recently, the design techniques mainly focus on two kinds of unequally spaced arrays: thinned arrays [5,28–29], which are derived by exciting some elements in an equally spaced array, and sparse arrays [2,3,6–11,30–31], in which the spacings of elements are random. In the sparse arrays with uniform excitation distribution, the element spacings with more degree of freedom are optimized to gain lower PSLLs, which makes this kind of arrays be studied widely. Practically the element positions of the sparse array possess the random errors. However, in the existing research, influence of the element position errors is always neglected. For sparse arrays, the low-sidelobe levels are directly related to position of each element, and presence of the position errors will have a great impact on peak sidelobe levels (PSLLs). Therefore, the tolerance synthesis problem considering random position errors is worth of research.
As a pioneering work, Schjaer-Jacobsen [13] studied the worst-case tolerance optimization of linear broadside arrays, by using algorithms for optimum tolerance assignment problems [12]. However, in his algorithms, derivative information is needed. In [14], Jiao et al. studied the worst-case tolerance optimization synthesis for low sidelobe linear arrays by using the regular polyhedron method, a direct local search method. However, a suitable initial value should be provided, and a local optimal solution is obtained. In [15], the interval arithmetic is applied to analyze only impact of the manufacturing tolerances of the excitation amplitudes on the radiated array patterns. In [32], the interval analysis is applied to analyze the pattern tolerances when phase errors exist in the excitation weights. A design of linear arrays in the presence of tolerances on the amplitude weights is optimized by interval analysis and convex optimization in [33]. The linear array’s excitation tolerances are optimized for the robust amplitude beamforming in [34]. However, these works in [33, 34] are designed for array’s excitations and not for sparse array’s position errors. In [16], the tolerable linear arrays are designed by using the genetic algorithm. The tolerance optimization is highly important and computationally difficult [16], so evolutionary algorithms are suitable for solving the problems.

In recent years, evolutionary algorithms, such as genetic algorithm (GA) [5, 6, 9, 23], particle swarm optimization (PSO) [7, 24–27], and differential evolution (DE) [8, 10, 11], are widely used to solve the antenna optimization problem. In [5, 6, 9, 23], GA and modified GA are used to synthesize the thinning array, sparse array, and adaptive array respectively. In [7], PSO is used to synthesis the linear array and control the null location. In [24–26], PSO algorithms are used to optimize a prefractal monopole antenna, broadband nanoplasmonic arrays and multiple band antenna. In [27], PSO algorithm is applied in microwave image. As a population-based stochastic global optimization algorithm, DE has been successfully applied to many optimization problems. In [8], the authors have been presented an overview of DE-based approaches used in electromagnetics, which pointing out that the DE algorithm and its modified algorithm are effective to solve the electromagnetics problem. However, tuning control parameters involved in DE is a challenging task. For tolerance optimization problems, it is certain to cause computationally difficult and waste time by tuning control parameters blindly. In some modified self-adaptive algorithms such as SaDE and jDE [17, 18], the control parameters are self-adapted to gain better optimal results, based on the learning experiences from previous generations. However, these self-adaptive algorithms do not make full use of the fitness values available in their self-adaptive strategies.

Simplified quadratic interpolation (SQI) is a powerful direct search method. In [19–22], the SQI method is used to improve performance of some evolutionary algorithms, such as Price’s algorithm, the genetic algorithm, and the DE algorithm. In [22], Zhang et al. proposed a hybrid differential evolution (HDE) algorithm with the SQI operator to solve complex antenna design problems. However, the self-adaptive technique is not used in [22].

In this paper, inspired by above ideas, we propose a novel self-adaptive hybrid differential evolution (SAHDE) algorithm for solving worst-case tolerance synthesis problem. For the low-sidelobe sparse linear array, taking the nominal element positions as design parameters, a worst-case tolerance synthesis model considering the random position errors is established. The errors are assumed to obey the Gaussian distributions. Sample points are chosen randomly within a fixed position tolerance region, and then the random tolerance synthesis model is converted to a deterministic optimization problem. In order to solve the problem easily, we presented a new self-adaptive parameter control technique, and a novel SAHDE algorithm. In the algorithm, the control parameters (difference scale factor $F$, and crossover probability factor $CR$) are produced by the SQI operator, and the connections between control parameters and fitness values available are established. 11 benchmark problems are used to validate effectiveness of the novel algorithm. Moreover, a sparse linear array is optimized to gain lower PSLLs by using the SAHDE algorithm. Simulated results show that the SAHDE algorithm is efficient. Finally, a fast way for calculating fitness values of the deterministic optimization problem is presented. The SAHDE algorithm is used to minimize the worst-case PSLLs and obtain optimal nominal element positions for the low-sidelobe sparse linear array. Simulated results illustrate that the worst-case peak sidelobe levels for the sparse linear arrays are improved evidently.

This paper is organized as follows. The worst-case tolerance synthesis model for the low-sidelobe sparse linear array is established in Section 2. The SAHDE algorithm and its numerical experiments are presented in Section 3. Worst-case tolerance synthesis of low sidelobe sparse linear arrays by using
the SAHDE algorithm and related simulation results are shown in Section 4. Conclusions are drawn in Section 5.

2. MODEL OF WORST-CASE TOLERANCE SYNTHESIS FOR THE LOW-SIDELOBE SPARSE LINEAR ARRAY

Geometry of the symmetrical sparse linear array with isotropic point source element is shown in Fig. 1, where \( N \) is the element number, and \( d_i \) represents position of the \( i \)th element, \( i = -M, \ldots, -1, 0, 1, \ldots, M \). Define

\[
D = (d_{-M}, \ldots, d_{-1}, d_0, d_1, \ldots, d_M)
\]  

(1)

The array pattern of the \( N \)-element sparse linear array with uniform excitation is expressed as

\[
E(\theta, D) = \sum_{i=-M}^{M} \exp(jk \sin \theta d_i)
\]

(2)

where \( k = 2\pi/\lambda \) is the propagation constant.

