



Medical Image Analysis Using Transfer Learning

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ABSTRACT

Medical image analysis using transfer learning being the foundation of this project, showcases a machine learning technique that saves time and resources in classifying lung diseases. It entails refining an already trained model to classify and extract features from images. Initially trained on a diverse dataset. Instead of starting from scratch for each new image classification task, practitioners can fine-tune the pre-trained model for specific applications. We yielded an initial accuracy of 81% through Inception V3 and ResNet models after which with ongoing efforts dedicated; we received better results which reached a remarkable accuracy of 96% on DenseNet121 model. The abstract encapsulates the essence of transfer learning as a dynamic tool amplifying model accuracy and efficiency, exemplified in our pursuit of excellence within the intricate landscape of lung disease detection. This study explores the transformative capability of transfer learning, shaping the future in image analysis for a context to Pre-trained CNN models for specific applications.

KEYWORDS: Accuracy, CNN models, Datasets, DenseNet121, Feature extraction, Features, Medical Image Analysis, Image Classification, Image Net, Inception V3, Pre-trained model, ResNet, Transfer learning

I. INTRODUCTION

In today's technologically fast-paced world the concept of CNN Models and Transfer Learning plays an important role in the sector of medical diagnosis and imaging, highlighting their advantages in accuracy and resource efficiency. It addresses challenges like data diversity and model interpretability, while suggesting future research directions. Ultimately, it underscores the scope in the concept of CNN models and transfer learning to revolutionize medical diagnosis, emphasizing the need for ongoing development to overcome current limitations. The project's primary focus revolves around addressing the scarcity of extensive medical image datasets suitable for training deep neural networks effectively. It aims to develop training methodologies utilizing a diverse aggregated medical image dataset, with the goal of extracting broadly applicable medical features that can enhance performance in analyzing lung diseases. In a broader context, medical image analysis encompasses important processes like object detection, classification of images, segmentation, and description, all which play integral roles in various applications like medical diagnosis, self-driving cars, and security systems. The increasing demand for accurate and efficient image analysis, especially in real-time scenarios like disease detection, symptom understanding underscores the importance of this endeavor.

Existing systems often struggle with accuracy and efficiency due to limited training data and reliance on manually crafted features. However, transfer learning offers a promising solution by enabling the development of precise and efficient systems, even with limited labelled data, while also improving generalization to new tasks and data. In the domain of image analysis, the concept of transfer learning is increasingly recognized as essential for diverse practical applications. A variety of widely used pre-trained models, including Xception, VGGNet, DenseNet121 and InceptionV3 are available for transfer learning in image analysis. Moreover, there are popular open-source frameworks such as TensorFlow Object Detection API, PyTorch Lightning, and Detectron2, which provide robust tools for performing image analysis tasks.

II. LITERATURE SURVEY

In the pursuit of advancing transfer learning processes, several impactful articles contribute unique methodologies. In [1] the topic delves into the concept of transfer learning (TL) with CNN for medical image identification. Examining 425 articles, it recommends deep models like Inception, emphasizing their efficiency in overcoming data scarcity. The [2] titled as "TL-med: A Two-stage transfer learning recognition model for medical images of COVID-19." Addresses the minimal number of labelled images by ideating transfer learning model (TL-Med) of two stages. It uses main features from diverse data which refines them with large-scale medical data, addressing the challenge of insufficient labelled COVID-19 data. The results on dataset demonstrate a recognition accuracy of 93.24%, showing the model's accuracy effectiveness in detecting disease images. In [3] the authors have studied the concept of CNN and process of transfer learning in medical diagnosis, highlighting their advantages in accuracy and resource efficiency. It addresses challenges like data diversity and model interpretability, while suggesting future research directions. Ultimately, it showcases the potential of CNN and the concept of transfer learning to revolutionize diagnosis practices, emphasizing the need for ongoing development to overcome current limitations.

