

Deep Learning-Driven Automated Detection of COVID-19 from Radiography Images: a Comparative Analysis

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Abstract

The COVID-19 pandemic has wreaked havoc on the whole world, taking over half a million lives and capsizing the world economy in unprecedented magnitudes. With the world scampering for a possible vaccine, early detection and containment are the only redress. Existing diagnostic technologies with high accuracy like RT-PCRs are expensive and sophisticated, requiring skilled individuals for specimen collection and screening, resulting in lower outreach. So, methods excluding direct human intervention are much sought after, and artificial intelligence-driven automated diagnosis, especially with radiography images, captured the researchers' interest. This survey marks a detailed inspection of the deep learning–based automated detection of COVID-19 works done to date, a comparison of the available datasets, methodical challenges like imbalanced datasets and others, along with probable solutions with different preprocessing methods, and scopes of future exploration in this arena. We also benchmarked the performance of 315 deep models in diagnosing COVID-19, normal, and pneumonia from X-ray images of a custom dataset created from four others. The dataset is publicly available at https://github.com/rgbnihal2/COVID-19-X-ray-Dataset. Our results show that DenseNet201 model with Quadratic SVM classifier performs the best (accuracy: 98.16%, sensitivity: 98.93%, specificity: 98.77%) and maintains high accuracies in other similar architectures as well. This proves that even though radiography images might not be conclusive for radiologists, but it is so for deep learning algorithms for detecting COVID-19. We hope this extensive review will provide a comprehensive guideline for researchers in this field.

Keywords COVID-19 · Deep learning · Radiography · Automated detection · Medical imaging · SARS-CoV-2

Introduction

COVID-19 has become a great challenge for humanity. Fast transmission, the ever-increasing number of deaths, and no specific treatment or vaccine made it one of the biggest problems on earth right now. It has already been characterized as a pandemic by the World Health Organization (WHO) and is being compared to the Spanish flu of 1920 that took millions of lives. Even though the fatality

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Department of Robotics and Mechatronics Engineering, University of Dhaka, Dhaka, Bangladesh rate of the disease is only 2–3% [1], the more significant concern is its rapid spreading among humans. The reproductive number of the virus is between 1.5 and 3.5 [2], making it highly contagious. Therefore, early diagnosis is essential to contain the virus. This, however, has proved to be very difficult as the virus can stay inactive in humans approximately 5 days before showing any symptoms [3]. Even with symptoms, COVID-19 is hard to be distinguished from the common flu.

At present, one of the most accurate ways of diagnosing COVID-19 is by a test called Reverse Transcription Polymerase Chain Reaction (RT-PCR) [4]. Since the coronavirus is an RNA virus, its genetic material is reverse transcribed to get a complementary DNA or cDNA. This can then be amplified by polymerase chain reaction or PCR, making it easier to measure. However, it is a complicated and time-consuming process, taking almost 2–3 h and demands the involvement of an expert. Newer technology can produce results in 15 min, but it is costly. Even then,

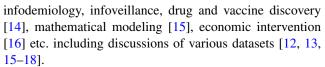


there have been studies showing that RT-PCR can yield false negatives [5]. There are newer machines that can autonomously carry on the tests, eliminating human errors and health risks associated with it. However, it is both costly and unavailable in many parts of the world. Moreover, RT-PCR only detects the presence of viral RNA. It cannot prove that the virus is alive and transmissible [6]. The testing material is also of scarcity due to the sheer number of cases in the pandemic, leading to increasing costs.

Another method of COVID-19 detection is antibody testing [7]. It aims to detect the antibody, generated as an immune response of the COVID-19-affected body. This testing method was designed for mass testing for the already affected. It is cheap and fast, producing results in 15 min and can be carried out in a modest laboratory. However, the problem is that the average incubation period of the coronavirus is 5.2 days [3], and antibodies are often not generated before a week from infection and sometimes even later than that. Thus, early containment is not possible. Also, this testing method is susceptible to both false positives and false negatives due to the cases of minor symptoms. Thus, in terms of early detection and containment, this method is not quite up to the task.

Since the outbreak of this disease, researchers have been trying to find a way to detect COVID-19 that is fast, cheap, and reliable. One of the prominent ideas is to diagnose COVID-19 from radiography images. Studies show that one of the first affected organs in coronavirus cases is the lungs [8]. Thus, radiography images of the lungs could give some insight on their condition. Radiologists, however, often fail to diagnose COVID-19 successfully solely from the images due to the similarity between COVID-19affected lung images and pneumonia-affected lung images and sometimes even normal lung images. Besides, manual interpretation may suffer from inter and intra-radiologist variance and be influenced by different factors such as emotion and fatigue. Recent advances in deep learning regarding such diagnostic problems allow Computer-Aided Diagnosis (CAD) to reach new heights with its ability to learn from high-dimensional features automatically built from the data. Especially during this pandemic, when expert radiologists are conclusively experiencing difficulties diagnosing COVID-19, CAD seems to be the top candidate to assist the radiologists and doctors in the diagnosis. Works like [9-11] and many more are showing the potential of deep learning-driven CAD to face this pandemic.

Not just in the diagnosis of the virus, but there have been many works done with deep learning applied to almost all the sectors affected by the coronavirus. And the flood of such works have resulted in a number of surveys relating to the role of artificial intelligence in this pandemic situation covering not just diagnosis but also clinical management and treatment [12], image acquisition [13],



The overall observation is that most of the surveys done to date tried to cover a wider extent of the domain instead of depicting an exhaustive overview in one direction. However, we are motivated to focus on what we consider to be the most important aspect of fighting the dreadful disease, and that is detection and diagnosis. Throughout this work, we dispensed a comprehensive discussion on available datasets, existing approaches, research challenges with probable solutions, and future pathways for deep learning empowered automated detection of COVID-19. In addition to the qualitative assay, we provided a quantitative analysis that comprises of extensive experimentation using 315 deep learning models. We tried to investigate some of the key questions here: (1) What are the key challenges to diagnose the disease from radiography data? (2) Which CNN architecture performs the best to extract distinct features from the X-ray modality? (3) How transfer learning can be utilized and how well does it perform being pretrained on the widely used ImageNet dataset [19]? To the best of our knowledge, this is the very first survey that includes a benchmark study of 315 deep learning models in diagnosis of COVID-19.

The rest of the paper is organized as follows. In "Related Work," we present a study on related works. Subsequently, Section "Radiography-Based Automated Diagnosis: Hope or Hype" describes the radiography based diagnosis process. We shed light on some challenges in radiographybased diagnosis in "Challenges." In "Description of Available Datasets," we describe the publicly available radiography datasets. Detailed analysis of deep learning-based approaches of COVID-19 diagnosis is presented in "Deep Learning-Based Diagnosis Approaches". Section "Image Prepossessing" reviews different data preprocessing techniques. Section "Comparative Analysis" provides a quantitative analysis of some state-of-the-art deep learning architectures on our compiled dataset. Finally, we conclude this paper with future research directions in "Discussion and Future Directions" and concluding remarks in "Conclusion."

Related Work

This paper gives a comprehensive overview of the automated detection of COVID-19 through data-driven deep learning techniques. Previously, Ulhaq et al. [12] discussed the existing literature on deep learning-based computer vision algorithms relating to the diagnosis, prevention, and control of COVID-19. They also discussed the infection management and treatment-related deep learning algorithms



along with a brief discussion about some of the existing datasets. However, they included a pre-print (non peer-reviewed) version of some papers which limit the acceptability of the work.

In [14], the authors discussed the role of AI and big data in fighting COVID-19. In addition to the existing deep learning architectures available for detection and diagnosis of COVID-19, they discussed the existing SIR (Susceptible, Infected, and Removed) models and other deep models for identification, tracking, and outbreak prediction. The survey also included different speech and text analysis methods, cloud-based algorithms for infodemiology and infoveillance, deep learning algorithms for drug repurposing, big data analysis—based outbreak prediction, virus tracking, vaccine, and drug discovery, etc. They did not, however, show any comparative quantitative analysis of the reviewed works.

Shi et al. [13] reviewed medical imaging techniques in battling COVID-19. Various contact-less image acquisition techniques, deep learning-based segmentation of lungs and lesion, X-ray and CT screenings, and severity analysis of COVID-19 along with some publicly open datasets are included in the work. Nevertheless, they did not provide any quantitative analysis of existing methods either. Also, their discussions on the existing datasets are somewhat inadequate.

In [15], the authors discussed mathematical modeling of the pandemic with SIR, SEIR (Susceptible, exposed, infected, and removed) and SIQR (Susceptible, infected, quarantined, and recovered) models.

In [17], the authors gave a comparative list of publicly available datasets consisting of the image data of COVID-19 cases. However, they did not shed any light on the works done so far and also did not provide any pathways for future research in this domain. In contrast to other studies, Latif et al. [16] discussed deep learning algorithms for risk assessment and prioritization of patients in addition to diagnosis.

In [18], Nguyen discussed deep learning—driven medical imaging, IoT-driven approaches for the pandemic management, and even Natural Language Processing (NLP)—based approaches for COVID-19 news analysis. However, like others, they did not give any comparative analysis either.

Most of the discussed papers have tried to cover a wide area of topics and, thus, lacked detailed discussions in a single domain. None of the papers provides a quantitative analysis of the works discussed, something that can be very helpful to the researchers. Our work aims to overcome these limitations by focusing on a single domain, COVID-19 detection. We also provide a quantitative and comparative analysis of 315 different deep learning algorithms, something that is yet to be done by any

other survey papers. A summary of all the survey papers covered in this section is presented in Table 1.

Radiography-Based Automated Diagnosis: Hope or Hype

Radiography images, i.e., chest X-ray and computed tomography (CT), can be used to diagnose COVID-19 as the disease primarily affects the respiratory system of the human body [8]. The primary findings of chest X-rays are those of atypical pneumonia [20] and organizing pneumonia [21]. The most common finding in chest radiography images is Ground Glass Opacity (GGO), which refers to an area with increased attenuation in the lung. As shown in Fig. 1, a chest X-ray image shows some hazy grey shade instead of black with fine white blood vessels. In contrast, CT scans show GGO [8] (Fig. 2a), and in severe cases, consolidation (Fig. 2b). Chest images sometimes also show something called "crazy paving" (Fig. 2c), which refers to the appearance of GGO with a superimposed interlobular and intralobular septal thickening. These findings can be seen in isolation or combination. They may occur in multiple lobes and affect in the peripheral area of the lungs.