When the array is symmetrical and the center element located in the origin of coordinates, \( d = 0 \), \( d_i = d_{-i} \), and the array pattern is expressed as

\[
E(\theta, D) = 1 + 2 \sum_{i=1}^{M} \cos(jk \sin \theta d_i)
\]

(3)

The design objective is to minimize the PSLL value and find the optimal element positions. For this purpose, the fitness function is defined as

\[
\text{fitness}_0(D) = \max_{\theta \in S} \left| \frac{E(\theta, D)}{\max(E)} \right|
\]

(4)

where \( S \) represents the sidelobe region corresponding to \( D \). As a constraint condition, \( d_i \) should satisfy [6]

\[
\min \{d_i - d_j, 0 \leq j \leq i \leq M \} > d_{\min}
\]

(5)

where \( d_{\min} \) is the minimum element spacing of the array.

By minimizing Eq. (4) under constraint of Eq. (5), proper \( D \) can be gained. In engineering applications, however, the element position errors always exist, and the random position errors deteriorate the PSLL of the sparse linear array. In the following, taking the nominal element positions as design parameters, a worst-case tolerance synthesis model considering random position errors with fixed tolerance region is established.

Usually, the element positions are symmetrical. However, the element errors may be asymmetrical. Define the actual element position vector as

\[
\text{Da} = (d_{a-M}, \ldots, d_{a-1}, d_{a0}, d_{a1}, \ldots, d_{aM})
\]

(6)

For every element in \( \text{Da} \), we assume that \( d_{ai} \) obeys the Gaussian distribution with mean \( d_i \) and standard deviation \( \sigma_i \)

\[
da_i \sim N(d_i, \sigma_i^2)
\]

(7)

Figure 1. Geometry of the symmetrical sparse linear array.
Define the position error vector as
\[ \mathbf{Dc} = (d_{c_{-M}}, \ldots, d_{c_{-1}}, d_{c_0}, d_{c_1}, \ldots, d_{c_M}) \] (8)
dc_i represents the ith element position error, i = −M, −(M − 1), ..., 0, ..., M − 1, M. Actual position coordinate vector \( \mathbf{Da} \) is expressed as
\[ \mathbf{Da} = \mathbf{D} + \mathbf{Dc} \] (9)
For every element in \( \mathbf{Dc} \), obviously \( dc_i \) obeys the Gaussian distribution with mean 0 and standard deviation \( \sigma_i \)
\[ dc_i \sim N(0, \sigma_i^2) \] (10)
In Gaussian distribution \( N(\mu, \sigma) \), 99.7% data are within interval \( (\mu - 3\sigma, \mu + 3\sigma) \), and only 0.3% data are outside. In fact, the 0.3% possibility is extremely small, and it can be ignored. Usually, interval \( (\mu - 3\sigma, \mu + 3\sigma) \) is considered instead of all possibilities. So, the tolerance region is defined as
\[ \Omega = \{ \mathbf{Dc} \mid -3\sigma_i < dc_i < 3\sigma_i, i = -M, \ldots, M \} \] (11)
For the \( N \)-element sparse linear array with uniform excitation considering the random position errors, the array pattern is expressed as
\[ E(\theta, \mathbf{D}, \mathbf{Dc}) = \sum_{m=-M}^{M} \exp[i k \sin \theta (d_m + dc_m)] \] (12)
The PSLLs can be calculated by
\[ SLL(\mathbf{D}, \mathbf{Dc}) = \max_{\theta \in S} \left| \frac{E(\theta, \mathbf{D}, \mathbf{Dc})}{\max(E)} \right| \] (13)
where \( S \) represents the sidelobe region corresponding to \( \mathbf{D} \) and \( \mathbf{Dc} \). In order to gain better PSLLs, taking the nominal element positions \( \mathbf{D} \) as design parameters, the optimization problem can be expressed as
\[ \min_{\mathbf{D}} SLL(\mathbf{D}, \mathbf{Dc}) \] (14)
Since \( \mathbf{Dc} \) is a Gaussian random vector, Problem (14) is a stochastic optimization problem, and it is difficult to solve Problem (14) directly. In order to solve Problem (14) easily, we randomly choose \( NS \) sample points in the tolerance region of Eq. (11)
\[ \mathbf{Dc}_i, \quad i = 1, 2, \ldots, NS \]
The optimization task is to minimize the PSLLs for all possible \( \mathbf{Dc} \) vectors, including the worst-case PSLLs. Thus Problem (14) can be formulated as the multi-objective optimization problem
\[ \min_{\mathbf{D}} SLL(\mathbf{D}, \mathbf{Dc}_i), \quad i = 1, 2, \ldots, NS \] (15)
Problem (15) is a multi-objective optimization problem with many objectives, and it is also difficult to solve Problem (15) directly. Usually, if the worst-case PSLL is minimized, the optimization task is reached. We should pay attention to the worst-case PSLLs. Therefore, Problem (15) can be converted to the single-objective optimization problem
\[ \min_{\mathbf{D}} \max_{1 \leq i \leq NS} SLL(\mathbf{D}, \mathbf{Dc}_i) \] (16)
The fitness function is expressed as
\[ \text{fitness}(\mathbf{D}) = \max_{1 \leq i \leq NS} SLL(\mathbf{D}, \mathbf{Dc}_i) \] (17)
We hope that Problem (16) can be solved by the HDE algorithm [22]. However, it is difficult to choose the control parameters in the HDE algorithm, and proper control parameters may be obtained after many times experiments. Since calculating the fitness function (17) is very time-consuming, it is necessary to reduce blind experiment times. A self-adaptive HDE algorithm is suitable for solving this problem. Aiming at this problem, we propose a novel SAHDE algorithm in the following section.
3. NOVEL SELF-ADAPTIVE HYBRID DIFFERENTIAL EVOLUTION ALGORITHM AND ITS NUMERICAL EXPERIMENTS

3.1. Hybrid Differential Evolution Algorithm

Differential evolution (DE) is a simple powerful population-based stochastic search algorithm. In the HDE algorithm (DESQI for short in [22]), the simplified quadratic interpolation operator is incorporated with DE to accelerate the convergence of DE. The main operations in the HDE algorithm are described as follows.

