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COVID-19 X-ray images classification based on enhanced fractional-order cuckoo search optimizer using heavy-tailed distributions

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ABSTRACT

Classification of COVID-19 X-ray images to determine the patient’s health condition is a critical issue these days since X-ray images provide more information about the patient's lung status. To determine the COVID-19 case from other normal and abnormal cases, this work proposes an alternative method that extracted the informative features from X-ray images, leveraging on a new feature selection method to determine the relevant features. As such, an enhanced cuckoo search optimization algorithm (CS) is proposed using fractional-order calculus (FO) and four different heavy-tailed distributions in place of the Lévy flight to strengthen the algorithm performance during dealing with COVID-19 multi-class classification optimization task. The classification process includes three classes, called normal patients, COVID-19 infected patients, and pneumonia patients. The distributions used are Mittag-Leffler distribution, Cauchy distribution, Pareto distribution, and Weibull distribution. The proposed FO-CS variants have been validated with eighteen UCI data-sets as the first series of experiments. For the second series of experiments, two data-sets for COVID-19 X-ray images are considered. The proposed approach results have been compared with well-regarded optimization algorithms. The outcomes assess the superiority of the proposed approach for providing accurate results for UCI and COVID-19 data-sets with remarkable improvements in the convergence curves, especially with applying Weibull distribution instead of Lévy flight.

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1. Introduction

The first cases of coronavirus disease (COVID-19) were registered in Wuhan, an important city of China, in December 2019. COVID-19 is originated by a virus called SARS-CoV-2, and currently, it is one of the major concerns worldwide. By the middle of April 2020, the affections of COVID-19 in humans can be summarized in more than 150,000 deaths and almost 1,700,000 confirmed cases worldwide [1]. These amounts are evidence of the fast dissemination in the population. The symptoms of COVID-19 known until now include fever, cough, sore throat, headache, fatigue, muscle pain, among others [2]. The test for detection of COVID-19 commonly used is invasive, and it is called a reverse transcription-polymerase chain reaction (RT-PCR) [3]. COVID-19 is not only affecting the health of the nations but also the consequences of the diseases are important (e.g., economic and psychological [4,5]). Another important consideration is that prompt detection could be reflected in early treatment. Then COVID-19 is a pandemic whit a large number of challenges to face. Based on the above information, they are required tools that permit the detection of this mortal disease in a timely manner.

On the other hand, the use of medical images to diagnose disease has increased in the latest years. Different computer vision and image processing tools can be used to identify the abnormalities produced by the illnesses. One of the main advantages of using this kind of system is that the detection is fast and accurate. However, the results must be validated by an expert. In
this way, medical image processing algorithms can be used as a primary diagnosis that provides a clue about a possible disease. In the case of COVID-19, the use of X-ray images permits to study how the virus is affecting the lungs. Since COVID-19 is a recent virus, only a few data-sets are available, and the number of works related to them is also reduced.

A number of studies have been proposed to classify COVID-19 from X-ray scan images using different techniques, such as DenseNet201 [6], MobileNet v2 [7], DarkNet [8], and others [9–12]. In general, the medical images can be directly analyzed to identify certain elements in the scene that permits the diagnosis. However, it is also common to extract different features from all the images and create a set that contained the information of the objects contained in the images [13]. In this case, not all the features extracted could provide essential information about the disease that is going to be detected; for that reason, it is necessary to use automatic tools that remove the non-desired (irrelevant or duplicated) elements.

Feature Selection (FS) is an important pre-processing tool that helps to extract the information desired from the irrelevant data [14–16]. FS helps in reducing the dimensionality of the data by selecting only the relevant information in the next steps. With this pre-processing, the machine learning could be applied more efficiently over the data-set by decreasing computational effort [17,18]. The use of FS has been extended for different domains, for example in bio-medicine for diagnosis of neuromuscular disorders [19], in signal processing for speech emotion recognition [20], in internet of things specially for medicine with EEG signals [21], human activity recognition [22] and text classification [23], to mention some. The FS works by maximizing the relevance of the features, but also the redundancy could be minimized. In comparison to other methods, FS preserves the data by not applying domain transformations; then, the FS operates by including and excluding the attributes in the data-set without any modification. The FS has three main advantages, (1) it improves the prediction performance, (2) it permits a better understanding of the processes that fastly generate the data, and (3) it has more effective predictors with lower computational cost [24]. According to Pinheiro et al. [25], they exist two kinds of FS method, namely the wrapper and filter methods. The most popular approaches are the wrapper-based because they provide more accurate results [26]. The wrapper FS method needs an internal classifier to find a better subset of features; this could affect their performance, especially with large data-sets. Moreover, they also have backward and forward strategies to add or remove features that become non-suited for a larger amount of features. Considering these drawbacks, meta-heuristic algorithms (MA) are used to increase the performance of FS processes.

MA have become more popular in recent years due to their flexibility and adaptability [27]. MA can be used in a wide range of applications, and FS is not an exception. Some classical approaches Genetic Algorithms (GA) [28], Particle Swarm Optimization (PSO) [29] and Differential Evolution (DE) [30] have been successfully used to solve the FS problem. More so, modern MA algorithms also have been applied for FS, such as Grasshopper Optimization Algorithm (GOA) [31], Competitive Swarm Optimizer (CSO) [32], Gravitational Search Algorithm (GSA) [33], and others. Since FS can be seen as an optimization problem, there is no MA able to handle all the difficulties in FS. This last fact is defined according to the No-Free-Lunch (NFL) theorem [34]; therefore, it is necessary to continue exploring new alternative MA.

An interesting MA is the Cuckoo Search (CS) algorithm introduced by Yang and Deb [35,36] as a global optimization method inspired by the breeding behavior of Cuckoo birds (CB). CS is a population-based method that employs three basic rules that mimic the brood parasitism of the Cuckoo bird. In nature, the female CB lays the eggs in the nest of other species. Some CB species can imitate the color of the eggs of the nest where they will be laid. In the CS context, a set of host nest containing eggs (candidate solutions) are randomly initialized, then only one Cuckoo egg is generated in each generation using Lévy Flights. After that, a random host nest is chosen, and its egg is evaluated using a fitness function. The fitness value of the Cuckoo egg and the one selected from a random host nest are compared, the best egg is preserved in the host nest. A fraction of the worst nest is abandoned in each generation, and the best solutions are preserved [35]. The use of CS for several applications has been increased in recent years [37,38]. The CS is an interesting alternative for solving complex optimization problems. However, the CS has the disadvantage that it can fall into sub-optimal solutions [39]; this is caused due to inappropriate balance between exploration and exploitation phases. To overcome such drawbacks, they have been introduced some modifications of the standard CS.