It is worth noting that chest CT is considered to be more sensitive [22] for early COVID-19 diagnosis than chest X-ray since chest X-ray may remain normal for 4–5 days after start of symptoms where CT scan shows a typical pattern of GGO and consolidation. Besides, CT scans can show the severity of the disease [23]. A recent study performed on 51 COVID-19 patients shows that the sensitivity of CT for COVID-19 infection was 98% compared to RT-PCR sensitivity of 71% [24].

However, CT scans are difficult to obtain in COVID-19 cases. It is mainly due to the difficulty of decontamination issues regarding patient transports to the CT suites. As a matter of fact, the American College of Radiology dictates that CT scan-related hassles can disrupt the availability of such radiology services [25]. Another problem is that CT is not available worldwide, and in most cases, expensive and, thus, has a low outreach. This is why, despite the lesser sensitivity and quality, chest X-rays are the most common method of diagnosis and prognosis of not only COVID-19 cases but most other lung-related abnormalities.

The main problem is that these findings are not only found in COVID-19 cases but also in pneumonia cases. In many mild COVID-19 cases, the symptoms are similar to that of the common cold and sometimes show no different than that of normal lungs. Even though Research in [27] has indicated that the radiography image of COVID-19-affected lungs differs from the image of bacterial pneumonia-affected lungs. In COVID-19 cases, the effect is more likely



Table 1 Related surveys on AI techniques for detecting COVID-19 from radiography images

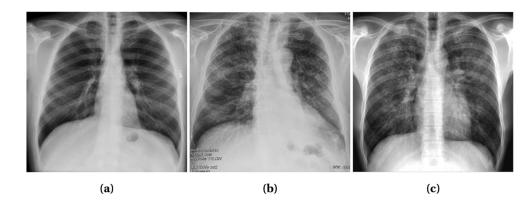
Study	Key topics	No. of reviewed papers ¹	No. of discussed datasets ²	Benchmarking deep models ³
Ulhaq et al. [12]	Vision-based diagnosis, control and treatment	21	6	-
Pham et al. [14]	AI and big data-based diagnosis, outbreak prediction, and biomedicine	32	2	-
Shi et al. [13]	AI-based image acquisition, segmentation, and diagnosis	14	4	-
Kalkreuth et al. [17]	COVID-19 dataset listing	4	12	_
Latif et al. [16]	AI-based COVID-19 diagnosis, pan- demic modeling, dataset description, and bibliometric analysis	25	5	-
Nguyen [18]	AI-based COVID-19 diagnosis, modeling, text mining, and dataset description	12	10	-
Mohamadou et al. [15]	Mathematical modeling of pandemic and COVID-19 diagnosis	20	6	-
Our study	Deep learning-based COVID-19 diagnosis	38	16	Benchmarked 315 deep models that comprises the combinations of 15 CNNs and 21 classifiers

¹Diagnosis-related papers, ²Radiography-based datasets, ³"-" means not applicable for the paper The Italic entries signify our contributions

to be scattered diffusely across both lungs, unlike typical pneumonia. However, in the early stages, even expert radiologists are often unable to detect or distinguish between COVID-19 and pneumonia.

Amidst such a frightful situation, deep learning-driven CAD seems a logical solution. Deep learning can extract and learn from high-dimensional features humans are not even able to comprehend. So, it should be able to deliver in this dire situation as well. Moreover, there has already been a flood of such approaches recently with good results, showing hope in this crisis period. However, many such hopeful ideas have turned into false hopes in the past. This work investigates the different works relating to deep learning aided CAD to resolve whether this is our hope, or if it is only another hype.

Fig. 1 X-ray images with different infection types: a Patchy GGOs present at both lungs; b Nuanced parenchymal thickenings; and c GGOs with some interstitial prominence. Images obtained from [26]



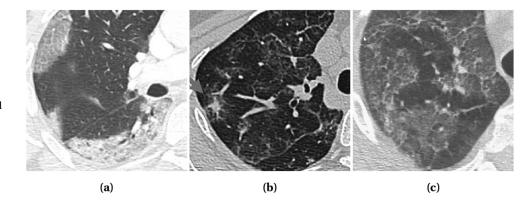
Challenges

As discussed in the previous section, radiography image can pave an efficient way to the detection of the COVID-19 at an earlier stage. However, the unavailability and quality issues related to COVID-19 radiography images introduce challenges in the diagnosis process while effecting the accuracy of the detection model. Here we discuss some of the major challenges faced by the researchers in the detection of COVID-19 from radiography images.

Scarcity of COVID-19 Radiography Images Significant numbers of radiography image datasets are made available by the researchers to facilitate collaborative efforts for combating COVID-19. However, they contain at most a few



Fig. 2 CT scan showing different infection types: a Subpleural GGOs with consolidations in all lobes; b GGOs with probable partially resolved consolidations; and c Scattered GGOs with band consolidations. Images obtained from [26]



hundreds of radiography images of confirmed COVID-19 patients. As a result, poor predictions are made by the models being over-fitted by insufficient data which puts a cap on the potential of deep learning [28]. Data augmentation techniques can be used to increase the dataset volume. Common augmentation techniques like flipping and rotating can be applied to image data with satisfactory results [29]. Transfer learning is another alternative to deal with insufficient data size while reducing generalization errors and over-fitting. Significant numbers of works integrated data augmentation and transfer learning into deep learning algorithms to obtain pleasing performances [30–40]. Few shot learning [41] and zero shot learning [42] are also plausible solutions to the data insufficiency problem.

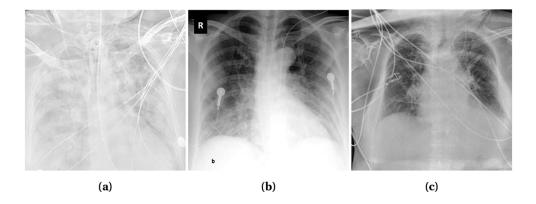
Class Imbalance This is a common problem faced while detecting COVID-19 from radiography images. Radiography datasets consisting of a sufficient number of Xray images of pneumonia-affected and normal lungs are available on a large scale. In contrast, for being a completely new disease, the number of images of COVID-19affected lungs are significantly less than that of normal and pneumonia-affected lungs. As a result, the model becomes prone to giving poor predictions to the minority class. Re-sampling of the datasets is often performed as a solution to the class imbalance problem, which attempts to make the model unbiased from the majority classes. In random under sampling strategy, samples for the minor classes are duplicated randomly, whereas samples from the majority classes are removed randomly in random over sampling methods to mitigate the class imbalance of a dataset [43]. However, over sampling may result in over-fitting [44], which can be reduced adopting any of the improved oversampling techniques-Synthetic Minority Over Sampling (SMOTE)[45], borderline SMOTE [46], and safe level SMOTE [47]. Another technique to deal with the class imbalance problem is to assign weights to the cost function, which ensures that the deep learning model gives equal importance to every class. Some of the works applied data augmentation, e.g., LV et al. [48] employed a Module Extraction technique (MoEx) where standardized features of one image are mixed with normalized features of other images. In another work, Punn et al. [49] manifested class average and random oversampling as an alternative method to data augmentation.

Artifacts/Textual Data and Low Contrast Images In radiography images, artifacts like wires, probes, or necklaces are often present as depicted in Fig. 3. Even image annotation with textual data (e.g., annotation of the right and left side of an image with "R" and "L," respectively) is a common practice. These artifacts hamper the learning of a model and lead to poor prediction results. Although textual data (Fig. 3b) can be erased manually by covering it with a black rectangle [32], it is time consuming. A more advanced efficient way is to use a mask of two lungs (for X-ray images) and concatenate with the original image [35, 48, 49]. Thus, the unnecessary areas are ignored and the focus is only given on the interested areas. Mask can be generated using U-Net[48] or binary thresholding [35, 49]. In CT images, the lungs are segmented to focus on the infectious regions [50]. Segmentation tools include U-Net, VB-Net, BCDU-Net, and V-Net which are used in [51–53] and [54], respectively. In some cases, image quality issues such as low contrast (Fig. 3a) introduces challenges in the detection process. To overcome this problem, histogram equalization and other similar methods can be applied [30, 35, 48]. Authors in [48] used Contrast Limited Adaptive Histogram Equalization (CLAHE) which is an advanced version of histogram equalization aiming to reduce over amplification of noise in near constant regions. Additionally, in [35], histogram equalization combined with Perona-Malik filter (PMF) and unsharp masking edge enhancement is applied to facilitate contrast enhancement, edge enhancement, and noise elimination on the entire chest X-ray image dataset. Some literature works are also observed to exclude the faulty images from their dataset [37, 52, 55, 56].

Similar Clinical Manifestations In many cases, viral pneumonia shows similar symptoms as COVID-19 which makes



Fig. 3 X-ray images of some faulty images. a Low Contrast with wire around Image. b
Textual data on top left corner and probes on chest. c Wires over the chest. Images obtained from [26]



it difficult to distinguish them. Additionally, mild COVID-19 cases often show no or mild symptoms, and thus, often indistinguishable from the normal lung images to the naked eye. In the worst-case scenario, these result in low detection probability, i.e., low true positive rate (TPR) and high false negative rate (FNR) for COVID-19 cases. The consequence is that the subjects who are screened as COVID-19 negative in false negative cases may end up contaminating others without attempting for a second test. This suggests that a trade-off should be made between sensitivity (true positive) and specificity (true negative) [57]. In [58], the authors argued that the trade-off should be kept as low as possible to make the model highly sensitive (in contrast to low specificity).

Description of Available Datasets

A large volume of well-labeled data can improve the network quality in deep learning while preventing over-fitting and poor predictions. It is a hard task to collect good-quality data then labeling those accordingly and for uncharted territory like the novel Coronavirus, the hurdles are even bigger. However, time demands to tackle this peril at hand. Therefore, many researchers around the world have been working on creating standard datasets. In this section, we discuss some of these datasets in detail.

COVID-19 Image Data Collection [26]: This publicly available dataset consists of chest X-ray and CT images of individuals suspected with COVID 19 and pneumonia patients. They were collected through doctors, hospitals, and other public sources. 434 images were labeled from the gathered 542 COVID-19 images, and among them, the X-ray and CT images numbered 462 and 80, respectively. There are about 408 Anterior-to-Posterior (AP) or Posterior-to-Anterior (PA) images and 134 Anterior-to-Posterior Supine (AP Supine) images of the patients. The metadata of 380 subjects

marked 276 COVID-19-positive cases where 150 male, 91 female patients, and the rest of them were unlabeled.