In [4] the authors have addressed the global impact of COVID which focuses on its challenges in detection. It proposes a deep learning method, specifically using (CNN) with models like VGG16. Results indicate VGG16's superior performance (98.00% accuracy), showing the use as a valuable tool for effective COVID-19 detection, aiding healthcare professionals in decision-making for optimal therapy with minimal resources. In 2016, Liu and authors [5] proposed the SSD algorithm for fast object detection. Both SSD and YOLO are highly efficient, but SSD uses multi-scale feature maps which aid in detecting objects at different scales. As deep networks reduce spatial resolution, detecting small objects at low resolutions becomes challenging, affecting accuracy. YOLO lacks multi-scale features and relies on smoothing lower-resolution maps. Recent studies applied deep CNNs for COVID-19 pneumonia detection from X-rays, achieving high accuracy, precision, and sensitivity. Some used transfer learning with ResNet50 on small 339 instance datasets, obtaining 96.2% accuracy models. The disease of COVID was detected using the concept CNN in a chest by the authors of [6]. They have investigated the use of different approaches for detecting COVID-19 from medical imaging data like chest X-rays, with most achieving an accuracy range of 90-94%. The approach involves fine-tuning the top layers of the pre-trained models while transferring the learned low-level features to the new customized models. One such approach by Xiao [7] on residual networks and multiple instances learning for binary classification of COVID-19 from CT scans. Another study by Panwaretal. proposed a binary classification model using a fine-tuned VGG model to detect COVID-19 from input images. Additionally, they employed Grad-CAM visualization techniques to provide color-coded explanations, making the deep learning model

more interpretable. The key idea is to utilize the powerful feature extraction capabilities of pre-trained models like ResNet50 and VGG, while customizing and fine-tuning them for the specific task of COVID-19 detection from CT images, aiming to develop accurate and explainable diagnostic models.

III.METHODOLOGY

Our research methodology for transfer learning involved training and implementing a CNN architecture, resulting in an accuracy of 96% using the DenseNet-121 model. Activation functions utilized in this approach were ReLU and Softmax.[8].

3.1 DATASET INFORMATION

The images of CT scans of human lungs are included in the dataset. CT scan is a type of X-ray that is used to diagnose the sensitive inner organs of human body precisely. A medical images collection from various sources comprises this dataset; a sum of 17,704 images, out of which 12,351 images are directed toward training and 5,353 images for the purpose of testing the model. The distribution of images across classes reveals that the dataset is relatively balanced, with each class having a substantial number of images. The classes include Normal, Mass, Pneumonia, and COVID, with Normal having the highest number of images at 4,685, followed closely by Mass at 4,528, Pneumonia at 4,273, and COVID at 4,219. The training set distribution is also balanced, with Normal having 3,280 images, mass having 3,170 images, Pneumonia having 2,992 images, and COVID having 2,909 images. The testing set distribution follows a similar pattern, with Normal, mass, Pneumonia, and COVID having 1,405, 1,358, 1,281, and 1,310 images, respectively. This dataset provides a comprehensive and \balanced collection of images for training and testing purposes in the context of diagnosing respiratory conditions using medical imaging. The dataset's reliability is ensured through acquisition from reputable organizations.

3.2 CNN ARCHITECTURE

Our model begins with an input layer, capturing raw pixel values of medical images. Convolutional layers extract features hierarchically, identifying patterns like edges and shapes. We use a pre-trained convolutional base, facilitating feature extraction from medical images. Pooling layers aid in retaining relevant spatial features, with Max pooling enhancing focus on critical details. Batch normalization stabilizes learning, particularly beneficial for diverse medical data. Dropout layers prevent overfitting by excluding neurons during training. The fully connected layer translates features into class probabilities, adapting specifically to medical image nuances. Fig. 1 represents the 3D CNN architecture of the model.