3.1.1. Initializing Parameters and Population

Initializing parameters, difference scale factor $F$, crossover probability factor $CR$, population size $NP$, and original population $X_0$, $X_0 = (X_{1,0}, X_{2,0}, \ldots, X_{i,0}, \ldots, X_{NP,0})$, where $X_{i,0}$ is a $D$-dimension vector. In [22], $F = 0.5$, $CR = 0.9$, and $NP$ is chosen based on the specific problem.

3.1.2. Mutation

At $G$th generation, $X_{i,G} = (x_{1,i,G}, x_{2,i,G}, \ldots, x_{j,i,G}, \ldots, x_{D,i,G})$. This operation creates the mutant trial vector $V_{i,G+1} = (v_{1,i,G+1}, v_{2,i,G+1}, \ldots, v_{j,i,G+1}, \ldots, v_{D,i,G+1})$ based on the current parent population $X_G$. The DE’s strategy is expressed as

$$V_{i,G+1} = X_{r1,G} + F \cdot (X_{r2,G} - X_{r3,G})$$

where $X_{r1,G}, X_{r2,G},$ and $X_{r3,G}$ represent three individuals in $G$th generation, and indexes $r1, r2,$ and $r3$ represent different random integers in range $[1, NP]$, which are also different from index $i$, $i = 1, 2, \ldots, NP$.

3.1.3. Crossover

After mutation, the offspring vector $U_{i,G+1} = (u_{1,i,G+1}, u_{2,i,G+1}, \ldots, u_{j,i,G+1}, \ldots, u_{D,i,G+1})$ is produced by the binomial crossover operation

$$u_{j,i,G+1} = \begin{cases} v_{j,i,G+1}, & \text{if } rand(0,1) < CR \text{ or } j = rand(1, D) \\ x_{j,i,G}, & \text{otherwise} \end{cases}$$

3.1.4. Selection

The selection operation chooses the better one from the parent vector $X_{i,G}$ and the trial vector $U_{i,G}$ according to their fitness values $f(\cdot)$. For example, if we deal with a minimization problem, the selected vector is given by

$$X_{i,G+1} = \begin{cases} U_{i,G+1}, & \text{if } f(U_{i,G+1}) < f(X_{i,G}) \\ X_{i,G}, & \text{otherwise} \end{cases}$$

3.1.5. The SQI Operator

As a powerful direct search method, the simplified quadratic interpolation (SQI) operator can be used to improve the local search ability in DE [22]. The SQI operator with three points, $x_1, x_2$ and $x_3$, is defined as follows.

$$p_{i,j} = 0.5 \cdot \frac{(x_{2i} - x_{3i}) \cdot f(x_{1}) + (x_{3i} - x_{1i}) \cdot f(x_{2}) + (x_{1i} - x_{2i}) \cdot f(x_{3})}{(x_{2i} - x_{3i}) \cdot f(x_{1}) + (x_{3i} - x_{1i}) \cdot f(x_{2}) + (x_{1i} - x_{2i}) \cdot f(x_{3})}$$

where $P = (p_{1,1}, p_{2,1}, \ldots, p_{D,j}), x_j = (x_{j1}, x_{j2}, \ldots, x_{jD}), j = 1, 2, 3$. $P$ is a new trial vector.

The SQI operator is described briefly as follows. Firstly, find the best and worst individuals, as well as two random individuals which are different from the best and worst ones, and set the best one as $x_1$, and two random individuals as $x_2$ and $x_3$. Secondly, compute $P$ by Eq. (21), and evaluate $P$. Thirdly, if $P$ is better than the worst one, replace the worst one by $P$ in the population.
3.2. Self-Adaptive Parameter Control Technique

In the DE algorithm, parameter $F$, the difference scale factor, is real, which regulates the difference vector. Larger $F$ values are helpful for improving the population diversity, while smaller $F$ values make the algorithm converge fast. Crossover probability factor $CR$ represents the proportion of parents in offspring. Besides, an inverse correlation between $CR$ and the local search ability exists. Therefore, the search ability of the algorithm can be regulated by monitoring $F$ and $CR$ effectively.

In the HDE algorithm [22], the self-adaptive technique is not used, and the control parameters are set as constants, thus the algorithm cannot adapt to different optimization problems. Some self-adaptive DE algorithms, such as SaDE and jDE, do not make full use of the fitness values available in their self-adaptive strategies. So far, connection between fitness values available and control parameters is not concerned widely.

Based on these observations, we propose a new self-adaptive parameter control technique. In $G$th generation, one individual corresponds to a set of control parameters $(F_{i,G}, CR_{i,G})$, which is shown in Table 1. Thus, the control parameters are no longer scalar quantities, but vectors. We use $F_G = (F_{1,G}, F_{2,G}, \ldots, F_{i,G}, \ldots, F_{NP,G})$ and $CR_G = (CR_{1,G}, CR_{2,G}, \ldots, CR_{i,G}, \ldots, CR_{NP,G})$ in the evolutionary process.