Actualmed COVID-19 Chest X-ray Dataset Initiative [59]: A similar dataset of Anterior-to-Posterior (AP) and Posterior-to-Anterior (PA) views of chest X-rays including metadata have been published recently. This open sourced dataset consists of 238 images where cases with COVID-19, no findings and inconclusive images tallied 58, 127, and 53 respectively.

Figure 1 COVID-19 Chest X-ray Dataset Initiative [60]: Another dataset with 55 images of Anterior-to-Posterior (AP) and Posterior-to-Anterior (PA) view of chest X-ray was released with public accessibility. The X-ray images from 48 subjects labeled 10 males and 7 females and the rest remained unlabeled. The dataset enlisted 35 confirmed COVID-19, 3 no findings, and 15 inconclusive images. The age of the subjects ranges from 28 to 77.

COVID-19 Radiography Database [61]: A substantial dataset is made based on the chest X-ray images. This dataset comprises such data of COVID-19-positive patients, normal individuals, and viral pneumonia-infected people. The latest release has Posterior-to-Anterior (PA) view images of 219 COVID-19 positives, 1341 normal, and 1345 viral pneumonia specimens. The Italian SIRM dataset[62], COVID Chest X-ray Dataset by Cohen J et al. [26], 43 different articles and chest X-ray (pneumonia) images database by Mooney P et al. [63] has led considerable facilitation for this dataset.

COVIDx [64]: This dataset was developed by combining the following 5 publicly available dataset: Cohen J et al. [26], ActualMed COVID-19 Chest X-ray Dataset Initiative [59], Figure1 COVID-19 Chest X-ray Dataset Initiative [60], RSNA Pneumonia Detection Challenge dataset [65], and COVID-19 Radiography Database [61]. The augmented



dataset consisted of 13989 images from 13870 individuals. The findings recorded 7966 normal images, 5459 images of pneumonia patients, and 473 images of COVID-19-positive patients.

Augmented COVID-19 X-ray Dataset [66]: This dataset accumulated an equal number (310) of positive and negative COVID-19 images from 5 well-known datasets. They are as follows: COVID Chest X-ray Dataset [26], Italian SIRM [62] radiography dataset, Radiopaedia [67], RSNA Pneumonia Detection Challenge [65], and Chest X-ray Images (pneumonia) [63]. Later using data augmentation methods like flipping, rotating, translation, and scaling, the number was enhanced to 912 COVID-19-positive images and 912 COVID-19-negative images.

COVID-CT Dataset [68]: This CT image-based radiography dataset has been well-acknowledged by radio experts. It consists of 349 and 463 CT images from 216 COVID-19-positive patients and 55 COVID-19-negative patients respectively. As the data collection from two categories implied, the labeling of the data has been done in two classes: COVID-19 positive and COVID-19 negative.

Extensive COVID-19 X-ray and CT Chest Images Dataset [69]: Unlike the other datasets, this contains both X-ray images and CT scans, with a total number of augmented images reaching 17,099. Amongst them, 9544 of them are X-rays, and the rest are CT scans. The dataset has been created for binary classification between COVID and non-COVID. Among the X-ray images, there are 5500 non-COVID images and 4044 COVID images. In the case of the CT scans, there are 2628 non-COVID images and 5427 COVID images.

COVID-19 CT Segmentation Dataset [70]: This dataset consists of 100 axial CT images of more than 40 COVID-positive patients. The images were segmented in three labels by radiologists. They are ground glass, consolidation, and pleural effusion. This dataset provides another nine volumetric CT scans that include 829 slices. Amongst them, 373 have been annotated positive and segmented by a radiologist, with lung masks provided for more than 700 slices.

COVID-19 X-ray Images [71]: This is a dataset in Kaggle that contains both chest X-ray and CT images with the tally of the number of images up to 373. This dataset includes 3 classes, COVID-19, streptococcus pneumonia, and others, including SARS, MERS, and ARDS. The dataset comes with metadata that includes information such as sex, age, medical status of the patients, and other related medical status.

Researchers around the world have made their dataset open for everyone to facilitate the research scopes. The Italian SIRM [62] Covid-19 dataset, a Twitter thread reader [72] with chest X-ray images, Radiopaedia [67], COVID-19 BSTI Imaging Database [73] etc. are some notable dataset which have been made public recently. The Italian SIRM records 115 confirmed COVID-19 cases. The thread reader from Twitter assembled 134 chest X-ray images of 50 COVID-19 cases along with gender, age, and symptom information of each. The Radiopaedia provides chest radiography images of 153 normal, 135 COVID-19 positives, and 485 pneumonia-affected patients. BSTI Imaging Database consists of comprehensive reports and chest radiography images (both X-ray and CT scans) of 59 COVID-19-positive patients. Coronacases.org [74] is a website dedicated to COVID-19 cases, which includes 10 such cases with extensive patient reports. Eurorad.org [75] contains 39 radiography images of COVID-19-positive patients. Images include both X-ray and CT scans along with extensive clinical reports. All the datasets mentioned above are being used for model training either directly or after going through some data augmentation process.

A summary of the discussed datasets is presented in Table 2.

Deep Learning–Based Diagnosis Approaches

In this section, we explore the recent literature on radiography-based COVID-19 diagnosis using deep learning-based methods by arranging it into two groups. The first group includes studies that detect COVID-19 from chest X-rays whereas the latter discuses CT scan-based works.

X-ray-Based Diagnosis

Recently, a lot of work have been published on COVID-19 detection from chest X-ray images. The detection problem is mostly modeled as a classification problem of 3 classes: COVID-19-affected lungs, normal lungs, and pneumonia-affected lungs. Here we discuss some recent works on X-ray based COVID-19 diagnosis grouping them according to the used deep learning approaches.

Transfer Learning There remains a scarcity of standard, large volume dataset to train deep learning models for COVID-19 detection. The existing deep convolutional neural networks like ResNet, DenseNet, and VGGNet have the setbacks of having a deep structure with excessively large parameter sets and lengthy training time. Whereas Transfer Learning (TL) surmounts most of these offsets. In transfer learning, knowledge acquired from the training on one dataset is reused in another task with a related dataset,



Table 2 Available radiography datasets to detect COVID-19. Here, NC means COVID-19 negative

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SI. No.	Dataset	Date of publication	Modality	Class	Description	Works on the dataset	Highest accuracy
_	COVID-19 Image Data Collection [26]	February 15, 2020 ^a	Chest X-ray CT scan	COVID-19	462 X-ray 80 CT scan	Apostolopoulos et al. [55] Narin et al. [76] Minaee et al. [36] LV et al. [48] Punn et al. [49]	99.18% [55]
7	Actualmed COVID-19 Chest X-ray Dataset initiative [59]	April 20, 2020 ⁴	Chest X-ray	COVID-19	238 X-ray	I	I
к	Figure 1 COVID-19 Chest X-ray Dataset Initiative [60]	April 3, 2020 ^a	Chest X-ray	COVID-19	55 X-ray	1	1
4	COVID-19 Radiography Dataset [61]	March 28, 2020	Chest X-ray	COVID-19 Normal Pneumoinia	219 COVID-19 1341 Normal 1345 Pneumonia	Chowdhury et al. [31] Bassi et al. [32] Zhang et al. [77]	98.3% [31]
N	COVIDx [39]	March 19, 2020⁴	Chest X-ray	COVID-19 Pneumonia Normal	473 COVID-19 5459 Pneumonia 7966 Normal	Ucar et al. [38] Farooq et al. [34] Wang et al. [39] Karim et al. [35]	98.26% [38]
9	Augmented COVID-19 X-ray-Images Dataset [66]	March 26, 2020	Chest X-ray	COVID-19 NC (COVID Negative)	912 COVID-19 912 NC	Alqudah et al. [78]	95.2%
٢	Italian SIRM COVID-19 Database[62]	March 3, 2020	Chest CT scan	COVID-19	115 COVID-19 cases	Apostolopoulos et al. [55] Bassi et al. [32] Yamac et al. [40] Apostolopoulos et al. [79] Hall et al. [33]	99.18% [55]
∞	Radiopaedia[67]	September 8, 2020 ⁶	Chest X-ray CT scan	COVID-19	135 COVID-19 485 Pneumonia 153 Normal	Apostolopoulos et al. [55] Yamac et al. [40] Hall et al. [33]	99.18% [55]



Table 2	(continued)						
SI. No.	Dataset	Date of publication	Modality	Class	Description	Works on the dataset	Highest accuracy
6	COVID-CT Dataset [68]	March 28, 2020^a	Chest CT scan	COVID-19 Normal	349 COVID-19 463 Normal	He at al. [80]	[08] %98
10	Thread reader [72]	March 28, 2020	Chest X-ray	COVID-19	50 COVID-19 cases	Yamac et al. [40]	95.90% [40]
=	Extensive COVID-19 X-ray and CT Chest Images Dataset [69]	June 13, 2020	Chest X-ray CT scan	COVID-19 Normal	9471 COVID 8128 NC		
12	COVID-19 CT segmentation dataset [70]	April 13, 2020	Chest CT scan	Positive (COVID-19) Negative	373 Positive 456 Negative	Chen et al. [81]	89% [81]
13	COVID-19 X-ray Images [71]	May 15, 2020	Chest X-ray CT scan	COVID-19 Streptococcus Other	309 COVID 17 Streptococcus 47 Other		1
41	COVID-19 BSTI Imaging Database [73]	September 8, 2020^b	Chest X-ray CT scan	COVID-19 Positive	59 COVID-19 cases	Maghdid et al. [82]	94.1% [82]
15	Coronacases.org [74]	September 8, 2020^b	Chest CT scan	COVID-19 cases	10 COVID-19 Positive cases		1
16	Eurorad.org [75]	September 8, 2020 ^{<i>b</i>}	Chest X-ray CT scan	COVID-19 Positive	39 COVID-19 cases		

^aFirst commit ^bLast accessed

yielding improved performance and faster convergence. In the case of COVID-19 detection, the models are often pre-trained on ImageNet [19] dataset and then fine-tuned with the X-ray images. Many researchers have used TL after pre-training their network on an existing standard dataset [30–40, 48, 49, 55, 76, 77, 79, 83, 84]. Some works even pre-trained their model twice. For example, ChexNet [85] is one such network with twofold pre-train on ImageNet [86] and ChestX-ray-14 [87].