Recently, an interesting approach called fractional-order cuckoo search (FO-CS) is proposed to include the use of fractional calculus to enhance the performance of the CS [40]. The FO-CS possesses better convergence speed than the CS; besides, the balance between exploration and exploitation produced by the fractional calculus permits more accurate solutions to complex optimization problems. Nevertheless, several other approaches can be included with FO-CS for further performance enhancement.

This paper presents the use of heavy-tailed distributions instead of Lévy flights in the FO-CS. This kind of distribution has been used to enhance the mutation operator in evolutionary algorithms and other MA [41,42]. Using the heavy-tailed distributions, it is possible to escape from non-prominent regions of the search space [41]. The distributions used in to enhance the FO-CS are the Mittag-Leffler distribution (MLD) [43], the Pareto distribution (Pd) [44], the Cauchy distribution (Cd) [45], and the Weibull distribution (Wd) [46]. The FO-CS based on heavy-tailed distributions is proposed as an alternative method for solving features selection problems. In this context, the experimental results include different tests considering eighteen data-sets from the UCI repository [47]. The comparisons included different MA from the state-of-the-art, where the proposed approach provides better results in terms of accuracy and convergence. Moreover, the most important contribution of this work is the use of the FO-CS based on heavy-tailed distributions for FS over a COVID-19 data-set. Over this experiment, the results obtained by the proposed method permits to accurately identify three different classes, namely normal patients, COVID-19 infected patients, and Pneumonia patients. Regarding the results of UCI and COVID-19 data-sets, different statistical and metrics validate the good performance of the FO-CS based on heavy-tailed distributions, especially Weibull distribution. The main contribution of the current study can be summarized as follows:

1. Provide an alternative COVID-19 X-ray image classification method which aims to detect the COVID-19 patient from normal and abnormal cases.
2. Extracting features from X-ray images then using a new feature selection method to select the relevant features.
3. Develop feature selection method using a modified CS based on the fractional concept and heavy-tailed distributions.
4. Evaluate the developed method using two data-set for real COVID-19 X-ray images.
5. Compare the new developed FS method with other recently implemented FS methods.

The rest of the paper is organized as follows. Section 2 presents the related works. Section 3 introduces the basics of heavy-tailed
distributions, CS, and FO-CS. Section 4 explains the proposed modified FO-CS, in Section 5 presents the experimental results and comparisons. Finally, in Section 6, the conclusions and future work are discussed.

2. Related works

In this section, we present a simple review of the existing works of COVID-19 medical image classification, the applications of MA for feature selection, and the recent applications of the CS.

Recently, some studies have been presented to classify and identify COVID-19 from chest X-ray images using different techniques. Pereira et al. [48] applied CNN to extract texture features from chest X-ray images. They applied resampling techniques to balance the distribution of the classes to implement a multi-classification process. Ozturk et al. [8] applied a deep learning-based approach for determining COVID-19 for X-ray scan images. They used DarkNet model to classify images into binary classification and multi-classification. For binary classification, images were classified into COVID or non-findings. For the multi-classification, images were classified into three classes, COVID, pneumonia, and no-findings. Elaziz et al. [49] proposed a machine learning-based model to classify COVID-19 X-ray images into two classes, COVID or non-COVID. They proposed a feature extraction method called fractional multichannel exponent moments to extract features from X-ray scan images. Thereafter, they applied an improved Manta-Ray Foraging Optimization algorithm as a feature selection method to select the most relevant features. In [9], the authors evaluated several convolutional neural network models with different X-ray scan images that include, COVID-19, normal incidents and bacterial pneumonia. Ouchicha et al. [50] proposed a deep learning-based approach for COVID-19 classification, called CVDNet, which is built by using the residual neural network. The evaluation outcomes showed that CVDNet could achieve a high detection accuracy on a small dataset. In [51], the authors proposed a deep learning approach, namely COVIDX-Net for COVID-19 diagnosis from chest X-ray scan images. COVIDX-Net depends on seven deep learning architectures. It was evaluated with 50 X-ray images, and it showed a good performance. In [52], a new model called CoroNet is proposed for COVID-19 detection from both of CT and X-ray images. Xception architecture is used to build CoroNet, and ImageNet datasets are applied to train it. It showed an acceptable performance for several classes with an average accuracy of 89.6 Toraman et al. [53] presented a Convolutional CapsNet model for COVID-19 detection from X-ray images. The proposed model was applied for binary classification (COVID-19, and None-COVID) and also for multi-class classification. Furthermore, Jain et al. [54] applied a deep learning model based on ResNet-101 for COVID-19 X-ray image classification. The proposed model achieved significant performance including several performance measures. Hassantabar et al. [55] applied deep neural network (DNN) and convolutional neural network (CNN) to detect infected lung tissue of COVID-19 patients form X-ray scans. Furthermore, there are different approaches for COVID-19 X-ray images classification, such as MobileNet v2 [7], DenseNet201 [8], and others [10–12]. In recent years, the modern MA have been widely applied for FS, for example, in [56] it has been proposed the use of the Grasshopper Optimization Algorithm (GOA) to identify the best subset of features, the use of Crow Search Algorithm (CSA) with chaotic maps has also been proposed for FS using different benchmark data-sets [31]. Another interesting work includes the implementation of the Competitive Swarm Optimizer (CSO) for FS [32]; the advantage of CSO is its capability to handle high-dimension optimization problems. In [33], an evolutionary version of the Gravitational Search Algorithm (GSA) is introduced for FS, and the use of mutation and crossover operators enhance the performance of the standard GSA. Besides using the hybrid Particle Swarm Optimization (PSO) to extract the optimal features of different benchmark data-sets. Too and Mirjalili [57] proposed an FS approach based on a hyper learning binary dragonfly algorithm. They applied the proposed approach for different FS datasets, including the COVID-19 patient health prediction dataset. All these approaches are novel, and their performance is relevant. However, they have some disadvantages; the first is that, like many other meta-heuristics, they can fall into sub-optimal solutions; the second is related to their use only in benchmark sets. Finally, since FS can be seen as an optimization problem, there is not MA able to handle all the difficulties in FS. This last fact is defined according to the No-Free-Lunch (NFL) theorem [34]; therefore, it is necessary to continue exploring new alternative MA.