<table>
<thead>
<tr>
<th>Individual</th>
<th>$F$</th>
<th>$CR$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$X_{1,G}$</td>
<td>$F_{1,G}$</td>
<td>$CR_{1,G}$</td>
</tr>
<tr>
<td>$X_{2,G}$</td>
<td>$F_{2,G}$</td>
<td>$CR_{2,G}$</td>
</tr>
<tr>
<td>$\ldots$</td>
<td>$\ldots$</td>
<td>$\ldots$</td>
</tr>
<tr>
<td>$X_{NP,G}$</td>
<td>$F_{NP,G}$</td>
<td>$CR_{NP,G}$</td>
</tr>
</tbody>
</table>

In order to solve a specific optimization problem efficiently, the control parameter values should adapt to the problem. Our self-adaptive parameter control technique should find a set of proper control parameter values corresponding to the problem, by using the fitness values available. The SQI operator can be used to achieve this goal. Now we introduce the new self-adaptive parameter control technique in detail. Define difference scale factor intervals: $F_t$ and $F_b$, $F_b \subseteq F_t$. Define the selection intervals of the crossover probability factor: $CR_t$ and $CR_b$, $CR_b \subseteq CR_t$.

In the first generation, $F_0$ and $CR_0$ are randomly produced in intervals $F_t$ and $CR_t$, respectively. In other generations, the control parameter vectors are generated by the SQI operator. However, the produced control parameters may lie in inappropriate intervals, which may reduce the solution accuracy. In order to overcome this shortcoming, we introduce a generation interval $G_t$. When the generation number is a multiple of $G_t$, the prior knowledge for the control parameters is used. Values in vectors $F_G$ and $CR_G$ are chosen randomly in intervals $F_b$ and $CR_b$, respectively. When the generation number is not a multiple of $G_t$, if the produced control parameters are not in intervals $F_t$ and $CR_t$, we choose the new control parameters as those corresponding to $r1$ in Eq. (18); Otherwise, the produced control parameters are chosen as the new ones.

It is worth to note that a selection operation for control parameters is also used for determining the appropriate control parameter values quickly. When the offspring is better than the corresponding parameters are chosen as the new ones. When the offspring is better than the corresponding parameters are chosen as the new ones. For example, in the current population, values in $F_G$ are generated by Eqs. (22) and (23).

$$F_{new_{i,G+1}} \begin{cases} F_{pi,G}, & \text{if } F_{pi,G} \in F_t \text{ & } \text{mod } (G, G_t) \neq 0 \\ F_{r1,G}, & \text{if } F_{pi,G} \notin F_t \text{ & } \text{mod } (G, G_t) \neq 0 \\ \text{rand}(F_b), & \text{if } \text{mod } (G, G_t) = 0 \end{cases}$$

$$F_{pi,G} = 0.5 * \frac{(F_{r2,G} - F_{r3,G})f(X_{r1,G}) + (F_{r3,G} - F_{r1,G})f(X_{r2,G}) + (F_{r1,G} - F_{r2,G})f(X_{r3,G})}{(F_{r2,G} - F_{r3,G})f(X_{r1,G}) + (F_{r3,G} - F_{r1,G})f(X_{r2,G}) + (F_{r1,G} - F_{r2,G})f(X_{r3,G})}$$ (22)
While values in $\text{CR}_G$ are also generated by Eqs. (24) and (25).

\[
CR_{p_i,G}^{i,G} = 0.5 * \frac{(CR_{r2,G}^2 - CR_{r3,G}^2)f(X_{r1,G}) + (CR_{r1,G}^2 - CR_{r2,G}^2)f(X_{r3,G})}{(CR_{r2,G}^2 - CR_{r3,G}^2) * f(X_{r1,G}) + (CR_{r1,G}^2 - CR_{r2,G}^2) * f(X_{r3,G})}
\]

(24)

\[
CR_{new_i,G+1} = \begin{cases} 
    CR_{p_i,G}, & \text{if } CR_{p_i,G} \in CR_t \text{ and } \text{mod}(G,G_t) \neq 0 \\
    CR_{r1,G}, & \text{if } CR_{p_i,G} \notin CR_t \text{ and } \text{mod}(G,G_t) \neq 0 \\
    \text{rand}(CR_b), & \text{otherwise}
\end{cases}
\]

(25)

where $F_{new_i,G+1}$ and $CR_{new_i,G+1}$ represent the produced control parameters for the $i$th individual in the $G+1$th generation. In order to search appropriate control parameter values quickly, the selection operation is also used for $F_G$ and $CR_G$. Better values in vectors $F_{G+1}$ and $CR_{G+1}$ are selected by Eqs. (26) and (27) respectively for the minimization problem.

\[
F_{i,G+1} = \begin{cases} 
    F_{new_i,G+1}, & \text{if } f(U_{i,G+1}) < f(X_{i,G+1}) \\
    F_{i,G}, & \text{otherwise}
\end{cases}
\]

(26)

\[
CR_{i,G+1} = \begin{cases} 
    CR_{new_i,G+1}, & \text{if } f(U_{i,G+1}) < f(X_{i,G+1}) \\
    CR_{i,G}, & \text{otherwise}
\end{cases}
\]

(27)

Based on the new self-adaptive parameter control technique, we propose a novel SAHDE algorithm, and its flow chart is shown in Fig. 2. And the detail procedure is presented as follows.

**Step 0:** Set parameters: Choose suitable $NP$, $F_t$, $CR_t$, $F_b$, $CR_b$ and $G_t$. Set $G = 0$.

![Figure 2. A flow chart of the SAHDE algorithm.](image-url)
Step 1: Initialize population: Choose randomly the initial population $X_0 = (X_{1,0}, X_{2,0}, \ldots, X_{NP,0})$ in the feasible region.

Step 2: Evaluate the fitness values for the initial population: Calculate $f(X_{i,0})$, $i = 1, 2, \ldots, NP$.

Step 3: Produce vectors $F_1$ and $CR_1$: Produce $F_{i,1}$ randomly in $F_t$, and $CR_{i,1}$ randomly in $CR_t$, $i = 1, 2, \ldots, NP$. Form $F_0 = (F_{1,1}, F_{2,1}, \ldots, F_{NP,1})$, and $CR_1 = (CR_{1,1}, CR_{2,1}, \ldots, CR_{NP,1})$.