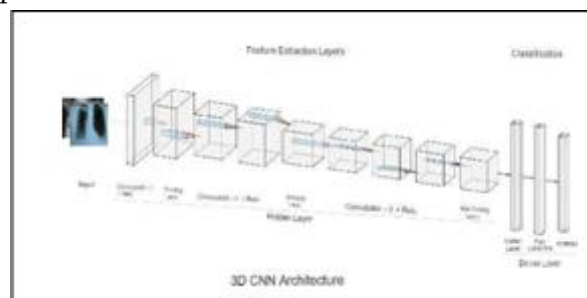


Fig. 13D CNN Architecture

3.3 TRANSFER LEARNING METHODOLOGY

The usage of pre-trained CNN models such as VGG, ResNet50, Xception, InceptionV3, MobileNetV2 and DenseNet, we evaluate their effectiveness on our dataset. These base models, sourced from TensorFlow, have their classification layers replaced to match our dataset's four output classes. While the pre-trained weights are retained, new classification layer weights are initialized randomly. Unlike freezing the convolutional layers, we fine-tune all layers on our dataset, allowing optimization for our specific task. To address the challenge of training neural networks with limited data, transfer learning was applied using a tiny training dataset to generate a feature set for managing lung diseases and Cancer cases. The dataset, being considerably smaller, takes greater time to build a perfect model. Fig. 2 given below represents the flowchart of how transfer learning process takes place. Therefore, the models described were sourced from pre-trained models to help identify lung diseases.[9].

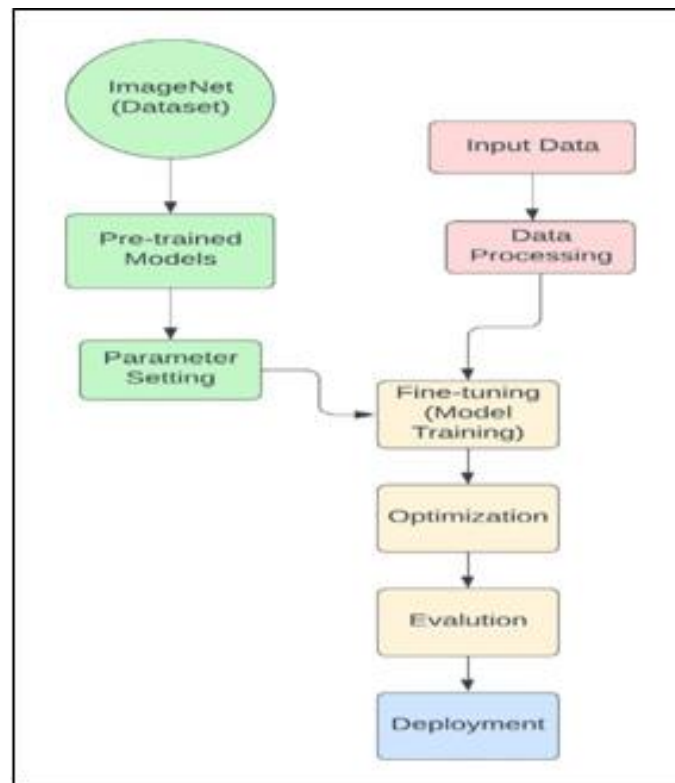


Fig. 2 Flowchart of Transfer Learning Process

3.4 MODELS USED

i. DenseNet121

DenseNet121, introduced by Huang [18], enhances the ResNet50 architecture by understanding connections, ensuring that every layer has connections with all the other layers present. This computationally built network allows for the usage of features while allocating and keeping the parameters as less as possible.

ii. InceptionV3

InceptionV3 addresses the issue of position variability in images by incorporating multiple types of kernels on the same pedestal, which results in increasing the network. The Inception contents allow for the simultaneous operation of many kernels. InceptionV3 is an extension of InceptionV2, addressing representational bottleneck concerns.

flattening the output from the earlier layer to a one-dimensional array and adding a Dropout layer to prevent overfitting.

The final output consisted of a dense layer with a Softmax function for predicting class probabilities. The Adam optimizer [20] and cross-entropy for a loss function were used for training. Overall, these techniques and models were crucial in developing an efficient and accurate model for detecting lung diseases. Models undergo 10 epochs of training with a batch size of 32 images, minimizing categorical cross-entropy loss. If validation fails to improve by at least 0.001 for three consecutive epochs, preventing overfitting and ensuring robust performance on unseen data.