Ensemble Learning Ensemble learning uses an augmentative learning technique that combines predictions from multiple models to generate more accurate results. It improves the model prediction results by reducing the generalization error and variance. In [88], ensemble learning technique is used by combining 12 models (Resnet-18,50,101,152, WideResnet-50,101, ResNeXt-50,101, MobileNet-v1, Densenet-121,169,201) to get better results. Karim et al. [35] have also ensembled 3 models (ResNet18, VGG19, DenseNet161). However, authors in [37], used this a bit differently by using only one model (i.e., ResNet18), then fine-tuning it with three different datasets and finally ensembling the three networks to get the final result. Ensemble learning has contributed significantly towards achieving an accurate result for COVID-19 detection.

Domain Adaptation Domain adaptation supports the modeling of a relatively new target domain by adapting learning patterns from a source domain, which has similar characteristics as the target domain. Chest X-ray image of a COVID-19 patient has a different distribution but similar characteristics as that of pneumonia; thus, Domain Adaptation technique can be used. This technique was used by Zhang et al. [77] for creating COVID-DA where the discrepancy of data distribution and task differences was handled by using feature adversarial adaptation and a novel classifier scheme, respectively. Employing this learning method marked a noticeably improved result in detecting COVID-19.

Cascaded Network Radiography-based COVID-19 detection suffers from the data scarcity. Introducing cascaded network architecture in a small dataset facilitates dense neurons in the network while avoiding the overfitting problem [89]. LV et al. [48] cascaded two networks (ResNet-50 and DenseNet-169) to classify COVID-19 samples. After a subject got classified as viral pneumonia from the 3 classes (normal, bacterial, and viral pneumonia) using ResNet, it was fed into DenseNet169 for the final classification as COVID-19. The infectious regions were concentrated on with an attention mechanism technique Squeeze-Excitation (SE) [90]. Contrast Limited Adaptive Histogram Equalization improved their image quality, and an additional Module Excitation (MoEx) [91] with the two networks enhanced

the imaging features. Both the cascaded networks, ResNet-50 and DenseNet-50 gained high accuracy of 85.6% and 97.1%, respectively.

Other Approaches Wang et al. [39] designed the COVID-Net architecture optimizing the human-machine collaborative design strategy. They also fabricated a customized dataset for the network training. Lightweight design pattern, architectural diversity, and selective long-range connectivity supported its reliability with an accuracy of 93.3% for detecting COVID-19. Optimizing the COVIDx dataset with Data Augmentation and pretraining the model with ImageNet also contributed to the high accuracy. Ozturk et al. [92] proposed Dark-COVIDNet evolved from DarkNet-19. It boasted 17 convolutional layers optimizing different filtering on each layer. Punn et al. [49] used NASNetLarge to detect COVID-19. Both proposed models performed well.

Computed Tomography-Based Diagnosis

CT scan images are scarce as it is expensive and unavailable in many parts of the world. CT scan images still have shown better performance in COVID-19 detection as it provides more information and features than X-ray. Many works have used deep learning for segmentation of CT images [81, 93, 94]. Authors in [52] even determined the severity of COVID-19 case followed by the conversion time from mild to severe case. In [51–54, 56, 93–95], authors used 3D images as input to detect the infectious regions from COVID-19 patients. Most works applied machine learning based methods. Here we discuss some of these works clustered according to the algorithms used.

Joint Regression and Classification, Support Vector Machine Machine learning (ML) algorithms, such as Logistic Regression (LR) and Support Vector Machine (SVM), hold superiority over CNN if we probe the complicacy of learning millions of parameters and ease of use. Even though LR and SVM are not that efficient in learning very high-dimensional features, the high-definition CT scan images redress the need for it in this application. So, it can be stated that LR and SVM provide effective results in COVID-19 detection with some attributive advantages. For example, in [50], authors formed four datasets by taking various patch sizes from 150 CT scans. For extracting features, they applied Grey Level Co-occurrence Matrix (GLCM), Local Directional Pattern (LDP), Grey Level Run Length Matrix (GLRLM), Grey-Level Size Zone Matrix (GLSZM), and Discrete Wavelet Transform (DWT) algorithms to the different patches of the images before applying SVM for COVID-19 classification with GLSZM, which resulted in a better performance. In [52], authors proposed a joint regression and classification



algorithm to detect the severe COVID-19 cases and estimate conversion time from mild to severe manifestation of the disease. They used logistic regression for classification and linear regression for estimating the conversion time. They also attempted to reduce the influence of outliers and class imbalance, by giving more weight to the important class. Both of the proposed models showed satisfactory performance.

Random Forest and Deep Forest Random forest algorithm relies on building decision trees and ensembling those to make predictions. Its simplicity and diversity made it applicable to a wide range of classification applications while reducing computational complexity. In [96], an infection Size-Adaptive Random Forest (iSARF) method is proposed where the location-specific features are extracted using VB-Net toolkit [97] and the optimal feature set is obtained using the Least Absolute Shrinkage and Selection Operator (LASSO). The selected features are then fed into the random forest algorithm for the classification. In [98], the authors proposed an Adaptive Feature Selection guided Deep Forest (AFS-DF) for the classification task. They also extracted the location-specific features using VB-Net and utilized these features to construct an N random forest. The adaptive feature selection was opted for minimizing redundancy before cascading multiple layers to facilitate the learning of a deep, distinguishable feature representation for COVID-19 classification.

Multi-view Representation Learning In classification and regression tasks single-view learning is commonly used as a standard for its straightforward nature. However, multi-view representation learning is getting attention in recent times because of its ability to acquire multiple heterogeneous features to describe a given problem with improved generalization. In [54], authors employed multi-view representation learning for COVID-19 classification among community-acquired pneumonia (CAP). V-Net[99] is used to extract different radiomics and handcrafted features from CT images before applying a latent representation based classifier to detect COVID-19 cases.

Hypergraph Learning In [100], a hypergraph was constructed where each vertex represents COVID-19 or community-acquired pneumonia (CAP) case. They extracted both radiomics and regional features; thus, two groups of hyperedges were employed. For each group of hyperedges, *k*-nearest neighbor was applied by keeping each vertex as centroid and found related vertices depending on features. Then the related vertices connected together by making edges. It is not impossible to have noisy, unstable or abnormal image in a dataset. To keep away these images, two types of uncertainty score were calculated: aleatoric uncertainty for

noisy or abnormal data and epistemic uncertainty for the model's inablity to distinguish two different cases. These uncertainty score referred to the quality of image and used as the weight of vertices. Label propagation algorithm [101] was run on the hypergraph generating a label propagation matrix, which was then used to test new cases.

CNN Architecture The quality of CT images coupled with the expansive computational power of recent technologies make deep learning the most potent candidate for COVID-19 detection. Many state-of-the-art deep CNN architectures like ResNet, DenseNet, and Alexnet have already been used for COVID-19 diagnosis [80-82, 94, 95]. These models are generally pre-trained on the ImageNet dataset before being fine-tuned using CT image datasets to avoid learning millions of parameters from scratch. However, the insufficiency of chest CT image data of COVID-19affected patients results in the augmentation of training data in many cases [51, 80, 81, 95]. In [53], authors proposed COVIDCT-Net where they used BCDU-Net [102] to segment infectious areas before feeding it into a CNN for classification. DeCovNet was proposed in [51] where a pretrained U-Net[103] is used to segment the 3D volume of the lung image before being fed into a deep CNN architecture.

Attention Mechanism To find the infectious regions better, attention mechanism was applied in [56, 81, 93, 94] and others. Authors in [81] used Locality Sensitive Hashing Attention [104] in their residual attention U-Net model. In [93], an online 3D class activation mapping (CAM)[105] was used with ResNet-34 architecture. Authors in [94] applied Feature Pyramid Network (FPN)[106] with ResNet-50 to extract top k-features of an image before feeding them to an attention module (Recurrent Attention CNN [107]) to detect the infections regions from the images. In [56], authors used location attention with the model learning the location of infectious patches in the image by calculating relative distance-from-edge with extra weight.

The details of the works that used deep learning for diagnosing COVID-19 is given in Table 3.

Image Prepossessing

Image prepossessing is a crucial step in radiography image classification with deep learning, simply because the smallest details in the images can veer the model to learn completely different features and make disparate predictions. Especially when it comes to COVID-19 images, where it is already difficult for models to learn to classify from two very similar classes (pneumonia and COVID-19), preprocessing becomes an important step. This is not only credited to the similarity between the classes but also the imbalance



Table 3 Overview of some COVID-19 detection approaches using deep learning. Trn train, Val validation, Tst test, TV train and validation, CV cross-validation, ACC accuracy, PRE precision, REC recall, SEN sensitivity, SPE specificity, F1-SCR F1 score, CM calculated from Confusion Matrix, AUC area under curve

SI. No.	Study	Method	Modality	Class	Dataset	Train-test split	Performance
	LV et al. [48]	Casecade SEMENet	Chest X-ray	COVID-19, Pneumonia, Normal	[26, 108]	Tm: 6386 images Val: 456 images Tst: 456 images	ACC: 97.14% F1-SCR: 97%
2	Bassi et al. [32]	CheXNet [85]	Chest X-ray	COVID-19, Pneumonia, Normal	[26, 61, 62, 108]	Tm: 80% Val: 20% Tst: 180 images	ACC: 97.8% PRE: 98.3% REC: 98.3%
С	Yamac et al. [40]	CSEN (CheXNet [85])	Chest X-ray	COVID-19, Viral, and Bacterial Pneumonia	[62, 63, 67, 72]	Stratified 5-fold CV	ACC: 95.9% SEN: 98.5% SPE: 95.7%
4	Zhang et al. [77]	COVID-DA (Domain Adaptation)	Chest X-ray	COVID-19, Pneumonia, Normal	[61, 65]	Trn: 10,718 images Tst: 945 images	AUC: 0.985 PRE: 98.15% REC: 88.33% F1-SCR: 92.98%
ĸ	Goodwin et al. [88]	12 Models Ensembled	Chest X-ray	COVID-19, Normal, Pneumonia	[26]	Trn: 80% Val: 10% Tst: 10%	ACC: 89.4% PRE: 53.3% REC: 80% F1-SCR: 64%
9	Misra et al. [37]	ResNet-18	Chest X-ray	COVID-19, Normal, Pneumonia	[26, 65]	Trn: 90% Tst: 10%	ACC: 93.3% PRE: 94.4% REC: 100%
7	Abbas et al. [30]	DeTraC- ResNet18	Chest X-ray	COVID-19, Normal, SARS	[26]	Tm: 70% Tst: 30%	ACC: 95.12% SEN: 97.91% SPE: 91.87%
∞	Chowdhury et al. [31]	SqueezNet (Best Model)	Chest X-ray	COVID-19, Viral Pneumonia, Normal	[61]	5-fold CV	ACC: 98.3% PRE: 100% REC: 96.7% F1-SCR: 100%