Step 4: Mutation: Calculate the mutant trial vector $V_{i,G+1}$ by using the DE mutation strategy

$$V_{i,G+1} = X_{i,G} + F_{i,G} \cdot (X_{r_2,G} - X_{r_3,G})$$

where indexes $r_1, r_2$, and $r_3$ are three different random integers in range $[1, NP]$, and also different from index $i$. Especially, among $r_1, r_2$, and $r_3$ individuals in current population, $r_1$ is the best, and $r_3$ is the worst.

Step 5: Crossover: Calculate the offspring vector $U_{i,G+1} = (u_{1,i,G+1}, u_{2,i,G+1}, \ldots, u_{j,i,G+1}, \ldots, u_{D,i,G+1})$ by the binomial crossover operation

$$u_{j,i,G+1} = \begin{cases} 
  v_{j,i,G+1}, & \text{if } rand(0,1) < CR_{i,G} \text{ or } j = rand(1, D) \\
  x_{j,i,G}, & \text{otherwise}
\end{cases}$$

Step 6: Evaluate the fitness values for the offspring vector: Calculate $f(U_{i,G+1})$, $i = 1, 2, \ldots, NP$.

Step 7: Selection: Determine a new vector $X_{i,G+1}$ by using the selection operation Eq. (20).

Step 8: The SQI operation: Execute the SQI operation, as shown in the original HDE algorithm.

Step 9: Termination condition judgement: If the stopping criterion is not met, go to Step 10. Otherwise, the algorithm is terminated.

Step 10: Produce vectors $F_{G+1}$ and $CR_{G+1}$: First, calculate $F_{new,i,G+1}$ by using Eq. (23), and $CR_{new,i,G+1}$ by Eq. (25), $i = 1, 2, \ldots, NP$, where indexes $r_1, r_2$ and $r_3$ are same as those in Step 4. Then, determine $F_{i,G+1}$ by Eq. (26), and $CR_{i,G+1}$ by Eq. (27), $i = 1, 2, \ldots, NP$. Finally, form $F_{G+1} = (F_{1,G+1}, F_{2,G+1}, \ldots, F_{NP,G+1})$, and $CR_{G+1} = (CR_{1,G+1}, CR_{2,G+1}, \ldots, CR_{NP,G+1})$. Set $G = G + 1$, and go to Step 4.

### 3.3. Numerical Experiments

The SAHDE algorithm is tested by using 11 benchmark problems listed in Table 2. In the table, $n$ is the dimension of the problem, $S$ the search space, $S \subseteq \mathbb{R}^n$, and $f_{\min}$ represents the minimum value of

<table>
<thead>
<tr>
<th>Test function</th>
<th>$n$</th>
<th>$S$</th>
<th>$f_{\min}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$f_1(x) = \sum_{i=1}^n x_i^2$</td>
<td>30</td>
<td>$[-100,100]^n$</td>
<td>0</td>
</tr>
<tr>
<td>$f_2(x) = \sum_{i=1}^n</td>
<td>x_i</td>
<td>+ \prod_{i=1}^n</td>
<td>x_i</td>
</tr>
<tr>
<td>$f_3(x) = \sum_{i=1}^n (\sum_{j=1}^n x_j)^2$</td>
<td>30</td>
<td>$[-100,100]^n$</td>
<td>0</td>
</tr>
<tr>
<td>$f_4(x) = \max{</td>
<td>x_i</td>
<td>, 1 \leq i \leq n}$</td>
<td>30</td>
</tr>
<tr>
<td>$f_5(x) = \sum_{i=1}^{n-1} (100(x_{i+1} - x_i)^2 + (x_i - 1)^2$</td>
<td>30</td>
<td>$[-30,30]^n$</td>
<td>0</td>
</tr>
<tr>
<td>$f_6(x) = \sum_{i=1}^n (x_i + 0.5)^2$</td>
<td>30</td>
<td>$[-100,100]^n$</td>
<td>0</td>
</tr>
<tr>
<td>$f_7(x) = \sum_{i=1}^n { 0.5 + \cos(2\pi x_i) + \text{random}[0,1] }$</td>
<td>30</td>
<td>$[-1.28,1.28]^n$</td>
<td>0</td>
</tr>
<tr>
<td>$f_8(x) = \sum_{i=1}^n</td>
<td>x_i - \sin(\sqrt{</td>
<td>x_i</td>
<td>})</td>
</tr>
<tr>
<td>$f_9(x) = \sum_{i=1}^n</td>
<td>x_i - 10 \cos(2\pi x_i) + 10</td>
<td>$</td>
<td>30</td>
</tr>
<tr>
<td>$f_{10}(x) = -20 \exp(-0.2 \sqrt{\frac{1}{n} \sum_{i=1}^n x_i^2}) - \exp(\frac{1}{n} \sum_{i=1}^n \cos(2\pi x_i) + 20 + e)$</td>
<td>30</td>
<td>$[-32,32]^n$</td>
<td>0</td>
</tr>
<tr>
<td>$f_{11}(x) = \sum_{i=1}^{n} x_i^2 - \prod_{i=1}^{n} \cos \left( \frac{x_i}{\sqrt{2}} \right) + 1$</td>
<td>30</td>
<td>$[-600,600]^n$</td>
<td>0</td>
</tr>
</tbody>
</table>
the problem. In our experiments, the population size $NP$ is set to 100. Other parameters are suggested as: $G_t = 25$, $F_t = (0.1, 2)$, $CR_t = (0.1, 1)$, $b_0 = (0.4, 1)$ and $CR_b = (0.5, 0.95)$. We use the maximum generation number $G_{\text{max}}$ as the stopping criterion if $G > G_{\text{max}}$, then the algorithm is terminated. For each problem, 30 independent runs of an algorithm are tested. In order to verify the superiority of the algorithm, the results obtained by the standard DE algorithm, the jDE algorithm [18], the DESQI algorithm [22] are shown simultaneously.