The cuckoo search (CS) is one of the algorithms that have been applied for several applications has been increased in recent years [37,38]. The CS is an interesting alternative for solving complex optimization problems. However, the CS has the disadvantage that it can fall into sub-optimal solutions [39]; this is caused due to inappropriate balance between exploration and exploitation phases. To overcome such drawbacks, they have been introduced some modifications of the standard CS. For example, in [58], the authors proposed the hybridization of CS with the Chaotic Flower Pollination algorithm (CFPA) for maximizing area coverage in Wireless Sensor Networks. The Reinforced Cuckoo Search Algorithm (RCSA) is presented by Thirugnanasambandam et al. [59] for multi-modal optimization, the RCSA contains three operators called modified selection strategy, Patron-Prophet concept, and self-adaptive strategy. An enhanced version of the CS that includes the quasi-opposition based learning and other two strategies is introduced in [60] for the parameter estimation of photovoltaic models. Another recent approach of fractional-order cuckoo search (FO-CS) including the use of fractional calculus was proposed for identifying the unknown parameters of the chaotic, hyper-chaotic fractional-order financial systems [40]. The proposed FO-CS exposed its efficiency in solving set of known mathematical benchmarks as well with a remarkable convergence speed in comparison with the basic CS.

3. Background

In this section, the essential knowledge about the features extraction and feature selection approaches from images are explained in detail. Moreover, we introduce an overall background about the integrated parts in the novel optimizer variants that are the heavy-tailed distributions and the novel fractional-order cuckoo search optimizer (FO-CS) as described below:

3.1. Image features extraction techniques

In this study, the image features are extracted using the following techniques, the Fractional Zernike Moments (FrZMs), Wavelet Transform (WT), Gabor Wavelet Transform (GW), and Gray Level Co-Occurrence Matrix (GLCM). Brief descriptions of these techniques are listed below.

3.1.1. Fractional Zernike Moments (FrZMs)

The FrZMs are used to extract gray images features as presented by [61]. This process is shown in Eq. (1) [62].

$$\text{FrZM}_{\alpha,n,m}(f) = \int_0^{\frac{2\pi}{\alpha}} \int_0^{2\pi} \text{FrR}_{\alpha,n,m}(r) g(r, \theta) r^{\alpha n/m} \, d\theta \, dr$$  \hspace{1cm} (1)$$

where $\alpha$ denotes the fractional parameter, $\alpha \in \mathbb{R}^+$, $g(r, \theta)$ is a gray image. Fr$R_{\alpha,n,m}(r)$ denotes the real valued radial polynomial
as, and it is calculated by Eq. (2). n and m define the order which \(|m| \leq n\) and \(n - |m|\) be even, \(j^2 = -1\).

\[ F_{R}(\omega, m)(r) = \frac{n + 1}{\pi} \sqrt{\alpha} \sum_{k=0}^{n-|m|/2} (-1)^{(n-k)/2} k \sum_{l=0}^{r^{k(n-2k-1)-1} 0 < r \leq 1} \]  

The calculation of FrZMs need a linear mapping transformation of image coordinates to a proper space inside a unit circle because they are determined in polar coordinates \((r, \theta)\) where \(|r| \leq 1\) [63]. This step is calculated by Eq. (3).

\[ r_{x,y} = \sqrt{c_{1}+c_{2}^{2}} + (c_{1}y+c_{2})^{2} \]  

\[ \theta_{x,y} = \tan^{-1} (\frac{c_{1}y+c_{2}}{c_{1}x+c_{2}}) \]  

where \(c_{1} = \sqrt{2}/(N - 1), c_{2} = -1/\sqrt{2}\). N defines the pixels number in the image.

3.1.2 Wavelet Transform (WT)

The WT is a method used for signal analysis and feature extraction [64]. It breaks up a signal into scaled and shifted versions of the mother wavelet. It is represented by the sum of the signal multiplied by the shifted and scaled versions of the wavelet function \((\psi)\). The continuous WT (CWT) of a signal \(x(t)\) can be computed as:

\[ CWT(x, b) = \int_{R} x(t) \psi_{x,b}(t) dt \]  

\[ \psi_{x,b}(t) = \frac{1}{\sqrt{a}} \psi(t - b \frac{a}{b}) \]  

where \(\psi\) denotes the mother wavelet, \(b, a \in \mathbb{R}\) and represent the shifted and scaled parameters, respectively. \(a\) is not equal to zero.

3.1.3 Gabor Wavelet Transform (GW)

The GW is a popular technique used filters for extracting image features [65]. It is calculated by Eq. (7).

\[ \psi_{\mu, v}(z) = \frac{|k_{\mu, v}|}{\sigma^{2}} e^{i [k_{\mu, v}] z} \sum_{i=0}^{2\pi} 2^{2} e^{i [k_{\mu, v}] z} - e^{2} / 2 \]  

where \(\mu\) and \(v\) represent the kernels' orientation and scale. \(k_{\mu, v}\) denotes the Gabor vector as in Eq. (8).

\[ k_{\mu, v} = k_{v} e^{i \phi} \]  

where, \(k_{v}\) equals \(k_{max}/v\). \(k_{max}\) represents the max frequency. \(f\) represents the spacing factor among kernels in the frequency space. \(\phi\) equals \(\pi/\phi\).

3.1.4 Gray Level Co-Occurrence Matrix (GLCM)

The GLCM technique is a statistical technique which applied to extract the texture features from an image [66].

The GLCM uses five equations to perform its task as follow:

- The contrast (con) equation represents the amount of local variations in an image \(i_{y}\). Eq. (9) calculates this equation.

\[ \text{con} = \frac{1}{L - 1} \sum_{i=0}^{L-1} \sum_{j=0}^{L-1} (i - j)^{2} \]  

- The correlation (corr) equation represents the relationship among the image pixels. Eq. (10) calculates this equation.

\[ \text{corr} = \frac{1}{2} \sum_{i=0}^{L-1} \sum_{j=0}^{L-1} \frac{(i - \mu_{i})(j - \mu_{j})}{\sigma_{i}^{2} \sigma_{j}^{2}} \]
Weibull distribution \( W_D \) \cite{46}:

A random variable has been shown to follow the Weibull distribution if it has the following tail formula as below:

\[
W_D(U) = \exp\left(\frac{U}{k}\right)\zeta
\]

(17)

where \( k \) and \( \zeta \) are the scale and shape parameters. The Weibull Distribution is a heavy-tailed when \( \zeta \) has values less than 1. Therefore in the current work, the tuned values of \( k \), and \( \zeta \) are 4 and 0.3, respectively.