Table 3 (c	(continued)						
SI. No.	Study	Method	Modality	Class	Dataset	Train-test split	Performance
6	Farooq et al. [34].	COVID-ResNet	Chest X-ray	COVID-19, Normal. Viral, and Bacterial Pneumonia	[33]	Tm: 13,675 images Tst: 300 images	ACC: 96.23% PRE: 100% REC: 100% F1-SCR: 100%
10	Alqudah et al. [78]	AOCT-Net	Chest X-ray	COVID-19, NonCovid-19	[99]	10-fold CV	ACC: 95.2% SEN: 93.3% SPE: 100% PRE: 100%
Π	Hall et al. [33]	3 Models Ensembled	Chest X-ray	COVID-19, Pneumonia	[26, 62, 67]	10-fold CV	ACC: 91.24% SPE: 93.12% SEN: 78.79%
12	Hemdan et al. [109]	COVIDX-Net	Chest X-ray	COVID-19, Normal	[26]	Tm: 80% Tst: 20%	ACC: 90% PRE: 83% REC: 100% F1-SCR: 91%
13	Apostolo- poulos et al. [79]	VGG19 (Best Model)	Chest X-ray	COVID-19, Normal, Pneumonia	[26, 62, 65]	10-fold CV	ACC: 93.48% SEN: 92.85% SPE: 98.75%
41	Apostolo- poulos et al. [55]	Mobile- Netv2.0 (from scratch)	Chest X-ray	COVID-19, nonCOVID-19	[26, 62, 65, 67]	10-fold CV	ACC: 99.18% SEN: 97.36% SPE: 99.42%
15	Karim et al. [35]	Deep COVID- Explainer	Chest X-ray	COVID-19, Normal. Viral, and Bacterial Pneumonia	[62, 65, 110]	5-fold CV	ACC: 96.77% (CM) PRE: 90% REC: 83%
16	Majeed et al. [83]	CNNX	Chest X-ray	COVID-19, Normal. Viral, and Bacterial Pneumonia	[26, 62, 65, 108]	Tm: 5327 images Tst: 697 images	SEN: 93.15% SPE: 97.86%



Table 3 (co	(continued)						
SI. No.	Study	Method	Modality	Class	Dataset	Train-test split	Performance
17	Minaee et al. [36]	SqueezNet (Best Model)	Chest X-ray	COVID-19, nonCOVID-19	[26, 111]	Tm: 2496 images Tst: 3040 images	ACC: 97.73% (CM) SEN: 97.50% SPE: 97.80%
18	Narin et al. [76]	ResNet50 (Best Model)	Chest X-ray	COVID-19, Normal	[26, 63]	5-fold CV	ACC: 98% PRE: 100% REC: 96% SPE: 100%
19	Punn et al. [49]	NASNetLarge (Best Model)	Chest X-ray	COVID-19, Normal, Pneumonia	[26, 65]	Tm: 1266 images Val: 87 images Tst: 108 images	ACC: 96% PRE: 88% REC: 91% SPE: 94%
20	Ozturk et al. [92]	DarkCovidNet	Chest X-ray	COVID-19, Normal, Pneumonia	[26, 112]	5-fold CV	ACC: 87.20% PRE: 89.96% REC: 92.18%
21	Sethy et al. [84]	ResNet50 +SVM	Chest X-ray	COVID-19, Normal	[26, 63]	Tm: 60% Val: 20% Tst: 20%	ACC: 95.38% F1-SCR: 95.52% MCC: 90.76%
22	Wang et al. [64]	COVIDNet	Chest X-ray	COVID-19, Normal, Pneumonia	[39]	Tm: 13,675 images Tst: 300 images	ACC: 93.30% PRE: 98.90% REC: 91%
23	Ucar et al. [38]	COVIDiag-nosisNet	Chest X-ray	COVID-19, Normal, Pneumonia	[26, 39, 108]	Tm: 80% Val: 10% Tst: 10%	ACC: 98.26% PRE: 98.26% REC: 98.26% SPE: 99.13%
24	Sun et al. [98]	AFS-DF (Deep-Forest)	Chest CT scan	COVID-19, Pneumonia	Not available	5-fold CV	ACC: 91.79% SPE: 89.95% SEN: 93.05% AUC: 96.35%
25	Javaheri et al. [53]	COVID-CTNet	Chest CT scan	COVID-19, Pneumonia, Normal	Not available	Tm: 90% Val: 10% Tst: 20 cases	ACC: 90.00% SEN: 83.00% SPE: 92.85%



Table 3 (con	(continued)						
SI. No.	Study	Method	Modality	Class	Dataset	Train-test split	Performance
26	Kang et al. [54]	Multiview Representaion Learning	Chest CT scan	COVID-19, Pneumonia	Not available	Trn: 70% Tst: 30%	ACC: 95.5% SEN: 96.6% SPE: 93.2%
72	Donglin et al. [100]	UVHL (Hypergraph Leaming)	Chest CT scan	COVID-19, Pneumonia	Not available	10-fold CV	ACC: 89.79% SEN: 93.26% SPE: 84% PPV: 90.06%
78	Zhu et al. [52]	Joint regression and Classification	Chest CT scan	COVID-19, Severity estimation	Not available	5-fold CV	ACC: 85.91%
29	Ouyang et al. [93]	Attention ResNet34 +Dual Sampling	Chest CT scan	COVID-19, Pneumonia	[65]	TV set: 2186 images Tst: 2796 images	AUC: 0.944 ACC: 87.5% SEN: 86.9% SPE: 90.1% F1-SCR: 82.0%
30	Chen et al. [81]	Residual Attention U-Net	Chest CT scan	COVID-19 (Segmentation)	[62]	10-fold CV	ACC: 89% PRE: 95% DSC: 94%
31	He et al. [80]	DenseNet169 (Self-supervised Transfer Learning)	Chest CT scan	COVID-19, nonCOVID-19	[88]	Tm: 60% Val: 15% Tst: 25%	ACC: 86% F1-SCR: 85% AUC: 94%
32	Maghdid et al. [82]	Modified AlexNet	Chest CT scan	COVID-19, Normal	[26, 73]	Tm: 50% Val: 50% Tst: 17 images	ACC: 94.1% SPE: 100% SEN: 90%
33	Maghdid et al. [82]	Modified AlexNet	Chest X-ray	COVID-19, Normal	[26, 73]	Tm: 50% Val: 50% Tst: 50 images	ACC: 94% SPE: 88% SEN: 100%



Table 3 (co	(continued)						
SI. No.	Study	Method	Modality	Class	Dataset	Train-test split	Performance
34	Butt et al. [56]	ResNet18 +Location Attention	Chest CT scan	COVID-19, Normal, Viral Pneumonia	Not available	TV set: 85.4% Tst: 14.6%	ACC: 86.7% PRE: 86.7% REC: 81.3% F1-SCR: 83.90%
35	Song et al. [94]	DRE-Net	Chest CT scan	COVID-19, Normal, Pneumonia	Not available	Tm: 60% Val: 10% Tst: 30%	ACC: 86% PRE: 79% REC: 96% F1-SCR: 97%
36	Zheng et al. [51]	DeCovNet	Chest CT scan	COVID-19	Not available	Tm: 499 images Tst: 131 images	ACC: 90.10% SEN: 90.70% SPE: 91.10%
37	Barstugan et al. [50]	GLSZM+SVM (Best Model)	Chest CT scan	COVID-19, nonCOVID-19	[62]	10-fold CV	ACC: 98.71% SEN: 97.56% SPE: 99.68% PRE: 99.62%
38	Shi et al. [96]	iSARF (Random Forest)	Chest CT scan	COVID-19 Pneumonia	Not available	5-fold CV	ACC: 87.9% SEN: 90.7% SPE: 83.3%
39	Gozes et al. [95]	2D and 3D CNN (ResNet-50)	Chest CT scan	COVID-19 Normal	Not available	Tm: 50 patients Tst: 157 patients	SEN: 98.2% SPE: 92.2% AUC: 0.996



present in the datasets currently available. In this section, we discuss the various preprocessing methods used by different works in COVID-19 detection. A summary of the discussed techniques is also given in Table 4.

Data Augmentation Imbalance dataset is a prevalent issue in COVID-19 classification due to the scarcity of COVID-19 images. As discussed before in Section "Challenges," it can cause wrongful predictions. One way to solve this is data augmentation, which is a strategy to increase the number of images by using various image transformation methods such as rotation, scaling, translation, flipping, and changing the pixels' brightness. It creates diversity in images without actually collecting new ones. It is one of the simplest ways to deal with imbalanced datasets, and thus, is used in most of the works regarding COVID-19 classification, the list of which, is given in Table 4.

Class Resampling This is a popular method and has seen much success in overcoming data imbalance. In [113], authors randomly omitted images from the majority class (viral pneumonia, bacterial pneumonia, and normal) to balance the dataset, which is called random undersampling or RUS. The opposite is done in [37, 49, 93] where the authors randomly oversampled the minority class (COVID-19), which is called random oversampling or ROS. Both methods are widely used for their simplicity and effectiveness. In [114], authors experimented several undersampling and oversampling methods with multiclass and hierarchical classification models.

Focal Loss and Weighted Loss Focal loss [115] is another method to solve the issue of class imbalance. It puts more weight on the hard to classify objects and decreases the weight on easy and correct predictions. A scaling factor is added to the cross-entropy loss function, and it decreases as the confidence in a prediction goes up. This was used in [77] for addressing the class imbalance issue. On the other hand, the weighted loss function is a more common technique to balance the data among different classes, which puts more weight on the minority class and less on the majority. In [49, 88], and [52], authors have used weighted loss function and obtained good results.

Image Quality Enhancement In medical image analysis, every pixel of the image is important for the proper diagnosis of the disease. This is even more applicable when it comes to diagnosing COVID-19 as it bears so much similarity with pneumonia. Thus, image quality enhancement is applied in many works regarding COVID-19 detection. For this, the most prominent technique is increasing the contrast of the image that allows the features to stand out more. Histogram equalization used in [30, 35, 48, 49, 51, 80, 93], is an example of that. In [38, 51, 80], authors applied brightness adjustment for enhancing image quality. In [35, 49], authors applied Perona-Malik Filter [117] and Adaptive total variation [118], respectively for noise removal. Unsharp masking is used in [35] for edge sharpening.