3.7 HYPERPARAMETERS

A learning rate of 0.001 safeguards previously learned features during fine-tuning, with the Adam optimizer chosen for adaptability and efficient convergence. A batch size of 32 balances efficiency and memory constraints. 10 epochs suffice for fine-tuning, with dropout regularization introduced to prevent overfitting. ReLU and Softmax activations are applied strategically throughout the network.

3.8 MATHEMATICAL EQUATION

In a completely connected layer, every neuron is connected to other neurons in the previous layer. The layer of output Y of a fully connected layer can be calculated using the value of the input vector X , the weight W , and finally the bias vector denoted by b . The activation function identifies the absence of linearity into the connection. One of the majorly used activation function in this study is ReLU (Rectified Linear Unit), [15] which shows the output as it is given if it is positive, if not then shows zero, effectively introducing absence of linearity concept to the model. Another majorly used activation function in this project is Softmax which is important for multi-class classification problems. Softmax transforms the raw scores into the probabilities, making it suitable for classification tasks. Eq. 1 given below represents the formula for the ReLU that is used in the transfer learning process. Along with it the Eq. 2 represents the formula for Softmax used in the project.

$$Y = \text{ReLU}(W \cdot X + b) \quad (1)$$

$$\text{Softmax}(x_i) = \frac{e^{x_i}}{\sum_{j=1}^N e^{x_j}} \quad (2)$$

IV. RESULTS AND DISCUSSION

The study calculates the results using the terminologies of Precision, Recall and Accuracy. [19] These parameters are important for studying a medical system for lung diseases identification.

I) Precision Value

Precision is a concept that studies how strong the model recognizes the right samples and is near to the expected output. The higher the value of precision shows, the higher is the positive sample. The below Eq. 3 identifies the precision for deriving results.

$$\text{Precision} = \frac{TP}{TP+FP} \quad (3)$$

II) Recall Value

The increased and better accuracy[16] in first identifying the target instance and the value of likelihood both result into a greater recall value. The formula to calculate recall is shown in Eq. 4.

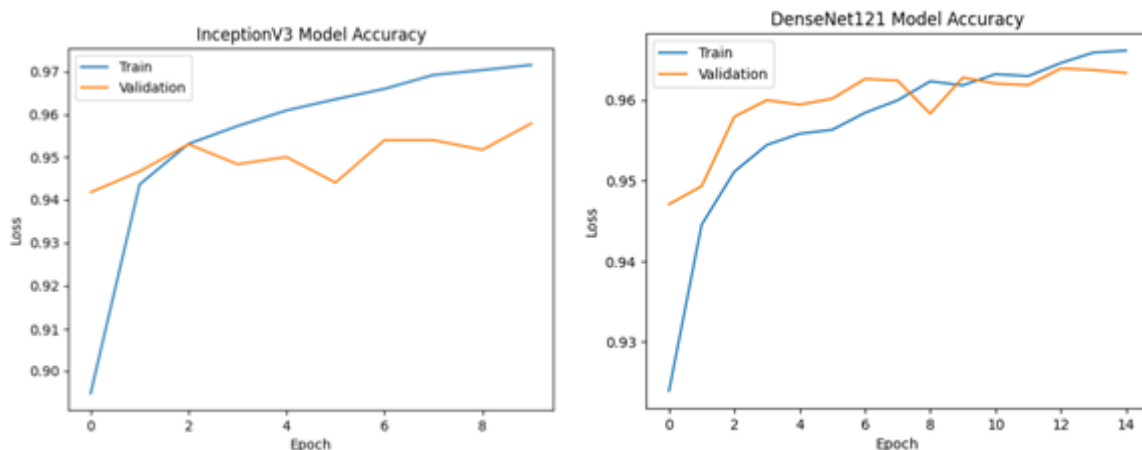
$$\text{Recall} = \frac{TP}{TP+FN} \quad (4)$$