Experimental results are given in Table 3. In the table, Mean and Std Dev represent the average function value found by the algorithm and the standard deviation of the found function values for 30 independent runs, respectively. As illustrated in Table 3, for problems $f_1, f_2, f_7, f_8$ and $f_{10}$, the SAHDE algorithm yields a smaller minimum objective function value. While for problems $f_5, f_6, f_9$ and $f_{11}$, the average function value found by the SAHDE algorithm is equal to that obtained by other algorithms. However, for problems $f_3$ and $f_4$, the average function value found by the SAHDE algorithm is a little bit larger than that obtained by the DESQI algorithm and the jDE algorithm. Therefore, the SAHDE algorithm yields relatively accurate solutions for all the problems, and performs favorably.

Table 3. Comparison results of average function values and the standard deviations for 11 benchmark problems.

<table>
<thead>
<tr>
<th>$f_i$</th>
<th>$G_{\text{max}}$</th>
<th>SAHDE Mean (Std Dev)</th>
<th>DESQI [22] Mean (Std Dev)</th>
<th>jDE [18] Mean (Std Dev)</th>
<th>DE/rand/1/bin Mean (Std Dev)</th>
</tr>
</thead>
<tbody>
<tr>
<td>$f_1$</td>
<td>1500</td>
<td>2.34E-45 (6.77E-45)</td>
<td>2.05E-23 (2.02E-23)</td>
<td>1.1E-28 (1.0E-28)</td>
<td>5.14E-14 (4.39E-14)</td>
</tr>
<tr>
<td>$f_2$</td>
<td>2000</td>
<td>3.39E-34 (3.39E-34)</td>
<td>7.92E-16 (3.31E-16)</td>
<td>1.0E-23 (9.7E-24)</td>
<td>3.75E-10 (1.96E-10)</td>
</tr>
<tr>
<td>$f_3$</td>
<td>5000</td>
<td>2.30E-12 (2.30e-11)</td>
<td>6.65E-18 (1.58E-17)</td>
<td>3.1E-14 (5.9E-14)</td>
<td>2.92E-11 (2.45E-11)</td>
</tr>
<tr>
<td>$f_4$</td>
<td>5000</td>
<td>4.86E-19 (1.75E-18)</td>
<td>2.17E-20 (3.07E-20)</td>
<td>00E+00 (0E+00)</td>
<td>1.62E-01 (4.29E-01)</td>
</tr>
<tr>
<td>$f_5$</td>
<td>2000</td>
<td>00E+00 (0E+00)</td>
<td>1.3E-01 (7.1E-01)</td>
<td>00E+00 (0E+00)</td>
<td>00E+00 (0E+00)</td>
</tr>
<tr>
<td>$f_6$</td>
<td>1500</td>
<td>00E+00 (0E+00)</td>
<td>00E+00 (0E+00)</td>
<td>00E+00 (0E+00)</td>
<td>00E+00 (0E+00)</td>
</tr>
<tr>
<td>$f_7$</td>
<td>3000</td>
<td>1.1E-03 (3.05E-04)</td>
<td>1.50E-03 (4.91E-04)</td>
<td>3.15E-03 (7.5E-04)</td>
<td>4.80E-03 (1.30E-03)</td>
</tr>
<tr>
<td>$f_8$</td>
<td>9000</td>
<td>9.125E+59 (1.81E-12)</td>
<td>4.1264.55 (242.34)</td>
<td>12569.5 (7.0E-12)</td>
<td>11890.53 (1447.78)</td>
</tr>
<tr>
<td>$f_9$</td>
<td>5000</td>
<td>00E+00 (0E+00)</td>
<td>00E+00 (0E+00)</td>
<td>00E+00 (0E+00)</td>
<td>7.29E+01 (3.08E+01)</td>
</tr>
<tr>
<td>$f_{10}$</td>
<td>1500</td>
<td>4.4E-15 (0E+00)</td>
<td>1.58E-12 (7.28E-13)</td>
<td>7.7E-15 (1.4E-15)</td>
<td>5.90E-08 (2.16E-08)</td>
</tr>
<tr>
<td>$f_{11}$</td>
<td>3000</td>
<td>00E+00 (0E+00)</td>
<td>00E+00 (0E+00)</td>
<td>00E+00 (0E+00)</td>
<td>2.46E-04 (1.3E-03)</td>
</tr>
</tbody>
</table>

In order to prove that the SAHDE algorithm is efficient for solving array synthesis problems, the SAHDE algorithm is used to minimize the fitness function (4) under constraint of Eq. (5). To compare the algorithm with other existing algorithms, a 37-element linear aperiodic array is considered, which has been synthesized by MGA [6], SaDE [11], and SHDE [10]. The array is also symmetric and with an aperture size is equal to $21.996\lambda$, where $\lambda$ is the wavelength. That means $d_m = 10.998\lambda$. Here, $d_{\text{min}}$ is set to $0.5\lambda$.

For the SAHDE algorithm, we choose $NP = 80$. Other parameters are chosen as suggested. The termination condition is that the maximum number of function evaluations reaches 40000. The optimal PSLLs and element positions are illustrated in Table 4. Compared with other algorithms, the SAHDE algorithm yields $-21.268$ dB PSLL, which is better than $-20.562$ dB by MGA [6], $-20.942$ dB by SaDE [11] and $-21.12$ dB by SHDE [10]. As shown in Fig. 3, the PSLLs obtained by the SAHDE algorithm for the array in 20 independent runs are stable. Comparison of the radiation patterns obtained by four algorithms for the array is also shown in Fig. 4. Simulated results indicate that the SAHDE algorithm yields better results than other existing algorithms in the same conditions. Results demonstrate the SAHDE algorithm is efficient for solving the array synthesis problem.

Since the self-adaptive parameter control technique is adopted, the SAHDE algorithm could find a set of proper control parameter values, making the algorithm converge fast and adapt itself to the problem being solved. The comparison results with some existing algorithms show that the self-adaptive parameter control technique could determine the appropriate control parameter values quickly, and that the performance of the SAHDE algorithm is efficient.
Table 4. PSLLs and element positions for the 37-element sparse linear array.