The charts for the jump-length of the proposed distributions with a scaling factor of 0.01 against the number of functions evaluation are plotted in Fig. 1. By inspecting the figures, once can detect that (I) the Lévy and Cauchy distributions are two-sided distributions where the random numbers have positive and negative values. (II) Mittag-Leffler, Pareto, and Weibull distributions are one-side distributions as the random numbers have positive values. (III) All the distributions expose long jumps (steps) frequently with different step values.
3.3. Cuckoo search

The cuckoo search (CS) has been innovated by Yang et al. [36] inspiriting from the natural behavior of cuckoo breeding parasitism. In [36], authors modeled that behavior mathematically via three hypotheses mainly (I) each cuckoo lays one egg at a time, and (II) cuckoo puts egg in a nest that chosen randomly, and the fittest is retained for the next generation (III) the available host nests number is bounded, and a host cuckoo can detect a stranger egg with a probability \( P_i \in [0, 1] \). These behaviors can be formulated mathematically using Le\^vy flights. For each cuckoo \( i \), a new solution \( U^{t+1} \) is obtained based on Le\^vy flight as in Eq. (18).

\[
U_i^{t+1} = U_i^t + \alpha \text{Levy}(s, \lambda),
\]

where

\[
\text{Levy}(s, \lambda) = \frac{\lambda \Gamma(\lambda)}{\pi} \frac{1}{s^{1+\lambda}}
\]

(18)

where \( \alpha > 0 \) is the step size, as per the literature, it is tuned as 1. Le\^vy flight is a random walk, and its jumps are computed from a Le\^vy distribution.

For the local random walk, the solution will be updated using the following equation;

\[
U_i^{t+1} = U_i^t + \alpha s \bigotimes H(P_i - \epsilon) \bigotimes (U_j - U_k),
\]

where the product \( \bigotimes \) denotes entry-wise multiplications. Where \( U_j \) and \( U_k \) are two randomly selected solutions. The \( H \) is a Heavy-side function, \( s \) is the step size, and \( \epsilon \) is a random number drawn from a uniform distribution. Yang et al. [36] utilized a switching probability \( P_i \) to transmit amongst the local and global random walk and assumed its values as 0.25 to achieve a balance in the transmission stage.

3.4. Fractional-order cuckoo search

Recently Yousri et al. [40] proposed a novel variant for CS algorithm to improve the global cuckoo walk of Eq. (18) via accounting the memory prospective of cuckoo during motion based on fractional-calculus (FC) method. In the FO-CS, four previous terms from memory are saved for each cuckoo during its motion, and that memory has been modified based on the first input first output concept. Moreover, the switching probability \( P_i \) has been computed based on Beta function and FC parameters as follow:

- The enhanced global random walk of cuckoo based on the previous four terms from memory \( (m = 4) \) is updated as follow;

\[
U_i^{t+1} = \frac{1}{1!} \sigma U_i^t + \frac{1}{2!} \sigma (1 - \sigma) U_i^{t-1} + \frac{1}{3!} \sigma (1 - \sigma) (2 - \sigma) U_i^{t-2} + \frac{1}{4!} \sigma (1 - \sigma) (2 - \sigma) (3 - \sigma) U_i^{t-3} + \alpha \text{Levy}(s, \lambda),
\]

(20)

where \( \alpha \) refers to the derivative order coefficient [40]. The authors strongly believe that with increasing the memory terms, the execution time has been increased; therefore, in the current study, the number of terms is selected as \( m = 4 \) to maintain an acceptable execution time.

- In the FO-CS, the switching probability \( P_i \) has generated using the \( B \) distribution. The \( B \) function is calculated for \( \sigma \) from 0.1 to 1 with step 0.1 and number of memory terms \( (m = 4) \) using Eq. (21b).

\[
P_i(l) = \frac{(Z - \min(Z)) \ast (d - c)}{(\max(Z) - \min(Z))} + c
\]

(21b)

where \( B \) refers to the beta function with inputs of the derivative order vector \( \sigma_{vec} \) that is varied from 0.1 to 1 with step 0.1 and memory terms is \( m = 4 \). The generated distribution \( Z \) has minimum and maximum values \( \min(Z) \) and \( \max(Z) \). The normalization interval ranges are \( c = 0.2 \) and \( d = 0.3 \). The \( l = 1, 2, 3, \ldots, 10 \) denotes the index of the derivative order. The recommended value of the derivative order is 0.3 therefore the index \( l \) is 3 [40].

4. Proposed methodology

The structure of the proposed COVID-19 classification approach is given in Fig. 2, and the following steps summarized this structure. In general, the proposed COVID-19 classification method consists of two stages. The first stage is to extract the features using a set of methods, as discussed in Section 3.1. The second stage is to select the relevant features from those extracted features. To achieve this task, a modified FO-CS based on heavy-tailed distributions is implemented as a feature selection method. Where combining among the heavy-tailed distributions and FO-CS in place of the lévy flight distribution, open the door for new four variants of the FO-CS algorithm that can be viewed as FO-CS_ML, FO-CS_P, FO-CS_P, FO-CS_C,, and FO-CS_W. For the first variant FO-CS_ML, the Mittag-Leffler used in place of lévy flight. For the FO-CS_P, FO-CS_P, FO-CS_C, and FO-CS_W variants, the Pareto, Cauchy, and Weibull distributions, respectively, merged with FO-CS in place of lévy flight. Sequentially, the mathematical equation of Eq. (20) can be rewritten as follows for the four variants:

**In case of FO-CS_ML**:

\[
U_i^{t+1} = \frac{1}{1!} \sigma U_i^t + \frac{1}{2!} \sigma (1 - \sigma) U_i^{t-1} + \frac{1}{3!} \sigma (1 - \sigma) (2 - \sigma) U_i^{t-2} + \frac{1}{4!} \sigma (1 - \sigma) (2 - \sigma) (3 - \sigma) U_i^{t-3} + \alpha ML(\gamma, \delta),
\]

(22)

**In case of FO-CS_P**:

\[
U_i^{t+1} = \frac{1}{1!} \sigma U_i^t + \frac{1}{2!} \sigma (1 - \sigma) U_i^{t-1} + \frac{1}{3!} \sigma (1 - \sigma) (2 - \sigma) U_i^{t-2} + \frac{1}{4!} \sigma (1 - \sigma) (2 - \sigma) (3 - \sigma) U_i^{t-3} + \alpha P(a, b),
\]

(23)

**In case of FO-CS_C**:

\[
U_i^{t+1} = \frac{1}{1!} \sigma U_i^t + \frac{1}{2!} \sigma (1 - \sigma) U_i^{t-1} + \frac{1}{3!} \sigma (1 - \sigma) (2 - \sigma) U_i^{t-2} + \frac{1}{4!} \sigma (1 - \sigma) (2 - \sigma) (3 - \sigma) U_i^{t-3} + \alpha C(\beta, \mu),
\]

(24)

**In case of FO-CS_W**:

\[
U_i^{t+1} = \frac{1}{1!} \sigma U_i^t + \frac{1}{2!} \sigma (1 - \sigma) U_i^{t-1} + \frac{1}{3!} \sigma (1 - \sigma) (2 - \sigma) U_i^{t-2} + \frac{1}{4!} \sigma (1 - \sigma) (2 - \sigma) (3 - \sigma) U_i^{t-3} + \alpha W(\kappa, \zeta),
\]

(25)

The details of the stages of the proposed method are given as presented below:

4.1. Stage 1: Extract features

In this stage, the X-ray COVID-19 images are used as input to our methods. Then, for each image in the current data-set, a set of features are extracted using Fractional Zernike moments,
Wavelet, Gabor Wavelet, and GLCM, as discussed in Section 3.1. All of these features are collected in one matrix, where each row represents the extracted features from its corresponding COVID-19 image. This matrix of extracted features is used as an input to the second stage (i.e., to the developed FO-CS method) to determine the irrelevant features that are removed and the relevant features that are kept.

