Application of Differential Evolution Algorithm in Prediction of Time Series Data

Satyendra Nath Mandal¹, Jit Saha²

Assistant Professor, Department of Information Technology, Kalyani Govt. Engg. College, Kalyani, Nadia, India¹

M.Tech Final Year, Department of Computer Science & Engineering, Kalyani Govt. Engg. College, Kalyani, Nadia, India²

ABSTRACT: Several statistical and soft-computing methods have been applied to predict the time series data in different fields. In this paper, an algorithm has been proposed to predict the time series data based on Differential Evolution algorithm. The performance of proposed algorithm has been measured by Mean Absolute Percentage Error (MAPE) metric. It has been observed that in most of the cases the performance of the proposed algorithm is better compare to other applied methods in same data set. Finally, the algorithm has been tested on several time series data predicted by the other methods.

KEYWORDS: Statistical and soft-computing methods, Differential Evolution, Time Series data, Mean Absolute Percentage Error (MAPE).

I. INTRODUCTION

In recent time several evolutionary algorithms have been applied to different real life optimization problems and the results given by them is satisfactory. Differential Evolution (DE) is one of the latest evolutionary based optimization algorithms, proposed by Storn& Price. Like other evolutionary-type algorithms, DE is population based algorithm and it represents the candidate solutions by chromosomes based on real numbers. In recent times this algorithm has been applied in several optimization problems from different field of science.

PuspanjaliMohapatra and Alok Raj have applied DE on Indian stock market prediction. They have compared performance of two algorithms differential evolution (DE) and Back propagation (BP) for training a Functional Link Artificial Neural Network (FLANN) based ANN to get optimized value of weights of underlined neural network. The DE optimized FLANN has been always performed well than BP optimized FLANN in terms of MAPE & RMSE.

Juan Peralta Donate, XiaodongLi,Germa’nGutie´rrezSa´nchez and AraceliSanchis de Miguel have applied DE to training automatic design of artificial neural networks (ADANN) for predicting time series data. The performance of DE has been compared with other two algorithms- GA, EDA (estimation of distribution algorithms) and famous time series prediction software Forecast Pro for predicting passenger, temperature, Dow-Jones and Quebec and Mackey–Glass data set. Results have revealed that for Mackey–Glass data set DE optimized ANN has outperformed over Forecast Pro software.

YuelinGao and Junmei Liu have proposed a new version of DE, Multi objective Differential Evolution Algorithm with Multiple Trial Vectors (MTVDE) for finding optimal values of different multi objective optimization function. To evaluate the performance of the proposed algorithm it has been applied in several benchmark functions along with another evolutionary based algorithm and most of the cases MTVDE has performed well than other multi-objective based evolutionary algorithm (MOEA).

Ching-Wei Chien, Zhan-Rong Hsu and Wei-Ping Lee have proposed Modified mutation Factor Differential Evolution (MFDE). They have adopted a new mutation factor which has been changed dynamically taking consideration of exploration-exploitation property of algorithm. Proposed method has been applied in some multimodal and single modal functions along with other standard versions of DE.MFDE have performed well in multimodal function and high-dimensional function.
In this paper, DE algorithm has been applied for time series prediction. Here at first initial populations of chromosomes have been created randomly in the search space. Here each chromosome is a D-dimensional vector, where D is the number of input data. The initial populations of chromosomes have been created randomly in search space. Each chromosome represents the predicted data of input. After that, initial fitness of each chromosome has been calculated. Next, at each iteration or each generation of the algorithm mutation, cross-over and selection, these three steps have been performed for every chromosome of the population. The whole procedure has been carried out for 1000 iterations. At the end of 1000th iteration, chromosome having the best fitness has been considered as the predicted values of the input data.

II.THEORY

A. DE OVERVIEW

The DE algorithm belongs to family of evolutionary algorithm. It uses the same operators- mutation, cross-over and selection like GA. But it has several differences with GA. Unlike GA the decision variables are real numbers and mutation step of DE is completely different than mutation step of GA. For constructing solution, GA depends on crossover while DE depends on mutation operation [9]. This mutation operator is a unique tool of DE for attacking any optimization based problem.

B. PARAMETERS AND TERMINOLOGY OF BASIC DE

B.1. Chromosome

In DE each element of the population is called as chromosome which is a concept of any evolutionary based algorithm. Each chromosome is represented as D-dimensional vector.