<table>
<thead>
<tr>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>D_1</td>
<td>0.5024λ</td>
<td>0.5000λ</td>
<td>0.5000λ</td>
<td>0.5002λ</td>
</tr>
<tr>
<td>D_2</td>
<td>1.0024λ</td>
<td>1.0000λ</td>
<td>1.0000λ</td>
<td>1.0004λ</td>
</tr>
<tr>
<td>D_3</td>
<td>1.5024λ</td>
<td>1.5000λ</td>
<td>1.5000λ</td>
<td>1.5008λ</td>
</tr>
<tr>
<td>D_4</td>
<td>2.0032λ</td>
<td>2.0000λ</td>
<td>2.0000λ</td>
<td>2.0008λ</td>
</tr>
<tr>
<td>D_5</td>
<td>2.5035λ</td>
<td>2.5000λ</td>
<td>2.5000λ</td>
<td>2.5009λ</td>
</tr>
<tr>
<td>D_6</td>
<td>3.0036λ</td>
<td>3.0000λ</td>
<td>3.0000λ</td>
<td>3.0016λ</td>
</tr>
<tr>
<td>D_7</td>
<td>3.5081λ</td>
<td>3.5000λ</td>
<td>3.5000λ</td>
<td>3.5034λ</td>
</tr>
<tr>
<td>D_8</td>
<td>4.0784λ</td>
<td>4.0000λ</td>
<td>4.0000λ</td>
<td>4.0575λ</td>
</tr>
<tr>
<td>D_9</td>
<td>4.6153λ</td>
<td>4.5000λ</td>
<td>4.5000λ</td>
<td>4.5742λ</td>
</tr>
<tr>
<td>D_{10}</td>
<td>5.1347λ</td>
<td>5.0510λ</td>
<td>5.0000λ</td>
<td>5.1236λ</td>
</tr>
<tr>
<td>D_{11}</td>
<td>5.7215λ</td>
<td>5.5780λ</td>
<td>5.6003λ</td>
<td>5.7760λ</td>
</tr>
<tr>
<td>D_{12}</td>
<td>6.2980λ</td>
<td>6.1620λ</td>
<td>6.1752λ</td>
<td>6.3461λ</td>
</tr>
<tr>
<td>D_{13}</td>
<td>7.0717λ</td>
<td>6.9040λ</td>
<td>6.7895λ</td>
<td>7.0343λ</td>
</tr>
<tr>
<td>D_{14}</td>
<td>7.7762λ</td>
<td>7.6640λ</td>
<td>7.6300λ</td>
<td>7.8534λ</td>
</tr>
<tr>
<td>D_{15}</td>
<td>8.7827λ</td>
<td>8.4880λ</td>
<td>8.3314λ</td>
<td>8.6103λ</td>
</tr>
<tr>
<td>D_{16}</td>
<td>9.6633λ</td>
<td>9.4880λ</td>
<td>9.6454λ</td>
<td>9.8212λ</td>
</tr>
<tr>
<td>D_{17}</td>
<td>10.4926λ</td>
<td>10.4880λ</td>
<td>10.4574λ</td>
<td>10.4949λ</td>
</tr>
<tr>
<td>D_{18}</td>
<td>10.9980λ</td>
<td>10.9880λ</td>
<td>10.9980λ</td>
<td>10.9980λ</td>
</tr>
</tbody>
</table>

Figure 3. PSLLs obtained by the SAHDE algorithm for the 37-element array in 20 independent runs.

Figure 4. Comparison of the radiation patterns obtained by four algorithms for the 37-element array.

4. WORST-CASE TOLERANCE SYNTHESIS OF LOW-SIDELOBE SPARSE LINEAR ARRAYS BY USING THE SAHDE ALGORITHM

In this section, for the low sidelobe sparse linear array with random element position errors, Problem (16) is solved by the SAHDE algorithm. The fitness function for Problem (16) is calculated by Eq. (17). Obviously, bigger NS leads to the more accurate results; however, bigger NS will increase the
computational burden greatly. In [14], Jiao et al. chose the vertices of the tolerance region as the sample point candidates for calculating the worst-case PSLLs of the linear arrays. However, Problem (16) is a non-convex problem, and the worst cases may not appear in the vertexes of the tolerance region. In order to solve this problem, we present a fast way for calculating the fitness function (17).

Although Problem (16) is a non-convex problem, the points around boundary of the tolerance region in Eq. (11) have greater contribution to the worst-case PSLLs of the sparse linear array, while the points far from its boundary have very little contribution. In order to calculate the fitness function (17) efficiently, we choose some sample points around boundary of the tolerance region in Eq. (11) randomly.

Since it is difficult to depict the multidimensional problem, the 2-dimensional problem is taken as an example to illustrate the sampling way. Fig. 5 shows schematic diagram of the sampling way. In the figure, \((d_1, d_2)\) denotes the nominal element position vector, and the dash-line box represents boundary of the tolerance region in Eq. (11). The sample points are chosen randomly around boundary of the region based on the Gaussian distribution. Define the Chebyshev distance between the sample point and the nominal element position vector as

\[
dch_j = \max \left( \left| \left( d_1 + dc_{j1} \right) - d_1 \right| , \left| \left( d_2 + dc_{j2} \right) - d_2 \right| \right) = \max \left( \left| dc_{j1} \right| , \left| dc_{j2} \right| \right)
\] (30)

The greater the Chebyshev distance is, the closer the sample point approaches the boundary. In the tolerance region, we recommend to choose some sample points with greater Chebyshev distances instead of choosing all the points. Only these sample points are used to calculate the fitness function (17). The detail procedure for calculating the fitness function is presented as follows.