Image Segmentation Image segmentation involves dividing an image into segments to simplify image analysis

Table 4 A summary of different prepossessing approaches employed in deep learning-based automated detection of COVID-19

Preprocessing	Approach	Method	Reference
Addressing data imbalance	Data augmentation	Rotation, Translation, Cropping, Flipping etc.	[30–40, 51, 55, 66, 79, 81, 88, 95]
	Class resampling	Undersampling	[113, 114]
		Oversampling	[37, 49, 93, 114]
	Loss function	Focal Loss[115]	[51]
		Weighted Loss[116]	[49, 52, 88]
Image segmentation	Segmentation model	VB-Net[97]	[52, 93, 96, 98, 100]
		U-Net[103]	[48, 51, 95]
		V-Net[99]	[54]
		BCDU-Net[102]	[53]
Image quality enhancement	Contrast enhancement	Histogram Equalization	[30, 35, 48, 49, 51, 80, 93]
	Brightness changing	Adding or sub- tracting every pixel by a constant value	[38, 51, 80]
	Noise removal	Perona-Malik filter [117]	[35]
		Adaptive Total Variation [118]	[49]
	Edge sharpening	Unsharp Masking	[35]



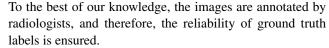
and focus only on the important part. Thus, in the case of COVID-19 detection, image segmentation is often a critical preprocessing step that can allow the model to learn from only the affected organ, namely, the lungs, to provide more accurate predictions. One of the most well-known algorithms for medical image segmentation is U-net [103], which has a superior design of skip connections between different stages of the network (used in [81] and [95]). Probably, the most famous derivations of U-Net is VB-Net [97], which is used in [52, 96, 98], and [100]. There are other architectures available for segmentations such as V-Net [99] used in [54], BCDU-Net [102] used in [53] and other works like [56, 81] and [94] that have incorporated image segmentation into their deep models rather than going through segmentation as a preprocessing step.

Comparative Analysis

As we have seen so far, researchers' continuous efforts have already elicited deep learning-based automated diagnosis of COVID-19 as a potential research avenue. However, despite having several publicly accessible datasets (See "Description of Available Datasets"), no "benchmark" dataset has been released yet that can be used to assess the performance of different methods' ability to detect COVID-19 using the same standard. Therefore, different authors reported the performance of their method based on different datasets and evaluation protocols. Being motivated by the urge to compare the models on the same scale, here we present a comparative quantitative analysis of 315 deep models that consists of the combinations of 15 convolutional neural network (CNN) models and 21 classifiers using our customized dataset. Note that, we release our dataset along with train-test split and models to be publicly available. ¹

Dataset Description

To conduct the analysis, we compile a dataset of our own that includes X-ray images from 4 different data sources, [26, 59, 60], and [61]. The dataset contains 7879 distinct images in total for 3 different classes: COVID-19, normal, and pneumonia, where the pneumonia class has both viral and bacterial infections. However, we have not taken all the available images of the 4 datasets, but only the frontal X-ray (Posterior-to-Anterior) images, leaving out the side views. We selected datasets from the above sources as they are fully accessible to the research community and the public.



The dataset is split into training (60%) and test (40%) sets using Scikit-learn's train_test_split module, which, given the split percentage, randomly splits the data between the two sets due to a random shuffle. The shuffling algorithm uses D.E. Knuth's shuffle algorithm (also called Fisher-Yates) [119] and the random number generator in [120]. The split is also stratified to ensure the presence of the percentage of class samples in each set. The distributions of samples across the classes and train-test set are shown in Table 5.

Method

The overall detection problem is posed as a multi-class classification problem that consists of two major components: feature extraction and learning a classifier.

Feature Extraction For the feature extraction step, we utilize transfer learning and select a CNN model which has been trained on ImageNet [86], a dataset containing 1.2 million images of 1000 categories. The last fully connected layer, classification layer, and softmax layer are removed from the model and the rest is considered as a feature extractor that computes a feature vector for each image.

Learning a Classifier We forward the extracted features to a learning model, which is then trained using 5-fold cross-validation on the training set. From Table 1, we notice an imbalance of sample distribution across the classes (COVID-19: 8.7%, normal: 37.1%, and pneumonia: 52.2%). Therefore, to analyze the deep models' performance at the presence of class imbalance problem, we have done experimentation with three different approaches: a weighted cost function, upsampling the training dataset, and downsampling the training dataset.

 Weighted Cost Function: Being inspired by the work in [116], we have applied a weighted cost function which ensures each class to be given same relative

Table 5 Data distribution of chest X-ray images among 3 different classes: COVID-19, normal, and pneumonia

Class	Total no. of samples	Training data	Test data
COVID-19	683	410 (5.2%)	273 (3.5%)
Normal	2924	1754 (22.3%)	1170 (14.8%)
Pneumonia	4272	2563 (32.5%)	1709 (21.7%)
Total	7879	4727 (60%)	3152 (40%)



¹The dataset and models are available at https://github.com/rgbnihal2/COVID-19-X-ray-Dataset

importance by assigning more weight on the minority class and less on the majority.

$$L = W_{\text{normal}} \times L_{\text{normal}} + W_{\text{pneumonia}} \times L_{\text{pneumonia}} + W_{\text{COVID}} \times L_{\text{COVID}}$$
 (1)

where L is total categorical cross-entropy loss. L_{normal} , $L_{\text{pneumonia}}$, and L_{COVID} denote the cross-entropy losses for normal, pneumonia, and COVID, respectively. Weight of each class is calculated using the following formula:

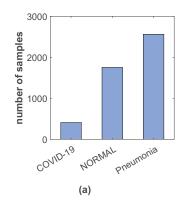
$$w_i = \frac{n}{k \times n_i} \tag{2}$$

where w_i is the weight of class i, n is the total number of observations in the dataset, n_i is the total number of observations in class i, and k is the total number of classes in the dataset.

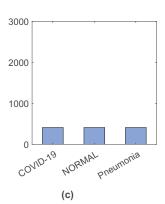
- **Upsampling:** Upsampling is the method of randomly replicating samples from the minority class. First, we separate observations from each class. Next, we resample the minority classes (normal and COVID-19), setting the number of samples to match the majority class (pneumonia). Thus, the ratio of the three classes becomes 1:1:1 as shown in Fig. 4b.
- **Downsampling:** Downsampling involves randomly eliminating samples from the majority class to prevent from dominating the learning algorithm. It has a similar process, like upsampling. After separating samples from each class, we resample the majority classes (pneumonia and normal), setting the number of samples to match that of the minority class (COVID-19). After that, the ratio of the three classes becomes 1:1:1 (see Fig. 4c).

Evaluation Process 15 CNN architectures are used to extract feature vectors, and these are fed to 21 classifiers. Therefore, $15 \times 21 = 315$ models are experimented with a general cross-entropy and the weighted cross-entropy given in (1). Thus, a total of 630 models are deployed with the dataset and ranked according to the performance metrics. In

Fig. 4 Distribution of samples among 3 different classes in a original, **b** upsampled, and **c** downsampled training dataset



3000 2000 1000 NORMAL Puenwouig (b)



addition, the performance of the top-5 models is evaluated with the upsampled and downsampled training dataset. This study thus includes a total of 640 models for performance

Following is the list of CNN models that have been used to assess the feature extraction step (number of features extracted from each image is given in the parenthesis):

Alexnet (4096) [121],

benchmarking.

- Xception (2048) [122],
- InceptionV3 (2048) [123],
- InceptionResNetV2 (1536) [124],
- VGG16 (4096) [125],
- ResNet50 (2048) [126],
- MobileNetV2 (1280) [127],
- DarkNet53 (1024) [128],
- DarkNet19 (1000) [129],
- GoogleNet (1024) [130],
- DenseNet-201 (1920) [131],
- ShuffleNet (544) [132],
- NasNetMobile (1054) [133],
- ResNet18 (512) [126], and
- VGG19 (4096) [125].

The architectures of these models are shown in Table 6. List of classifiers we tried for the comparative analysis:

- Tree-Based[134]: Fine, Medium, Coarse
- **Discriminant Analysis**[135]: Linear, Quadratic
- **Support vector machine**[136]: Linear, Quadratic, Cubic, Coarse Gaussian, Medium Gaussian
- K-Nearest Neighbor [137]: Fine, Medium, Cubic, Cosine, Coarse, Weighted
- Ensemble: BoostedTree[138], BaggedTree[139], SubspaceDiscriminant, Subspace KNN[140], RusBoostedTree

The overall performance is evaluated using accuracy, COVID-19 accuracy, weighted precision, weighted recall, weighted specificity, and weighted F1-score. Here we have used weighted average of performance metrics instead of

