

Summarization Application Testing

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Control: Abstract from the paper

ABSTRACT COVID-19 outbreak has put the whole world in an unprecedented difficult situation bringing life around the world to a frightening halt and claiming thousands of lives. Due to COVID-19's spread in 212 countries and territories and increasing numbers of infected cases and death tolls mounting to 5,212,172 and 334,915 (as of May 22 2020), it remains a real threat to the public health system. This paper renders a response to combat the virus through Artificial Intelligence (AI). Some Deep Learning (DL) methods have been illustrated to reach this goal, including Generative Adversarial Networks (GANs), Extreme Learning Machine (ELM), and Long /Short Term Memory (LSTM). It delineates an integrated bioinformatics approach in which different aspects of information from a continuum of structured and unstructured data sources are put together to form the user-friendly platforms for physicians and researchers. The main advantage of these AI-based platforms is to accelerate the process of diagnosis and treatment of the COVID-19 disease. The most recent related publications and medical reports were investigated with the purpose of choosing inputs and targets of the network that could facilitate reaching a reliable Artificial Neural Network-based tool for challenges associated with COVID-19. Furthermore, there are some specific inputs for each platform, including various forms of the data, such as clinical data and medical imaging which can improve the performance of the introduced approaches toward the best responses in practical applications.

Occurrences of Keywords: 7

Summary using only weighting system:

INDEX TERMS Artificial intelligence, big data, bioinformatics, biomedical informatics, COVID-19, deeplearning, diagnosis, machine learning, treatment. I. Appropriating AI techniques to deal with COVID-19 related issues can fill the void between AI-based methods and medical approaches and treatments. In this paper, our team relies on the findings of the most recent research focusing on COVID-19 and its various challenges to generalize and suggest a variety of strategies relevant but not limited to high-risk groups, epidemiology, radiology and etc. Objectives of data understanding include understanding data attributes and identifying main characteristics such as data volume and the total number of variables to summarize the data. Before processing and analysis comes data preparation that is the process through which raw data are refined and converted. Humans' contribution at this stage is important because their knowledge and potentials are not available to an ML solution that unlike humans is able to deal with huge data sets far beyond the extent that humans could handle or observe in a simultaneous manner. DL methods, as Fig. As a subset of machine learning, DL consists of numerous layers of algorithms that provide a different interpretation of the data it feeds on. . categorized into 3 parts, including high-risk groups, outbreak and control, recognizing and diagnosis. . The input layer as the initial layer is related to the database and is designed for database access. Taking advantage of a good number of microprocessors with database software database machines can send huge packets of data to the mainframe. If physicians confirm the decisions made by this layer, the recommended techniques in the third layer take the required images. The conventional optical microscope has come to be the dominant tool in pathological examinations. The fourth layer is dedicated to the optimization and improvement of the images. for network structure, and the classical ResNet was used to extract features [24]. DL technologies, such as a convolutional neural network (CNN), are supposed to be the right option for achieving these goals. . III. Since reaching the best possible results is the main objective, we will try to demonstrate ways through which ANN-based methods could. As [29] suggested it is necessary to keep patients involved COVID-19 registry that highlights clinical variables and cardiovascular complications because it facilitates the identification of the pattern of cardiovascular complications, further developing a risk model for cardiac complications, and assists with identification and/or prediction of the response to different types of treatment modalities. . data that [29] presents to predict the ways that cardiovascular system is affected by the Coronavirus. Therefore, the suggested model is capable of reducing the risk of possible cardiovascular complications. Hence, considering their properties and multiple advantages, ELMs are recommended for such problems. Another complication that COVID-19 causes in the elderly is heart failure, which requires heart failure specialists stay on guard and design a structured approach to these type of patients and include them in developing algorithms for the care of these patients in early stages until the time when definite universal COVID-19 examinations or clinical trials of antivirals are in place, and deeper understanding of final stages of the disease is realized [35]. Reference [35] and biomarkers, especially in high-risk elderly patients with underlying structural cardiac disease should be used with care and caution. As such, defining and managing advanced heart failure in the phase of hyper inflammation are important issues for heart specialists [35]. Fig. E LM ANN can use previous examples applied to the model to predict desired outputs. This means that training the supervised model happens through the

(Only a sample of summary, total summary saved as aiCovid_weightOnlySummary.txt)

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Another complication that COVID-19 causes in the elderly is heart failure, which requires heart failure specialists stay on guard and design a structured approach to these type of patients and include them in developing algorithms for the care of these patients in early stages until the time when definite universal COVID-19 examinations or clinical trials of antivirals are in place, and deeper understanding of final stages of the disease is realized [35]. Despite much praise that such data has received because of its role in improving efficiency, productivity and processes in different sectors, it has been criticized for its small number of users who collect, store, manage the data and have access to them [76]. Collecting, analyzing and leveraging the data such as consumer, patient, physical, and clinical data ends in big data. Different steps in the application of AI-based methods employed to overcome COVID-19 challenges are presented in the flowchart shown in Fig.1. The first step is the preparation of the data which are necessary for data mining during data understanding, data preparation and big data. In other words, it is a process in which data are reformatting, corrected and combined to enriched data. Moreover, Deep Learning (DL) methods could be employed in cases where enormous or complex data processing challenge ML or traditional means of data processing. Before processing and analysis comes data preparation that is the process through which raw data are refined and converted. Objectives of data understanding include understanding data attributes and identifying main characteristics such as data volume and the total number of variables to summarize the data. makes it possible to tell when wrong things are happening, or actions are to be taken regarding COVID-19 because it monitors and collects data coming from social media, news-feeds, and airline ticketing systems [77]. It is at this stage that human intervention, as a part of machine learning methods, takes place and experts investigate and analyze the data to extract the data with finest structures, patterns and features. Reference [64] has studied clinical and biological data of five COVID-19 patients. Advanced machine learning algorithms can integrate and analyze large-scale data related to COVID-19 patients to facilitate a deeper understanding of viral spread pattern, improve the speed and accuracy of diagnosis, develop fresh, effective therapeutic approaches, and even identify individuals who, depending on their genetic and physiological features, are most susceptible to the disease [75].

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