**Step 0:** Set parameters: Set the fixed standard deviation \(\sigma_i\), the tolerance region for every element position \((-3\sigma_i, 3\sigma_i), i = -M, -(M - 1), \ldots, 0, \ldots, M - 1, M\), the total sample point number \(J\), and the used sample point number \(NS (NS < J)\) in Eq. (17).

**Step 1:** Produce error vectors: Produce \(JDc\) vectors in the tolerance region (11) based on Eq. (10).

<table>
<thead>
<tr>
<th>Table 5. Optimal element positions for the 37-element sparse linear array.</th>
</tr>
</thead>
<tbody>
<tr>
<td>(3\sigma_i = 0.01\lambda)</td>
</tr>
<tr>
<td>(d_1)</td>
</tr>
<tr>
<td>(d_2)</td>
</tr>
<tr>
<td>(d_3)</td>
</tr>
<tr>
<td>(d_4)</td>
</tr>
<tr>
<td>(d_5)</td>
</tr>
<tr>
<td>(d_6)</td>
</tr>
<tr>
<td>(d_7)</td>
</tr>
<tr>
<td>(d_8)</td>
</tr>
<tr>
<td>(d_9)</td>
</tr>
<tr>
<td>(d_{10})</td>
</tr>
<tr>
<td>(d_{11})</td>
</tr>
<tr>
<td>(d_{12})</td>
</tr>
<tr>
<td>(d_{13})</td>
</tr>
<tr>
<td>(d_{14})</td>
</tr>
<tr>
<td>(d_{15})</td>
</tr>
<tr>
<td>(d_{16})</td>
</tr>
<tr>
<td>(d_{17})</td>
</tr>
<tr>
<td>(d_{18})</td>
</tr>
</tbody>
</table>
**Step 2:** Calculate the Chebyshev distance: Calculate the Chebyshev distance between every $\mathbf{Dc}$ and the nominal element position vector by Eq. (30), sort the distances in the descending order, and choose $NS$ sample points with larger Chebyshev distances.

**Step 3:** Evaluate the fitness function: The $NS$ sample points are used to calculate the fitness function value by Eq. (17).

Parameters $J$ and $NS$ are set as 50000 and 2500, respectively. The SAHDE algorithm is used to solve Problem (16). For its control parameters and the termination condition, refer to Section 3.

When the tolerance regions are fixed to three levels $3\sigma_i = 0.01\lambda$, $0.05\lambda$, $0.1\lambda$, the optimal element positions are given in Table 5, and the optimal PSLLs are shown in Table 6. The worst PSLLs are

<table>
<thead>
<tr>
<th>Tolerance level</th>
<th>Without tolerances</th>
<th>With tolerances</th>
</tr>
</thead>
<tbody>
<tr>
<td>$3\sigma_i = 0.01\lambda$</td>
<td>$-21.268$</td>
<td>$-20.477$</td>
</tr>
<tr>
<td>$3\sigma_i = 0.05\lambda$</td>
<td>$-17.734$</td>
<td>$-20.56$</td>
</tr>
<tr>
<td>$3\sigma_i = 0.1\lambda$</td>
<td>$-15.011$</td>
<td>$-19.04$</td>
</tr>
</tbody>
</table>

Table 6. Optimal PSLLs for the 37-element sparse linear array.

**Figure 5.** Schematic diagram of the sampling way.
Figure 6. Comparison of the radiation patterns for the 37-element sparse linear array with three fixed position tolerance levels. (a) $3\sigma_i = 0.01\lambda$. (b) $3\sigma_i = 0.05\lambda$. (c) $3\sigma_i = 0.1\lambda$.

$-20.84\text{ dB}, -20.09\text{ dB}$ and $-17.12\text{ dB}$, and the nominal PSLLs are $-20.92\text{ dB}, -20.56\text{ dB}$ and $-19.04\text{ dB}$. As comparison, without considering the element position errors, the nominal PSLL is $-21.268\text{ dB}$, as shown in Table 4. Analysis results show that the worst PSLLs are $-20.477\text{ dB}, -17.734\text{ dB}$ and $-15.011\text{ dB}$, corresponding to three tolerance levels. Although the nominal PSLLs $-20.92\text{ dB}, -20.56\text{ dB}$ and $-19.04\text{ dB}$ obtained by the tolerance optimization are higher than $-21.268\text{ dB}$ without considering the errors, the worst-case PSLLs are improved from $-20.477\text{ dB}$ to $-20.84\text{ dB}$, $-17.734\text{ dB}$ to $-20.09\text{ dB}$ and $-15.011\text{ dB}$ to $-17.12\text{ dB}$, as shown in Table 6. Comparison of the radiation patterns considering the errors are also shown in Fig. 6. The compared results show that the worst-case peak sidelobe levels for the sparse linear arrays are improved evidently.

5. CONCLUSION

In this paper, a worst-case tolerance synthesis for low-sidelobe sparse linear arrays with random position errors has been studied. First, the worst-case tolerance synthesis model considering random position errors is established. Through the random sampling, the random optimization problem is converted to a deterministic optimization problem. In order to solve the problem, a novel self-adaptive hybrid differential evolution algorithm is proposed then, in which a novel self-adaptive parameter control technique is adopted. The control parameters have been produced by the simplified quadratic interpolation operator, and the connections between control parameters and fitness values available are established. In order to determine the appropriate control parameter values quickly, a selection operation for control parameters is also used. 11 benchmark problems have been used for verifying the effectiveness of the algorithm. Moreover, a sparse linear array is synthesized by using the SAHDE algorithm. Simulated results demonstrate that the SAHDE algorithm is efficient not only for global optimization problems, but also for synthesizing low sidelobe sparse antenna arrays. Finally, the worst-case tolerance synthesis problems with three fixed position tolerance levels are solved by the SAHDE algorithm. Simulated results illustrate that the worst PSLLs have been improved evidently. The SAHDE algorithm can also be used to solve the worst-case tolerance synthesis for other antenna arrays.

REFERENCES