4.2. Stage 2: Features selection

This stage is considered the main core of the proposed COVID-19 detection using X-ray images. Since it aims to normal and abnormal images first, then determines COVID-19 patients from abnormal images. This aim is achieved by using the following steps:

4.2.1. Step 1: Construct initial population

The developed feature selection FO-CS method begins by dividing the data-set (features from the first stage) into training and testing sets, which represent 80% and 20% from the data-set, respectively. Then constructing a population of \( N \) real-valued solutions generated from a uniform random number. This step is formulated as:

\[ U_{ij} = LB_{ij} + r_1 \times (UB_{ij} - LB_{ij}), \quad j = 1, 2, \ldots, D, \quad i = 1, 2, \ldots, N \]  

(26)

In the above equation, the lower and upper boundaries of search space at dimension \( j \), and \( r_1 \) is a random number drawn from uniform distribution in the interval of \([0, 1]\).

4.2.2. Step 2: updating solutions

In this step, the solutions starting their updating by computing the fitness value of each of them. Since the FS problem is a discrete problem, so we convert the real-value of each solution to binary to deal with this type of problem. This process is formulated using the following equation.

\[ BU_{ij} = \begin{cases} 1 & \text{if } U_{ij} > 0.5 \\ 0 & \text{otherwise} \end{cases} \]  

(27)

Then features that corresponding 1's in \( BU \) are selected, and the other features are removed. Followed by using Eq. (28) to evaluate the performance of those selected features.

\[ Fit_i = \lambda \times \eta_i + (1 - \lambda) \times \left( \frac{|BU_i|}{\text{Dim}} \right) \]  

(28)

\( Fit_i \) represents the fitness value of the \( i \)th solution, \( \eta_i \) denotes the error of classification using the KNN classifier, \( |BU_i| \) is the number of selected features (i.e., the number of features which corresponding to ones in \( BU_i \)), \( \lambda \in [0, 1] \) denotes a uniform random number used to make a balance between two parts of the fitness value.

The next process is that the solutions \( U \) will be updated using the operators of the FO-CS algorithm and one of the heavy tail distributions. This performed by generating a step size in basis of the utilized distribution (i.e., Mittag Leffler Eq. (22), or Pareto Eq. (23), or Cauchy Eq. (24), or Weibull Eq. (25), or Levy flight Eq. (20)), respectively to update the current solution. Then using Eq. (21b) to compute the switching probability \( P_s \) and applied to discover the solution that will be replaced by new ones. Before starting a new iteration, the memory window can be updated using first in first out approach, as depicted in Algorithm 1.

4.2.3. Step 3: Terminal conditions of training

The main target of this step is to check whether the second step will be conducted again or not. This is achieved by checking if the terminal conditions are reached or not. In the case they are not reached, then step 2 (i.e., updating solution) is repeated again; otherwise, it will return by best solution \( U_b \).

4.2.4. Step 4: Evaluate best solution \( U_b \)

In this step, the testing set is used as input to KNN; however, their features that are corresponding to 0’s in \( U_b \) will be removed. Then predict the target of the testing set and compute the performance of this output using different metrics.

Algorithm 1: Pseudo-code of FO-CS algorithm

1: Set the population size \( N \), maximum number of iterations \( T \), the used distribution \((\text{Dis})\), derivative order coefficient \( \beta \) and number of terms \( r \).
2: Initialize the random population of \( n \) host nests \( U(i = 1, 2, \ldots, N) \).
3: Calculate the fitness values.
4: Initialize the past solutions (memory property).
5: \textbf{while} \(( t < T \) (Stop criteria is not satisfied)) \textbf{do} 
6: Get a cuckoo randomly.
7: \textbf{if} \( \text{Dis} \) is \( M_L \) \textbf{then}
8: Utilize the modified equation of motion Eq. (22) to update the solutions.
9: \textbf{else if} \( \text{Dis} \) is \( P \) \textbf{then}
10: Utilize the modified equation of motion Eq. (23) to update the solutions.
11: \textbf{else if} \( \text{Dis} \) is \( C \) \textbf{then}
12: Utilize the modified equation of motion Eq. (24) to update the solutions.
13: \textbf{else if} \( \text{Dis} \) is \( W \) \textbf{then}
14: Utilize the modified equation of motion Eq. (25) to update the solutions.
15: \textbf{else}
16: Utilize the equation of motion Eq. (20) to update the solutions.
17: \textbf{end if}
18: Update the past solutions (memory terms \((r)\)) on FIFO bases FIFO.
19: Compute the fitness function \( \text{Fit}_i(i = 1, 2, \ldots, N) \).
20: Select a nest from \( n \) randomly and compute the fitness at this index (index = \( j \)).
21: \textbf{if} \( \text{Fit}_i < \text{Fit}_j \) \textbf{then}
22: Replace \( U_j \) by the new one \( U_i \).
23: \textbf{end if}
24: Calculate \( P_s \) based on \( r = 4 \) and \( \beta = 0.3 \) via Eq. (21b).
25: Abandon a fraction \( P_s \) of worst nest and build new ones via Eq. (19).
26: Retain the best solutions.
27: Sort the solutions and determine the current best.
28: \textbf{t} = \textbf{t}+1.
29: \textbf{end while}
30: \textbf{Return the best solution}

5. Simulation and results

To validate the performance of the developed COVID-19 detection method, a set of experimental series is performed. The main aim of the first experimental series is to evaluate the performance of the main core of our COVID-19 detection (i.e., FO-CS based on heavy-tailed distributions), a set of eighteen UCI data-sets is used. Meanwhile, the second experimental series aims to test the applicability of the developed COVID-19 detection method by using two real-world COVID-19 images which have different characteristics.