III) Accuracy Value

Accuracy can be stated as the ratio of correct predictions that are made to the value of total number of samples used under study. Eq. 5 is depicting the formula used to find the accuracy of models in the project.

$$\text{Accuracy} = \frac{TP+TN}{TP+TN+FP+FN} \quad (5)$$

In the evaluation, DenseNet121 showed the best performance among the tested models. Following closely were VGG19 and ResNet50.[17] Fig. 4 and Fig. 5 given below depict the graphs of Accuracy and Loss Value of DenseNet121 Model respectively. These results highlight the impact of using transfer learning in process of detecting lung diseases in CT scans. By using transformation-based bespoke models, it is possible to further improve the performance of deep CNN models to achieve accuracy levels exceeding 90% across all performance criteria. The low False Positive Rate (FPR) of the proposed method makes it suitable for real-world screening settings.



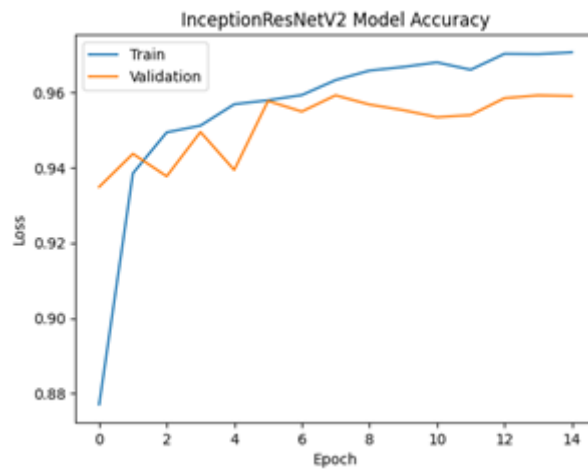
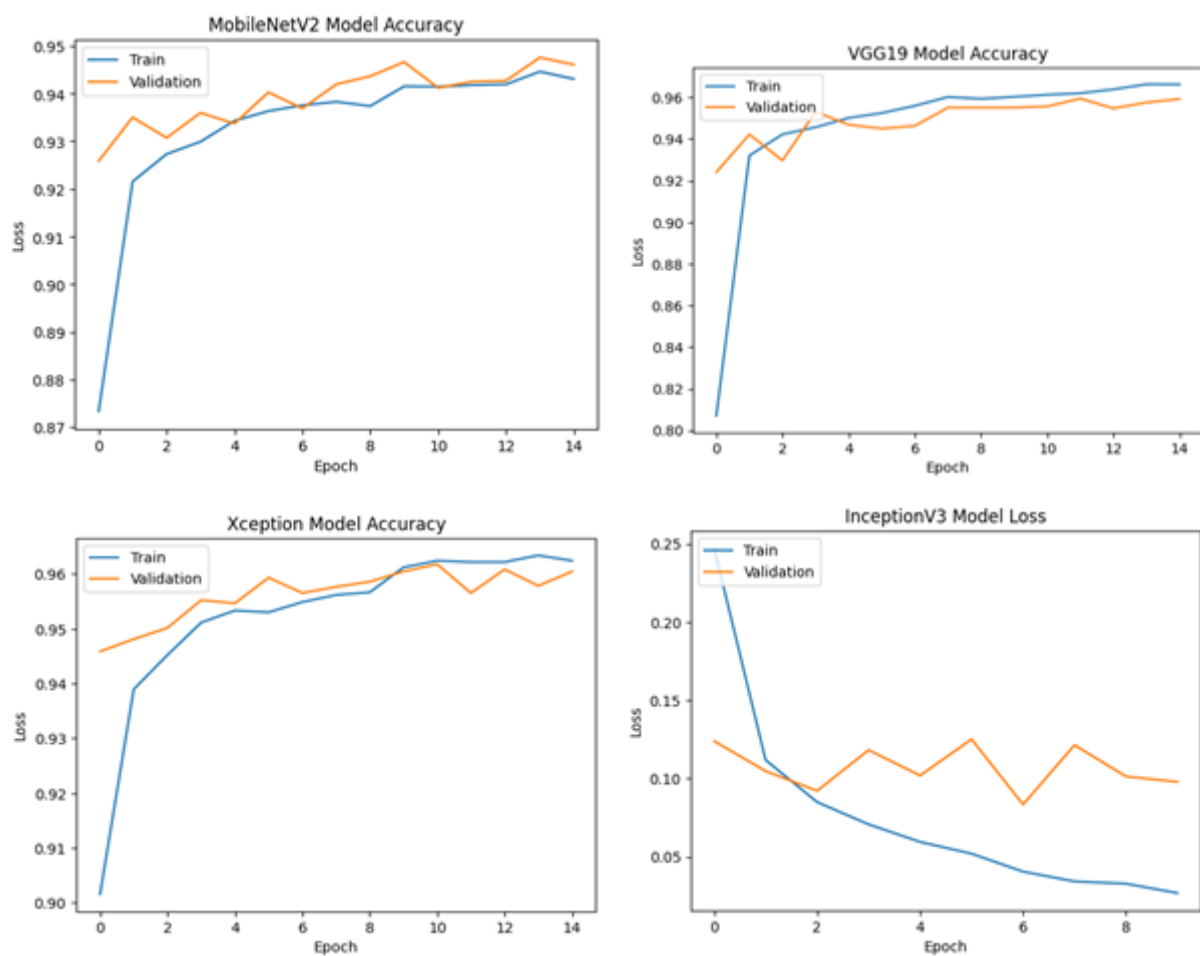