Table 6 Archit mance benchma visualization)

visualization)	ing in COVID-19 de	`		Layer	Patch size/stride	Depth	Output size
Layer	Patch size/stride	Depth	Output size	Convolution	3×3×512/1	2	28×28×512
() D . M . 50					$1\times1\times512/1$	1	
(a) ResNet-50				Max pool	$3\times3/2$	1	14×14×512
Convolution	7×7×64/2	1	112×112	Convolution	3×3×512/1	2	14×14×512
Max pool	3×3/2	1	56×56		$1 \times 1 \times 512/1$	1	
Convolution	1×1×64	3	56×56	Max pool	3×3/2	1	7×7×512
	3×3×64			Fully connected	-	2	1×4096
Convolution	1×1×256 1×1×128	4	28×28	Softmax	_	1	1×1000
Convolution	3×3×128	4	20 X 20	Sortmax		1	1 × 1000
	1×1×512			(d) AlexNet			
Convolution	1×1×312 1×1×256	6	14×14	Convolution	11×11/4	1	55×55×96
Convolution	3×3×256	U	14 × 14		3×3/2	1	27×27×96
	1×1×1024			Max pool			
Convolution	1×1×1024 1×1×512	3	7×7	Convolution	5×5/1	1	27×27×256
Convolution	3×3×512	3	1 ~ 1	Max pool	3×3/2	1	13×13×256
	$1\times1\times2048$			Convolution	3×3/1	1	$13 \times 13 \times 384$
Average pool	-	1	1×1000	Convolution	3×3	1	$13\times13\times384$
Fully connected		-	17/1000	Convolution	3×3	1	$13 \times 13 \times 256$
Softmax				Max pool	$3\times3/2$	1	$6\times6\times256$
				Fully connected	-	2	1×4096
(b) GoogleNet				Softmax	-	1	1×1000
Convolution	$7 \times 7/2$	1	$112\times112\times64$				
Max pool	$3\times3/2$	0	$56 \times 56 \times 64$	(e) DarkNet-53			
Convolution	$3\times3/1$	2	$56 \times 56 \times 192$	Convolution	$3\times3\times32/1$	1	256×256×32
Max pool	$3\times3/2$	0	$28 \times 28 \times 192$	Convolution	$3\times3\times64/2$	1	$128 \times 128 \times 64$
Inception(3a)	-	2	$28 \times 28 \times 256$	Convolution	$1\times1\times32/1$	1	128×128
Inception(3b)	-	2	$28 \times 28 \times 480$	Convolution	3×3×64/1		
Max pool	$3\times3/2$	0	$14 \times 14 \times 480$	Residual	-		
Inception(4a)	-	2	$14 \times 14 \times 512$	Convolution	3×3128/2	1	64×64
Inception(4b)	-	2	$14 \times 14 \times 512$	Convolution	1×1×64/1	2	64×64
Inception(4c)	-	2	$14 \times 14 \times 512$	Convolution	3×3×128/1	-	01,701
Inception(4d)	-	2	$14 \times 14 \times 528$	Residual	3 × 3 × 120/1		
Inception(4e)	-	2	$14 \times 14 \times 832$	Convolution	3×3×256/2	1	32×32
Max pool	$3\times3/2$	0	$7 \times 7 \times 832$				
Inception(5a)	-	2	$7 \times 7 \times 832$	Convolution	1×1×128/1	8	32×32
Inception(5b)	-	2	$7 \times 7 \times 1024$	Convolution	$3\times3\times256/1$		
Average pool	$7 \times 7/1$	0	$1\times1\times1024$	Residual	-		
Dropout(40%)	-	0	1×1×1024	Convolution	$3\times3\times512/2$	1	16×16
Linear	-	1	1×1×1024	Convolution	$1\times1\times256/1$	8	16×16
Softmax	-	0	$1\times1\times1024$	Convolution	$3\times3\times512/1$		
(c) VGG-16				Residual	-		
Convolution	3×3×64/1	2	224×224×64	Convolution	$3\times3\times1024/2$	1	8×8
Max pool	3×3×04/1 3×3/2	1	112×112×64	Convolution	$1\times1\times512/1$	4	8×8
Convolution	3×3×128/1	2	112×112×04 112×112×128	Convolution	$3 \times 3 \times 1024/1$		
Max pool	3×3/2	1	56x56x128	Residual	-		
Convolution	$3 \times 3 \times 256/1$	2	56×56×256	Average pool	-	1	1×1000
	$1\times1\times256/1$	1	,	FC			
3.6	22/2	-	2020256	-			



Max pool

 $3 \times 3/2$

 $28{\times}28{\times}256$

Softmax

ntinued)

Layer	Patch size/stride	Depth	Output size	
(f) ShuffleNet				
Convolution	$3\times3/2$	2	112×112	
Max pool	$3\times3/2$	2	56×56	
Stage 2	-/2	1	28×28	
	-/1	3	28×28	
Stage 3	-/2	1	14×14	
	-/1	7	14×14	
Stage 4	-/2	1	28×28	
	-/1	3	7×7	
Global pool	7×7	-	1×1	
FC	-	-	1×1000	
(g)MobileNetV2				
Convolution	$3\times3/2$	1	112×112×32	
Bottleneck	-/1	1	112×112×16	
Bottleneck	-/2	2	$56 \times 56 \times 24$	
Bottleneck	-/2	3	$28 \times 28 \times 32$	
Bottleneck	-/2	4	$14 \times 14 \times 64$	
Bottleneck	-/1	3	$14 \times 14 \times 96$	
Bottleneck	-/2	3	$7 \times 7 \times 160$	
Bottleneck	-/1	1	$7 \times 7 \times 320$	
Convolution	$1 \times 1/1$	1	$7\times7\times1280$	
Average pool	7×7/-	1	$1\times1\times1280$	
Convolution	$1 \times 1/1$	-	k	
(h) DenseNet-201				
Convolution	7×7/2	1	112×112	
Max pool	$3\times3/2$	1	56×56	
Dense block (1)	1×1	6	56×56	
	3×3			
Transition layer (1)	1×1	1	56×56	
	$3\times3/2$, avg pool		28×28	
Dense block (2)	1×1	12	28×28	
	3×3			
Transition layer (2)	1×1	1	28×28	
	$3\times3/2$,avg pool		14×14	
Dense block (3)	1×1	48	14×14	
	3×3			
Transition layer (3)	1×1	1	14×14	
	$3\times3/2$, avg pool		7×7	
Dense block (4)	1×1	32	7×7	
	3×3			
Average pool	7×7	1	1×1	
FC	1000			
Softmax	-			

Table 6 (continued)

Layer	Patch size/stride	Depth	Output size
(i) Xception			
Convolution	$3\times3\times32/2$	1	$149{\times}149{\times}32$
	$3\times3\times64$		$147 \times 147 \times 64$
Separable convolution	$3\times3\times128/1$	2	$147 \times 147 \times 128$
Max pool	3×3/2	1	$74 \times 74 \times 128$
Separable convolution	$3\times3\times256/1$	2	$74 \times 74 \times 256$
Max pool	3×3/2	1	$37 \times 37 \times 256$
Separable convolution	$3\times3\times728/1$	2	$37 \times 37 \times 728$
Max pool	3×3/2	1	$19 \times 19 \times 728$
Separable convolution	$3\times3\times728/1$	24	$19 \times 19 \times 728$
Separable convolution	$3\times3\times728/1$	1	$19 \times 19 \times 728$
	$3 \times 3 \times 1024/1$		$19{\times}19{\times}1024$
Max pool	3×3/2	1	$10{\times}10{\times}1024$
Separable convolution	$3 \times 3 \times 1536/1$	1	$10 \times 10 \times 1536$
	$3 \times 3 \times 2048/1$		$10\times10\times2048$
Average pool	-	-	$1\times1\times2048$
Optional FC	-	-	$1\times1\times1000$
Logistic regression	-	-	$1\times1\times1000$

- (a) In ResNet-18, the convolution layers 2 to 5 contain 2 successive convolutions instead of 3 and each is repeated only 2 times
- (b) InceptionV3 incorporates factorized 7×7 convolutions and takes in different sized inputs. InceptionResNetV2 incorporates residual networks with factorized convolutions, as well as several reduction blocks
- (c) In VGG-19, the 3rd, 4th, and 5th convolution layer has all 3×3 convolutions and is repeated 4 times each instead of 3 successive convolutions
- (e) DarkNet-19 has no residual networks, with only max pooling layers reducing the image

average as the dataset is imbalanced. This gives same weight to each class.

Analysis of Result

We present a comparative analysis in Table 7 among 5 top performing models in four categories, (a) models that employ a general cross-entropy cost function, (b) models that employ a weighted cost function, (c) models that are trained on the upsampled dataset, and (d) models that are trained on the downsampled dataset. The models are evaluated using different evaluation metrics on the test dataset.

From Table 7, we can see that the Densenet201-Quadratic SVM model outperforms other model in terms of performance metrics. By adopting the weighted cost function as



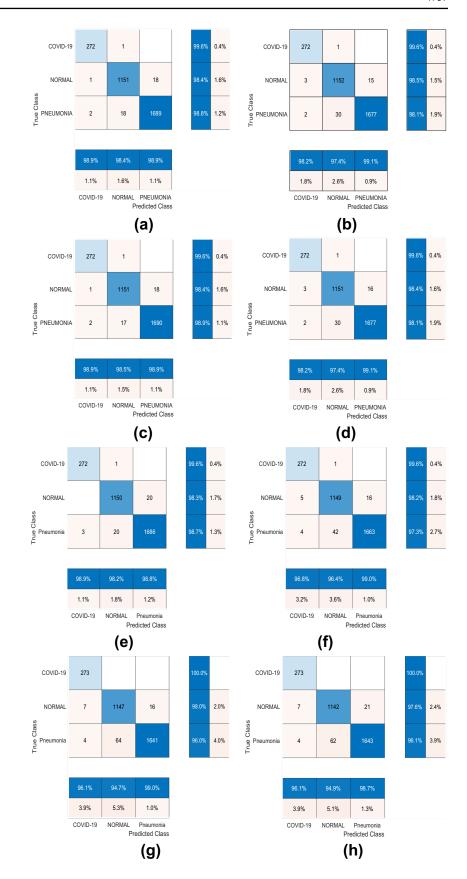
 Table 7
 Evaluation of top-5 models in terms of performance metrics

	Cost fur	Cost function unchanged	hanged			Weighted	ighted cost function	ıction			Up-sampling	ling				Downsampling	npling			
Architecture Dense- Dense- Res- net201 net201 net50	Dense- net201	Dense- Dense- Res- Dark- Dense- Dense- Dense- Dense- Dense- Dense- Dense- Res- Dark- Dense- Bes- Dense- Bes- Dense- Bes- Dark- Dense- Dense- Bes- Dark- Dense- Dense- Bes- Dark- Dense- Dense- Bes- Dark- Dense- Dense- Dense- Bes- Dark- Dense- Bes- Dense- Dense- Bes- Dark- Dense- Bes- Dark- Dense- Bes- Dense- Bes- Dense- Bes- Dark- Dense- Bes-	Res- net50	Dark- net53	Dark- Dense- Dense- Resnet53 net201 net201 net501 net501	Dense- net201	Dense- net201	Res- net50	Dark- net53	Dense- net201	Dense- Dense- Res- net201 net201 net501 net50	Dense- net201	Res- net50	Dark- Dense- net53 net201	Dense- net201	Dense- Dense- Resnet 201 net 50	Dense- net201	Res- net50		Dense- net201
Classifier	QSVM ESD	ESD		OSVM QSVM LSVM QSVM	LSVM	OSVM	ESD	1	QSVM	LSVM	QSVM LSVM QSVM ESD		QSVM	OSVM	LSVM		ESD	OSVM	1	LSVM
Validation accuracy	98.18	98.18 97.91		97.60 97.75 97.52 98.	97.52	91	97.82	97.93	97.61	97.54	99.39	98.47	99.36	99.23	98.56	97.31	97.39	97.61	97.97	96.50
Test accuracy	98.73	98.45 98.41	98.41	98.19 98.22		98.76	98.35	98.41	98.19	98.22	98.60	97.79	98.16	97.53	97.93	97.14	96.92	89.96	96.29	96.44
Weighted precision	98.73	98.45 98.41		98.19	98.23	98.76	98.36	98.41	98.19	98.23	98.60	97.78	98.16	97.55	97.95	97.20	86.96	96.72	96.32	96.56
Weighted recall	98.89	98.58 98.64		98.32	98.31	98.93	98.47	98.62	98.32	98.31	98.76	98.76	98.35	09.76	00.86	97.19	97.02	88.96	96.57	96.58
Weighted F1-score	98.81	98.51	98.53	98.25	98.27	98.84	98.41	98.51	98.25	98.26	98.68	97.81	98.24	97.59	96.76	97.17	76.76	96.77	96.42	96.51
Weighted specificity	98.73	98.44	98.34	98.35	98.23	98.77	98.43	98.42	98.23	98.31	98.60	97.83	98.17	97.58	94.98	97.28	97.03	96.78	96.36	89.96