5.1. Experimental Series 1: FS using UCI data-sets

In this experimental, eighteen well-regarded data-sets of UCI [47] source are used. These data-sets specifications are listed in
Table 1. As noted from Table 1, the examined benchmark problems diverse from the small to the high dimension to evaluate the efficiency of the proposed algorithm. In this study, the first stage of the proposed classification COVID-19 X-ray images (i.e., feature extraction) not used.

5.2. Comparative algorithms

We compared the improved FO-CS variants with several existing MA, including the original CS, Henry gas solubility optimization (HGSO) \[67\], Harris hawks optimization (HHO) \[68\].
5.3. Performance measures

In this section, the results of the proposed fractional-order cuckoo search algorithm using five types of distributions approaches are given; Levy distribution (FO-CS), Mittag-Leffler distribution (FO-CS\textsubscript{ML}), Pareto distribution (FO-CS\textsubscript{P}), Cauchy distribution (FO-CS\textsubscript{C}), and Weibull distribution (FO-CS\textsubscript{W}). Table 2 describes the accuracy results of the proposed methods compared to other competitor optimization methods.

It is apparent from Table 2 that the proposed FO-CS\textsubscript{ML} and FO-CS\textsubscript{P} got almost the same response over the proposed methods; each method got five best results out of eighteen. They gave better results than the other proposed methods (i.e., FO-CS, FO-CS\textsubscript{C}, and FO-CS\textsubscript{W}) in terms of the classification accuracy measure. The proposed FO-CS and FO-CS\textsubscript{W} also show almost the same performance over the proposed methods; each method excelled in three cases. They gave better results than only one of the other proposed methods (i.e., FO-CS\textsubscript{C}) according to used measure. The conventional CS performed better than only one of the proposed methods (FO-CS) over all the data-sets utilized in this paper in terms of the classification accuracy measure.

It is worthwhile to mention that the proposed FO-CS\textsubscript{ML} and FO-CS\textsubscript{P} got better results in five out of eighteen compared to all other comparative methods (i.e., HHO, HGSO, WOA, SSA, GWO, and GA). Meantime, SSA got the best results in four cases (i.e., DS1, DS11, DS17, and DS18) compared to all other methods. Generally, in terms of the accuracy measure, the proposed FO-CS outperformed the other state-of-the-art methods using various distribution approaches, mainly when used FO-CS\textsubscript{ML} and FO-CS\textsubscript{P}.

Table 3 shows the number of selected features achieved by the feature selection algorithms. From this Table, we can notice that the proposed FO-CS\textsubscript{W} made the smallest number of features selected in seven data-sets (DS6, DS7, DS11, DS13, DS16, DS17, and DS18). In comparison with the proposed FO-CS methods, the FO-CS\textsubscript{W} got the smallest optimal number of the selected features according to the results of the proposed FO-CS with the used distribution approaches, followed by FO-CS\textsubscript{SSA}. As well, in comparison with all tested methods, most of the best cases (in terms of the best number of selected features) have been accomplished by the FO-CS\textsubscript{W}, WOA achieved the smallest number of selected features in three data-sets, FO-CS\textsubscript{W} BDA gave a better performance than other methods on most of the used data-sets according to the number of selected feature measures. The high-performance of the FO-CS with Weibull distribution (FO-CS\textsubscript{W}) enhances its capability in reducing the number of selected features.

Table 4 summarizes the mean fitness function values obtained by the proposed algorithms. According to this measure, we can
reveal that the proposed FO-CSW is a superior method compared to the other proposed methods that are based on using the modified fractional-order cuckoo search algorithm with various distributions approaches. The proposed FO-CSW got the best mean fitness values in four cases, and it is not worse than any other method on most of the use data-sets. In comparison with all of the given methods in Table 4, FO-CSStat also outperformed the comparative methods in terms of the mean fitness function except HGSO; they got an equal number of best-obtained results (each method got the best results in two cases).

The standard deviation (STD) values of the proposed algorithm’s fitness function are given in Table 5. From these results, it can be seen that the proposed FO-CSStat exceeded the other proposed methods on most of the used data-sets. According to the STD measure, FO-CSW achieved the best results in four cases in terms of the STD measure. Moreover, FO-CSW achieved better
Table 8
Friedman test results for the first experiment.

| Measure    | CS | FO-CS | FO-CSML | FO-CSF | FO-CSW | HHO | HGSO | WOA | SSA | GWO | GA  |
|------------|----|-------|---------|--------|--------|-----|------|-----|-----|-----|-----|
| Fitness    | 4.47| 6.11  | 5.94    | 5.42   | 6.67   | 4.33| 5.56 | 8.11| 7.61| 8.72| 11.00|
| Rank       | 3   | 7     | 6       | 4      | 8      | 1   | 2    | 5   | 10  | 9   | 11  |
| Attributes | 6.36| 6.83  | 7.36    | 8.31   | 7.53   | 3.72| 4.58 | 4.50| 8.89| 4.44| 11.94|
| Rank       | 6   | 7     | 8       | 10     | 9      | 1   | 2    | 5   | 4   | 11  | 3   |
| Accuracy   | 5.72| 7.22  | 7.78    | 8.86   | 7.42   | 7.69| 7.14 | 6.36| 3.69| 7.17| 3.86|
| Rank       | 9   | 5     | 2       | 1      | 4      | 3   | 7    | 8   | 12  | 6   | 11  |

Fig. 3. Average of accuracy overall tested for each algorithm.

Fig. 4. Average of fitness value overall tested for each algorithm.

Fig. 5. Average of number of selected feature overall tested for each algorithm.
fitness function results with excellent STD values compared to other comparative methods in the majority of data-sets. Compared to all comparative methods in Table 5, HGSO got the best cases in most data-sets, which means that HGSO is a powerful competitor method in this research. Thus, the proposed FO-CS$_W$ beats all the comparative methods according to the STD measure. For the all feature selection algorithms, the minimum results of the fitness function values are given in Table 6. The obtained results proved that the proposed FO-CS$_W$ and FO-CS$_M$ are able to get better results compared to other proposed methods. It got the best results in six out of eighteen cases. However, FO-CS$_W$ provides average of the best fitness value better than FO-CS$_M$. 

Fig. 6. Minimum of fitness values by the recommended FO-CS$_W$ and the other FO-CS variants as well as the basic CS for (a) DS1, (b) DS2, (c) DS3, (d) DS4, (e) DS5, (f) DS6, (g) DS7, (h) DS8, (i) DS9, (j) DS10, (k) DS11, and (l) DS12.
overall the tested datasets. The obtained high-performance by the proposed FO-CSW algorithm demonstrates its capability to equilibrium the exploration and exploitation during the optimization process. As per the obtained-results, the performance of the proposed FO-CSW algorithm is proven on the small-sized data-sets and large-sized data-sets. The three data-sets (i.e., DS17, and DS16) are comparatively large, and the minimum fitness values of the proposed FO-CSW is obviously less compared to all other comparative methods.