Figure 4: Graph of Results of Accuracy of CNN Models



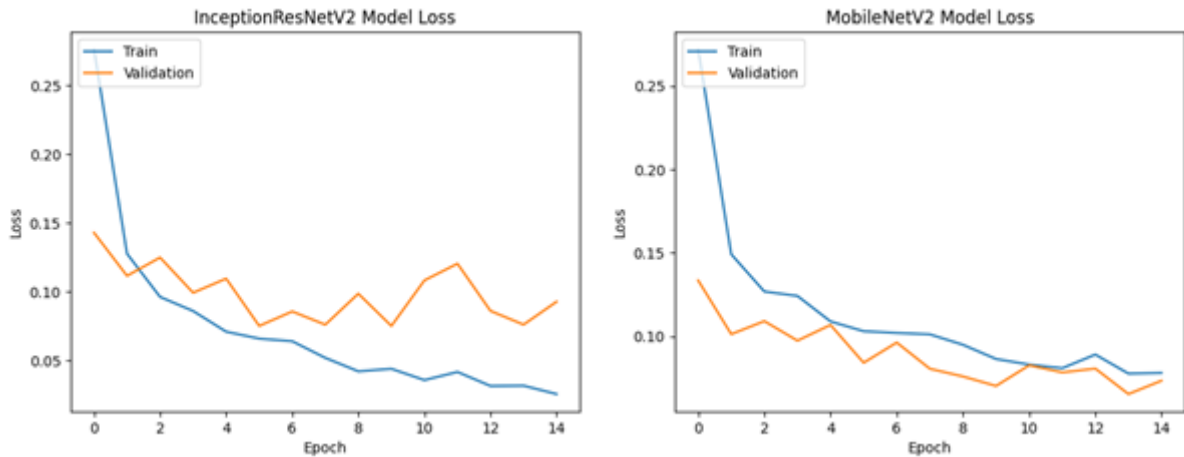
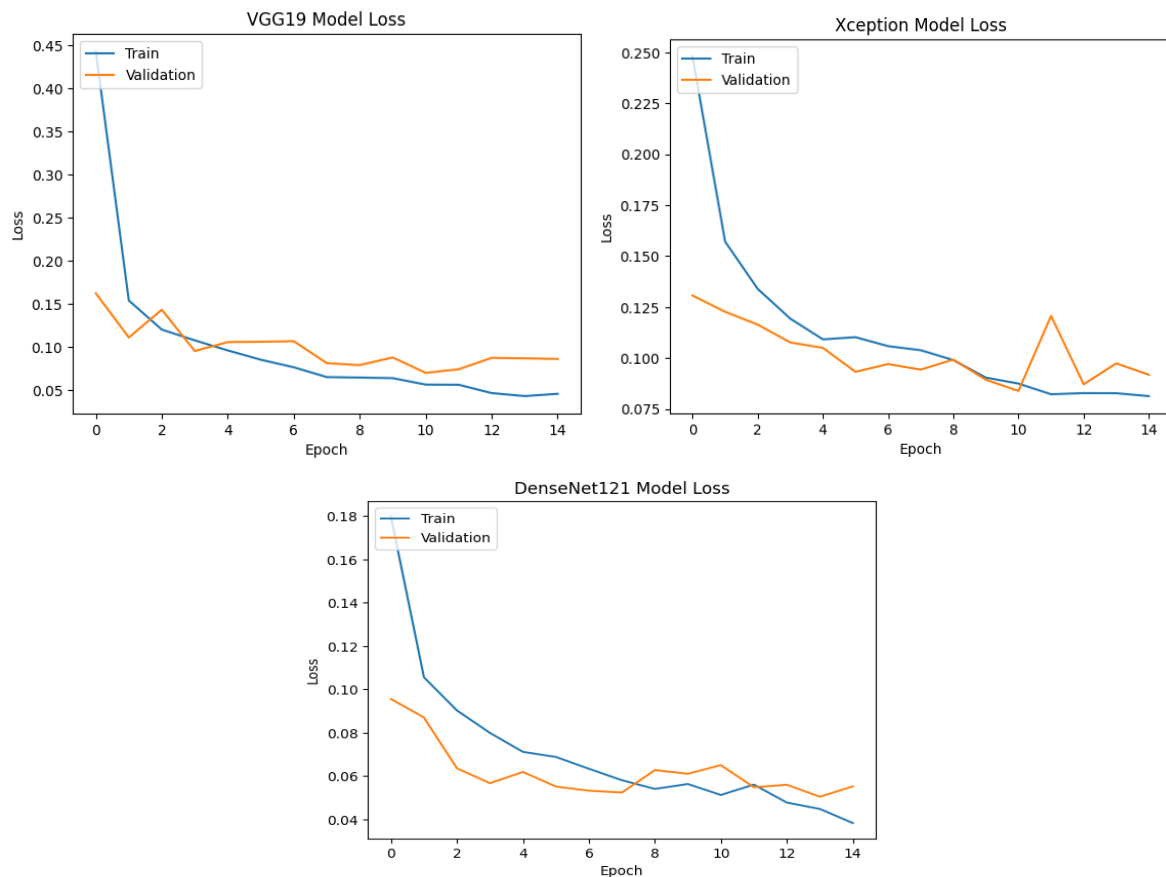


Figure 5: Graphs of Results of Loss Value Models



V. CONCLUSION

This paper serves as a valuable starting point for individuals in the deep learning field who are utilizing transfer learning. It offers guidance on selecting optimal methodologies to improve model accuracy, as evidenced by achieving 96% accuracy using the DenseNet-121 model. Additionally, the study conducts an analysis of different CNN architectures employed in classifying lung diseases from medical images. It showcases how advancements in deep learning algorithms yield promising results, augmenting radiologists' capabilities. The

models based on CNN have found an impactful use in the domain of healthcare and medical diagnosis. The study shows usage of multiple CNN image processing and classification models to detect lung diseases using images of CT scan of patients using transfer learning. The findings indicate that the method used achieves exceptional findings, with accuracy percentage exceeding 90% and low false positive rate. The results suggest that CNN-based methods can impact the control of lung disease spread by providing rapid screening. Given the widespread use of DL-based approaches in other medical imaging applications, their implementation in lung disease screening is timely. The analysis highlights DenseNet121 as a standout performer due to its smaller parameter size and minimal training time, surpassing other CNN models. Finally, the study shows the results of transfer learning-based transformations to detect lung diseases in suspected patients using CT scan images.

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