QSVM quadratic SVM, ESD ensemble subspace discriminant, LSVM linear SVM The Italic entries signify the highest performing architecture for that metric



Fig. 5 a–h Confusion matrix of two top performing models generated using four different settings: general cost function, weighted cost function, upsampling training-set, and downsampling training-set. In the figure, GC, WC, QSVM, and ESD denote general cost function, weighted cost function, Quadratic SVM and Ensemble Subspace
Discriminant respectively





given in (1), we observe slight improvement in the result for the best model. However, the change varies differently across the models. Some models' performance get worse (e.g., Alexnet-CoarseTree, Shufflenet-MediumTree, Alexnet-QuadraticSVM), some obtain better results (e.g., GoogleNet-BaggedTrensemble, Resnet50-CoarseKNN, NasNetMobile-Bagged Tree Ensemble etc.) and others do not change significantly. Note that, we consider at least 0.6% change (minimum 20 samples) in test accuracy as a significant change. We also show confusion matrix of some of the top performing models in Fig. 5.

Here we note few observations that are based on the performance analysis of 640 models:

- As we extracted features from the last pooling layer of each CNN architecture, the number of features is in mid range (between 1000 and 5000). For this dimension of the feature vector, SVM (Linear,Quadratic,cubic) and Ensemble classifiers (Subdiscriminant, Bagged, Boosted) are found to outperform other classifiers.
- Using Densenet201 as CNN architecture gives better result than other CNN architectures.
- 3. Weighted cost function performs better than the two alternative approaches, i.e., upsampling and downsampling.
- 4. With Darknet53-Quadratic SVM model, the accuracy of COVID-19 class is 100%. The model however detects three samples of pneumonia class to be COVID-19.
- 5. The validation accuracy and the test accuracy of the models are almost the same.

Misclassified COVID Samples

We present some of the COVID samples in Fig. 6 that have been misclassified by 78% of the models when trained with the general cost function and the weighted cost function. For COVID-positive cases, generally a chest X-ray shows high opacity in the peripheral part of the image. For patients with serious condition, multiple consolidations present across the image. In contrast, discontinuity of diaphragmatic shadow and subtle consolidation in chest X-ray differentiates pneumonia from other diseases. On the other hand, normal samples do not have any signs of consolidation or high opacity or any discontinuity of the diaphragmatic shadow.

According to an expert radiologist, the X-ray sample shown in Fig. 6a has low opacity in the left and right upper lobes and absence of consolidation even in the peripheral part of the lung, which is similar to normal lung condition. In contrast, the COVID samples shown in Fig. 6b—e are misclassified as pneumonia. These samples, according to the radiologist, are hard to diagnose even by an expert in this domain. More tests like CT scan and RT-PCR are required to confirm the status.

Feature Visualization

As we mentioned earlier in "Method," the CNN models we utilize here as feature extractor are pre-trained on the

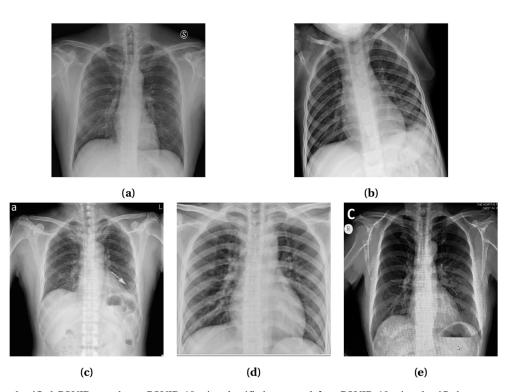


Fig. 6 Some miss-classified COVID samples. a COVID-19 miss-classified as normal; b-e COVID-19 miss-classified as pneumonia



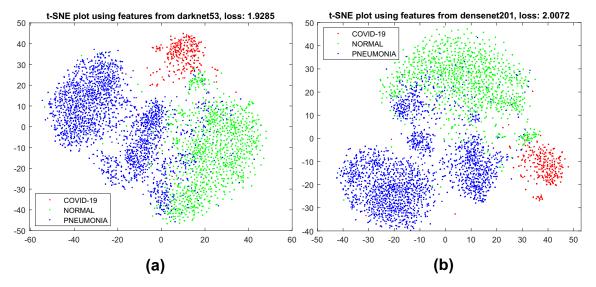


Fig. 7 2D t-SNE [141] visualization of the extracted features obtained by a DarkNet53 and b DenseNet201 from our X-ray image dataset

ImageNet dataset. Even though none of the classes we consider here is included in Imagenet, the CNN architectures produce very good features that are distinctive across the classes.

We visualize the features we obtain via transfer learning in Fig. 7 using 2D t-SNE (t-distributed stochastic neighbor embedding) [141] plot. One can notice that the features of three different classes are quite well separated and easy for a simple machine learning model to do the classification.

Performance with Lower Dimensional Features

Here we re-investigate the performance of the models using lower dimensional features. We apply t-SNE to the originally extracted higher dimensional features and fed the two-dimensional feature generated by t-SNE into the

classifiers we have mentioned in "Method." Table 8 shows the performance measures of top-5 models.

From Table 8, one can see that the Resnet50-Weighted KNN model gives the best result. It is to be noticed that, with the originally obtained higher dimensional features, Denenet201-Quadratic SVM performs the best. Here are our observations:

- 1. Using only 2D features, the best accuracy we obtained is 96.5%. This model gives about 2% more error than the previously obtained one. However, this method is quicker and simpler than the previous one.
- We found KNN classifiers (Weighted, Fine, Medium) and Ensemble Bagged Tree to perform better than other classifiers. We think ese classifiers' ability to learn from lower dimensional space make these perform better than the alternatives.

 Table 8
 Performance analysis of the models using t-SNE output as feature (top-5 models)

Architecture Classifier	Res-net50 WKNN	Res-net50 EBT	Dense-net201 WKNN	Dark-net53 WKNN	Dense-net201 EBT
Validation accuracy	95.37	95.20	95.43	94.39	95.37
Test accuracy	96.51	96.26	95.94	95.88	95.84
COVID-19 accuracy	97.80	96.70	98.53	99.63	98.53
Weighted precision	96.52	96.29	95.97	95.91	95.89
Weighted sensitivity	96.88	96.69	96.28	96.37	96.16
Weighted F1-score	96.70	96.47	96.10	96.12	95.99
Weighted specificity	96.56	96.34	96.23	95.97	95.93

WKNN weighted KNN, EBT ensemble bagged tree

The Italic entries signify the highest performing architecture for that metric



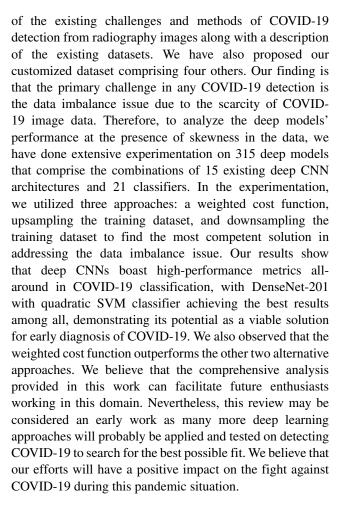
Discussion and Future Directions

One of the major issues in the deep learning-based automated diagnosis of COVID-19 from the radiography images is the requirement of providing a large annotated dataset, prepared by an expert physician or radiologist to train the deep model. Although a major number of recent works made their dataset of radiography images publicly available, they contain at most a few hundreds of images in the COVID-19 class. Compiling a dataset with sufficient images in the COVID-19 class requires collecting radiography images of confirmed COVID-19 patients from reliable and authentic sources which is a challenging task. The challenge is further intensified due to the requirement of the proper annotation of the collected data. To deal with the aforementioned challenges, Few-shot, Zero-shot, and Deep Reinforcement Learning (DRL) can be adopted in this domain for ensuring minimal human intervention in the diagnosis.

Few-shot learning enables a deep model to learn information from only a handful of labeled examples per category. Recently, Few-shot learning has been applied on chest X-ray images to classify some lung diseases where it performed promisingly well [142]. However, Few-shot learning has great potential to detect COVID-19 as well, which has not been explored yet. On the other hand, Zero-shot learning has the amazing capacity of recognizing a test sample which it has not seen during training. Therefore, Zero-shot learning can also be a resort to minimizing the issue regarding the scarcity of training data of COVID-19 class. In addition, DRL can also alleviate the need for good-quality images and the hassle of proper annotations. If the problem can be modeled as a Markov's Decision Process, DRL algorithms can perform remarkably well. As far as we know, the application of DRL in the detection of COVID-19 is not done yet. However, the success of DRL in varied fields, e.g., image processing tasks [143, 144]; image captioning [145]; video captioning [146]; automated diagnosis [147, 148], shows the potential of DRL to be effectively utilized in the diagnosis of COVID-19.

Conclusion

Due to scientists' and researchers' constant conscientious effort to unbolt new methods for the effectuation of COVID-19 detection with the power of deep learning, the research in this domain has made significant progress in a short time. However, due to the absence of a benchmark dataset, researchers have reported performances based on different datasets and evaluation protocols, rendering the idea of comparative analysis far-fetched. This work is inclined towards addressing this issue. Here we discussed some



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Compliance with Ethical Standards

Conflict of Interest The authors declare that they have no conflict of interest.

Ethical Approval This article does not contain any studies with human participants or animals performed by any of the authors.

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