B.2. Scaling factor or mutation factor (F)

In mutation operator, F is used to amplifying the difference of two chromosomes. It is a very important parameter of mutation operator. Value of this parameter vastly affects the performance of the algorithm. Value may be fixed or it can be adaptive during the execution of the algorithm. The main job of this parameter is to balance exploration and exploitation property of the algorithm. In initial stage it finds much more global optimal solutions; and has strong local search ability in later stages [6]. Generally value of F ε (0, 2] [3]. In our proposed method tuning of the F value has been guided by the following formula,

\[ F_{\text{iter}} = \frac{F_{\text{initial}}}{\text{itr}} \]  

where \( F_{\text{initial}} \) is the initial value of mutation factor (F) and \( \text{itr} \) is the current iteration number and \( F_{\text{iter}} \) is the value of the mutation factor at current iteration t.
B.3. Mutation Operator
In this step for each chromosome $X^G_i$, a mutant chromosome, $V^{G+1}_i$ is produced from three parent chromosomes of the current generation or iteration $G$. The simple mutation operator is defined as,

$$V^{G+1}_i = X^G_i + F(X^G_j - X^G_k).$$

Where $(G+1)$ represents next generation, $X^G_i$, $X^G_j$ and $X^G_k$ are three parent chromosome of the current generation. $X^G_i$, $X^G_j$ and $X^G_k$ are differ from each other. $F$ is scaling factor and $X^G_i$ is also called as donor vector.

B.4. Crossover Operator
In this step for each chromosome $X^G_i$, a trial chromosome or child chromosome $U^{G+1}_i$ is produced by the cross-over between parent chromosome $X^G_i$ and mutant chromosome $V^{G+1}_i$. The simple cross-over operator is binomial cross over and it is defined as,

$$U^{G+1}_i = \begin{cases} 
V^{G+1}_i, & \text{if (rand<CR).} \\
X^G_i, & \text{otherwise.}
\end{cases}$$

Where CR is crossover ratio and $CR \in [0,1]$. rand is random number, generated in between 0 and 1.

B.5. Selection Operator
In this step for each chromosome $X^G_i$, fitness comparison is performed between child chromosome, $U^{G+1}_i$ and parent chromosome $X^G_i$. During fitness comparison if parent, $X^G_i$ is superior to child, it remains in the search space, otherwise parent will be replaced by child, $U^{G+1}_i$. In case of maximization problem fitness function is proportional to the objective function value.

This can be defined as,

$$X^{G+1}_i = \begin{cases} 
U^{G+1}_i, & \text{if (fitness (U^{G+1}_i)) > fitness (X^{G+1}_i))}. \\
X^{G+1}_i, & \text{Otherwise.}
\end{cases}$$

III. VARIATION OF DE

Many variants of the basic DE have been proposed by Price & Storn (2003) based on following criteria-donor vectors used in mutation operator, number of vector difference used in mutation operator, type of cross over operator used in cross-over operation[10]. In order to distinguish among these variants, DE/x/y/z naming convention has been used, where $x$ denotes the donor vector during mutation operator, $y$ denotes the number of vector difference in mutation operator and $z$ denotes the type of crossover used.

Those variation are:
1. DE/best/1/exp
2. DE/rand/1/exp
3. DE/rand-to-best/1/exp
4. DE/best/2/exp
5. DE/rand/2/exp
6. DE/best/1/bin
7. DE/rand/1/bin
8. DE/rand-to-best/1/bin
9. DE/best/2/bin
10. DE/rand/2/bin

In this paper 7th variation is used for time series prediction problem.
IV. TIME SERIES FORECASTING

Quantities that represent the values have been taken by a variable over a period such as a month, quarter, or year. Time series data occurs wherever the same measurements are recorded on a regular basis. Time series data is a series of statistical data that is related to a specific instant or a specific time period. It is drawn from a data of monthly bookings for an airline in shown in Fig1.

Fig 1. Monthly bookings for an airline

V. PROPOSED ON BASED ON DIFFERENTIAL EVOLUTION ALGORITHM

Input: Time series data, maximum iteration number, population size per data element.

Output: Predicted data.

Method:
Step 1:
Step 1.1: To Compute upper bound (UB) and lower bound (LB) of the population from the given time series
Step 1.2: To initialize the parameters F, CR, population size of the algorithm.

Step 2: To generate population in search space between upper and lower bound. It has been calculated by the formula, 
\[ X_{i,j} = LB + (UB - LB) \times \text{ran}(0, 1). \]
Where \( X_{ij} \) is the j\textsuperscript{th} predicted data element of chromosome i.

Step 3: To compute initial fitness of each chromosome \( X_i \). It has been calculated by the formula,
\[ F_{Xi} = \sum_{j=1}^{D} \left| X_{i,j} - \text{data}_j \right|, \]
where \( data_j \) is the j\textsuperscript{th} input data element of the input and D is the number of input data. Here lower value of \( F_{Xi} \) indicates higher fitness value.

Step 4: To repeat step 5 to step 7.