According to the last evaluation measure (worst fitness function values), the proposed algorithms using the FO-CS with the used distribution approaches, as shown in Table 7, got better results than the other comparative methods relatively. The proposed FO-CSW achieved better results in comparison with the proposed algorithms as well as in comparison with other comparative methods. It obtained high-performance results (worst mean fitness values) in five out of eighteen experimented data-sets.

In general, the previous results showed that the best values obtained by FO-CS variants equal to 60% of all measures namely, they obtained the best values in 67% of the accuracy measure in all datasets, 0.50% of the selected features, and 78% of the standard deviation, also 56% of the fitness function values, the minimum, and the worst values. These results indicate that the variants of the proposed method can effectively compete with other algorithms in selecting the most relevant features.

Besides, from the given results, it can be recognized that the fractional-order calculus has increased the efficacy of the proposed algorithms, especially with the Weibull distribution approach in terms of fitness function values. The main reason here is that the proposed FO-CSW has evaded from converging towards the local area and improved its exploration functions to address more complex cases. Hence, it can make an excellent trade-off between the exploration and exploitation strategies.

Furthermore, Figs. 3–5 depict the results of the average of the accuracy, the fitness value, and the number of selected features, respectively, for each algorithm overall the tested data-sets. These figures provide extra evidence about the high performance of the proposed FO-CS variants with heavy-tailed distributions as feature selection approaches, which provide better results than the other methods.

For further analysis of the results of FO-CS variants, as feature selection approaches, the non-parametric Friedman test [73] is used as in Table 8. From this table, it can be noticed that the FO-CSp provides better results in terms of accuracy overall the tested data-set. Whereas, in terms of the number of selected features, FO-CSW allocates the first rank with a mean rank of nearly 3.5, and it has the smallest mean rank in terms of fitness value. Moreover, we can see that the FO-CSW obtained the first rank in the fitness function and the smallest selected attributes (in these measures, the smallest value is the best), whereas it was ranked third in the accuracy measure (in this measure the largest value is the best). Although the FO-CSW was ranked third, the first and the second rank were for the other variants of the proposed method (i.e., FO-CSp and FO-CSML); therefore, the proposed method variants obtained the first five ranks compared with the other algorithms.

Generally, the proposed FO-CSW has a superior exploration search capability due to the used fractional-order calculus between the candidate solutions, which encourage its exploration abilities over the search processes when it is needed and in the following phase, it can efficiently converge on the neighborhood of explored locations by using the Weibull distribution approach, chiefly, through the final iterations. Consequently, this proposed method (FO-CSW) has stabilized a balance between the exploration and exploitation strategies, so its influence can be recognized in the enhanced fitness results and the smallest number of selected features compared to all other comparative methods in this research.

As the acceleration convergence speed is the other aspect that should be discussed and evaluated for the recommended FO-CS variant, the convergence curves based on the optimal fitness
Fig. 8. Average of fitness values by the recommended FO-CS\textsubscript{W} and the other FO-CS variants as well as the basic CS for (a) DS1, (b) DS2, (c) DS3, (d) DS4, (e) DS5, (f) DS6, (g) DS7, (h) DS8, (i) DS9, (j) DS10, (k) DS11, and (l) DS12.

function and the mean convergence curves for the basic CS and FO-CS variants have been drawn for the eighteen datasets to demonstrate the efficiency of the recommended FO-CS\textsubscript{W} as in Figs. 6, 7 and 8, 9, respectively. By inspecting the convergence curves of the minimum fitness functions of Figs. 6–7, it can be seen that the CS suffers from stagnation in several functions in nearly 80% of the studied datasets. In contrast, the FO-CS variants expose high qualified performance, especially FO-CS\textsubscript{W}, FO-CS\textsubscript{ML}, and FO-CS\textsubscript{C} as they show successful performance for 6, 6 and 5 datasets, respectively. The average convergence curves across the set of independent runs are drawn as in Figs. 8–9 for achieving unbiased comparison. By inspecting the figures, it can be seen
Fig. 9. Average of fitness values by the recommended FO-CSW and the other FO-CS variants as well as the basic CS for (a) DS13, (b) DS14, (c) DS15, (d) DS16, (e) DS17 and, (f) DS18.

Table 9
FS results of the FO-CS variants along with the compared algorithms in both COVID-19 data-sets.

| Data-set1 | CS | FO-CS | FO-CS_{W} | FO-CS_{P} | FO-CS_{C} | HHO | HGSO | WOA | SSA | GWO | SGA |
|-----------|----|-------|-----------|-----------|-----------|-----|------|-----|-----|-----|-----|
| **Accuracy** | Best | 0.7133 | 0.8067 | 0.8467 | 0.8333 | 0.8267 | 0.8400 | 0.7600 | 0.8267 | 0.7833 | 0.7533 |
|           | Mean | 0.6907 | 0.7613 | 0.8267 | 0.8133 | 0.8093 | 0.8267 | 0.7653 | 0.7587 | 0.7680 | 0.7080 |
|           | STD | 0.0167 | 0.0307 | 0.0249 | 0.0130 | 0.0105 | 0.0267 | 0.0173 | 0.0417 | 0.0238 | 0.0272 |
| **Fitness value** | Best | 0.2747 | 0.0144 | 0.0118 | 0.0144 | 0.0175 | 0.0123 | 0.0455 | 0.0389 | 0.0491 | 0.2638 |
|           | Mean | 0.2814 | 0.0180 | 0.0159 | 0.0166 | 0.0187 | 0.0098 | 0.0328 | 0.0324 | 0.0388 | 0.0255 |
|           | STD | 0.0048 | 0.0027 | 0.0047 | 0.0028 | 0.0023 | 0.0023 | 0.0089 | 0.0060 | 0.0078 | 0.0072 |
| **Attr** | 177.2 | 25.6 | 27.6 | 27.6 | 25.4 | 27.4 | **23.4** | 25.4 | 26.4 | 172.4 | 72.2 |