Step 5: To calculate mutation factor for the current iteration using the formula,
\[ F_{\text{inv}} = \frac{F_{\text{initial}}}{\text{itr}}. \]

Step 6: for each chromosome \( X_i \) in the search space perform step 6.1 to 6.3
  
  Step 6.1: mutation
  Step 6.2: cross-over
  Step 6.3: selection

Step 7: if the termination condition is satisfied return the chromosome with best fitness as predicted value of the input data.

Step 8: Otherwise go to Step 7.

**V. EXPERIMENTAL SET-UP**

The proposed method has been applied on Fedora Core 15.i686 platform with CORE i5 processor and 8GB RAM. The coding has been done by C language.

Parameter value:
- Number of population (NP) = 30*number of data input.
- Mutation factor \((F)\) = Initial value is 0.9.
- Cross-over ratio \((CR)\) = 0.9.
- Maximum number of iteration = 1000.

The whole procedure has been done for 10 trials with different initial population and finally mean prediction has been calculated for all data from prediction results of 10 trials and it has been given as final prediction of input.

**TABLE 6. THE PERFORMANCE COMPARISON OF THE PROPOSED METHOD WITH OTHER PREDICTION METHOD IN DIFFERENT TIME SERIES DATA FROM DIFFERENT JOURNALS.**

<table>
<thead>
<tr>
<th>Serial Number</th>
<th>Data Set</th>
<th>Journal Name</th>
<th>Method used in journal</th>
<th>Error in Journal's method</th>
<th>MAPE in DE</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Real estate prices of detached houses</td>
<td><em>(Application of Fuzzy Neural Network for Real Estate Prediction)</em> J. Wang et al. ISNN 2006, LNCS 3973, pp. 1187 – 1191, 2006</td>
<td>Fuzzy neural network</td>
<td>0.894838%</td>
<td>0.074196%</td>
</tr>
<tr>
<td>Page</td>
<td>Title</td>
<td>Author(s)</td>
<td>Methodology</td>
<td>Accuracy</td>
<td>Error</td>
</tr>
<tr>
<td>------</td>
<td>----------------------------------------------------------------------</td>
<td>---------------------------------------------------------------------------</td>
<td>-----------------------------------------------------</td>
<td>----------</td>
<td>-------</td>
</tr>
<tr>
<td>3</td>
<td>LCD TV demand</td>
<td>Ruey-Chyn Tsaur. SoftComput</td>
<td>Fuzzy grey regression model</td>
<td>3.071154%</td>
<td>0.042545%</td>
</tr>
<tr>
<td>5</td>
<td>Forecasting engine system reliability</td>
<td>K. Xu et al. Engineering Driver 2, 117576</td>
<td>RBF neural network for time series prediction</td>
<td>0.419024%</td>
<td>0.021231%</td>
</tr>
<tr>
<td>6</td>
<td>Education capital in China</td>
<td>H. Guo et al. China University of Geosciences, 430074</td>
<td>Fuzzy Neural Network algorithm FNN2</td>
<td>41.555765%</td>
<td>2.457342%</td>
</tr>
</tbody>
</table>
### International Journal of Innovative Research in Science, Engineering and Technology

*(An ISO 3297: 2007 Certified Organization)*

**Vol. 5, Special Issue 13, October 2016**

<table>
<thead>
<tr>
<th>Page</th>
<th>Title</th>
<th>Authors</th>
<th>Algorithm</th>
<th>MSE</th>
<th>RMSE</th>
</tr>
</thead>
<tbody>
<tr>
<td>8</td>
<td><strong>Initial shoot length of Ken type mustard plant</strong></td>
<td>Satyendra Nath Mandal, J. Pal Choudhury, Dilip De &amp; S. R. Bhadrachaudhuri</td>
<td>Fuzzy soft computing model</td>
<td>0.8527%</td>
<td>0.168998 %</td>
</tr>
<tr>
<td>9</td>
<td><strong>Tourism Demand in Taiwan</strong></td>
<td>Chang-Jui Lin, Hsueh-Fang Chen, Tian-Shyug Lee</td>
<td>ARIMA</td>
<td>0.28%</td>
<td>0.050054 %</td>
</tr>
</tbody>
</table>

### VII. CONCLUSION & FUTURE WORK

It has been observed that the proposed algorithm has given better output as compare to others. The algorithm has been also tested on different time series data published in different International journals. The values of parameters of DE have been collected from different published journals. A multithreaded version of this method may be proposed in future to explore the search space efficiently and quickly and a more efficient mutation operator may be proposed for saving the algorithm from premature convergence for finding global optimum value of the underlying problem.

### REFERENCES


---

Copyright to IJIRSET www.ijirset.com 316