| Data-set2 | CS | FO-CS | FO-CS_{W} | FO-CS_{P} | FO-CS_{C} | HHO | HGSO | WOA | SSA | GWO | SGA |
|-----------|----|-------|-----------|-----------|-----------|-----|------|-----|-----|-----|-----|
| **Accuracy** | Best | 0.9895 | 0.9842 | 0.9895 | 0.9947 | 0.9717 | **1.0000** | 0.9789 | 0.9789 | 0.9526 | 0.9895 |
|           | Mean | 0.9779 | 0.9779 | 0.9874 | 0.9842 | 0.9705 | **0.9926** | 0.9632 | 0.9579 | 0.9317 | 0.9789 |
|           | STD | 0.0094 | 0.0069 | **0.0029** | 0.0064 | 0.0047 | 0.0060 | 0.0134 | 0.0144 | 0.0177 | 0.0064 |
| **Fitness value** | Best | 0.2814 | 0.0180 | 0.0159 | 0.0166 | 0.0187 | **0.0098** | 0.0328 | 0.0324 | 0.0388 | 0.2555 |
|           | Mean | 0.2814 | 0.0180 | 0.0159 | 0.0166 | 0.0187 | **0.0098** | 0.0328 | 0.0324 | 0.0388 | 0.2555 |
|           | STD | 0.0048 | 0.0027 | 0.0047 | 0.0028 | 0.0023 | 0.0023 | 0.0089 | 0.0060 | 0.0078 | 0.0072 |
| **Attr** | 123.8 | 23.2 | 24.4 | 24 | **21.8** | 22.8 | 26.6 | 25.2 | 22.6 | 112.2 | 68 |

from this convergence that, in most cases, the FO-CSW has faster convergence, for example, DS2, DS9, DS11, DS12, DS16, DS17, and DS18. However, FO-CSW provides better convergence at DS3, DS5, and DS13. Also, CS has the best convergence rate at DS1 and DS7. From all these graphs, it can be observed that the high effect of Fractional-order and Heavy-tailed distributions on the convergence CS.

5.4. Experimental series: FS using COVID-19 data-sets

In this section, the FO-CS variants are evaluated using two real-world COVID-19 images which have different characteristics.

5.4.1. Data-sets description

In this study, two data-sets are used to assess the performance of the developed FO-CS as the COVID-19 X-ray classification method. These data-sets are contained only two classes (i.e., COVID and No-Findings), which are used for binary classification, or they have three classes (i.e., COVID, No-Findings, Pneumonia) which used for multi-class classification. The description of these data-sets are given in the following.

1. Data-set 1: This is an X-ray image data-set which collects from different two sources. The first source is the COVID-19 X-ray images collected by Cohen JP [74]. This source has 127 COVID-19 X-ray. The second source named Chest
X-ray, which collected by Wang et al. [58]. This dataset contains normal and pneumonia X-ray images. From this source, 500 no-findings and 500 pneumonia are used. Fig. 10 depicts sample from this data-set (Data-set 1).
5.4.2. Results and discussion

In this section, the comparison results between the variants of FO-CS and other algorithms are discussed. All of the algorithms used the extracted features from algorithms introduced in Section 5.1. In this section, the extracted features from the COVID-19 x-ray images are the input for the proposed approach that aimed to reduce these features by removing the irrelevant ones. The parameter settings for each algorithm are similar to the previous experiments of the UCI data-set.

Table 9 shows the average of results in terms of accuracy, fitness value, and the number of selected features. One can be observed from this table that the FO-CS_w algorithm provides better results than other algorithms in terms of accuracy. In general, it has the first rank according to the STD, and the worst of accuracy. However, according to the Best value of the accuracy, the FO-CS_M has the higher. By analyzing the results of COVID-19 data-set1 in terms of fitness value, it has been observed that the FO-CS_w has better value according to the Mean and Best fitness value. However, the GA and FO-CS_M provide better results in terms of STD and worse of fitness value, respectively.

According to the results of the algorithms using the second COVID-19 data-set (i.e., Data-set 2), it can be observed that mean and Best of the accuracy of FO-CS_w is better than others. Also, in the worst value of accuracy, FO-CS_M and FO-CS_w have the same performance, but FO-CS_M is more stable in terms of accuracy. In addition, the FO-CS_w provides better performance in terms of Mean, Best, and Worst fitness value, but FO-CS_C is more stable than FO-CS_w in terms of fitness value. From all of these results, it can be noticed that the following observation, firstly, the variants of FO-CS, in general, provide better results than other algorithms. Secondly, the heavy-tailed distribution has the largest effect on FO-CS, which has high accuracy and smaller fitness value than levy flight distribution. Finally, using the content of FO leads to improve the performance of CS and the conclusion from the results of FO-CS when compared with traditional CS algorithms applied to COVID-19 X-ray images, and they converge faster than other algorithms, followed by FO-CS_M and FO-CS_P, respectively for the first data-set. The FO-CS_w is the superior one in the case of the second data-set.

From the previous results, we noticed that the proposed FO-CS based on heavy-tailed distributions provide better results than other algorithms for both UCI datasets and COVID-19 X-ray images. However, there are some limitations of these variants, such as determine the optimal value of the parameters for heavy-tailed distributions. In addition, determine the number of terms and derivative order coefficient.

6. Conclusions

In this study, we present four variants of the recent fractional-order cuckoo search optimization algorithm (FO-CS) using heavy-tailed distributions instead of Lévy flights to classify the extracted features from COVID-19 x-ray data-sets. The considered heavy-tailed distributions included the Mittag-Leffler distribution, the Pareto distribution, the Cauchy distribution, and the Weibull distribution. The application of these distributions can be used to improve mutation operators of MA, and it can be used to escape from non-prominent regions of the search space. For appraising the performance of the proposed variants before applying to the COVID-19 classification approach, we used eighteen data-sets from the UCI repository, and we compared their results with several well-regarded MA utilizing several statistical measures. At this end, the FO-CS based on heavy-tailed distributions variants have been employed for the COVID-19 classification optimization task to classify the data-sets for normal patients, COVID-19 infected patients, and Pneumonia patients. Two different data-sets have been studied, and the FO-CS novel approaches have been compared with numerous MA to evaluate the proposed approach for classifying such of those important data-sets to introduce a reliable and robust technique that helped in classifying the COVID-19 data-sets efficiently and with high accuracy. The FO-CS based on Weibull distribution showed its superiority compared to the other proposed variants and recent well-regarded MA. In future work, we will try to evaluate the proposed method in different applications, such as parameters estimation and solving various engineering problems.

CRediT authorship contribution statement

Dalia Yousri: Conceptualization, Data curation, Formal analysis, Methodology, Software, Visualization, Writing - review &
editting. Mohamed Abd Elaziz: Data curation, Formal analysis, Methodology, Software, Visualization, Writing - original draft. Laith Abualigah: Formal analysis, Supervision, Writing - review & editing. Diego Oliva: Conceptualization, Validation, Writing - review & editing. Mohammed A.A. Al-qaness: Conceptualization, Investigation, Writing - original draft. Ahmed A. Ewees: Formal analysis, Project administration, Software, Writing - original draft.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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