

# Lungs Tumor Detection Using Convolutional Neural Network on Histopathological Images

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## Abstract:

Diseases that affect the lungs have the effect of making breathing difficult. One of the biggest causes of death among people is lung cancer.

Human survival rates can be increased through early detection. In the event of a positive diagnosis in lung cancer patients has increased from 14% to 49% throughout that period. When computed tomography (CT) is more accurate than X-ray images, complete diagnosis requires the use of many imaging modalities that complement one another. In this study, we create and test a deep neural network to identify lung cancer in CT scans. An adaptive boosting algorithm and a densely connected convolutional neural network (DenseNet) were used to determine whether a given lung image was healthy or cancerous. For this task, we use a dataset containing 201 images of the lungs, with 85% of the images are used for training and 15% used for testing and classifying. According to experimental data, the suggested approach has a 90.85% success rate.

**Keywords--** Convolution Neural Networks (CNN), Image Processing, Deep Learning, Machine Learning, Artificial Intelligence

## I. Introduction

One of the leading causes of death globally is lung cancer [1]. It's one of the worst kind of cancer to strike people. It has the highest mortality rate of any tumor and is the leading cause of cancer-related death of men and women [2, 3]. Nearly 2.1 million new cases of lung cancer are diagnosed each year (14.3 percent of all cancers), with 1.8 million deaths occurring as a result (20.4 percent among all cancers). When abnormal cells in the lungs grow and divide uncontrollably, a tumor forms. Lung cancer has the highest mortality rate of all cancers. Around 85% of male and 77% of female lung cancer cases are caused by smoking cigarettes. With a mortality rate of 20.4 percent, lung cancer ranks among the dangerous diseases plaguing developing nations. A growing number of people lost their lives every year to lung cancer, making it one of the deadliest forms of the disease [4-6]. The Pros of Early forecasts using fuzzy logic will lead to outcome-focused investigation [5]. Lung cancer survival rates after diagnosis are proportional to the disease's stage. Nonetheless, people's chances of success are highest in their younger years. Lymph fluid, which bathes the lung tissues, acts as a conduit for the dissemination of cancer cells into the bloodstream. The lymph travels through lymph vessels before draining to the lungs and chest lymph nodes. One of the biggest challenges facing humans today is examining and treating lung disease. The survival rate of people with tumors can be

significantly increased through consistent early detection efforts. This study presents a technique for classifying lung cancers as malignant or benign using a convolutional neural network (CNN). CNN's success rate is 81%, making it the neural network with the best results. The lung is a common site for malignant tumors to form. The result is lung tissue damage. Lung cancer is characterized by the uncontrolled growth of abnormal cells in the lungs. Both lungs are equally at risk. Lung cancer symptoms include coughing up blood and having trouble breathing. Smokers are disproportionately affected by this problem. Cigarette smoke is a major contributor to lung cancer. Cigarette smoking is a leading cause of death in the United States [14, 15, 16]. The World Health Organization estimated in May 2020 that tobacco use causes the deaths of more than 8 million people each year. Tobacco consumption is responsible for the deaths of about seven million people worldwide, while secondhand smoke contributes to the deaths of an additional 1.2 million [17].

More than four thousand chemicals have been identified in cigarette smoke, according to [18]. Most of these chemicals have been linked to cancer. People who smoke more than one pack of cigarettes per day have a 20-25 times higher risk of developing lung cancer than nonsmokers [19], but early detection increases the chance of survival by 73%. Around 90% of lung cancer fatalities are attributable to tobacco use or smoking [1]. Although cigarette smoking is still the leading cause of lung cancer, there are other risk factors to consider. Lung cancer risk factors include environmental pollution (primarily air pollution), excessive alcohol consumption, and exposure to tobacco smoke (passive smoking).

Many people would benefit from early detection or prediction of lung cancer. This might be an efficient preventative measure. Lung cancer diagnosis often involves a battery of tests, including computed tomography, chest radiography, sputum cytology, magnetic resonance imaging, etc.

Late or delayed cancer identification, however, dramatically reduces a patient's likelihood of survival, and most of these procedures are expensive and time-consuming. [21] A technological solution for early diagnosis is required to address this challenge, since it will lead to more rapid outcomes.

The system uses deep learning methods for cancer detection. To distinguish between cancerous and noncancerous tissue, the system employs the Convolutional Neural Network algorithm on whole slide images [3]. [23] The algorithm in this system for early detection is responsible for extracting features. The machine will make a lung cancer diagnosis based on these characteristics. The system was developed to supply a quick and cheap answer for lung cancer early detection. Certain criteria and thresholds, to be detailed later [22], will be used to make the detection. Medical image processing [12, 13] is a popular application of Deep Learning.

## II. Existing System

Histopathological classification of lung cancer now frequently makes use of medical imaging [5]. In this study, we provide a novel CNN-based architecture for classification of lung cancer images using histopathology. One input layer, various blocks of convolutional layers, a drop-out layer, a max-pooling layer, and a fully connected layer are all part of the CNN architecture presented in this study. Sixteen convolutional layers, five dropout layers, five max-pooling layers, and one fully connected layer make up the total network architecture. The proposed architecture thus validates the superior performance when compared to baseline models.

MRI images are used as a dataset in [6]. Nevertheless, the dataset is quite tiny, hence transfer learning is applied. Instead of starting with a blank slate, models built using transfer learning make advantage of pre-trained, deep convolutional Neural Network architectures.

AlexNet, GoogleNet, VGG-19, VGG16, ResNet-18, ResNet-50, ResNet-101, ResNet-inception-v2, and SENet are just some of

architectures used on the dataset in the paper. Epochs of 25, 50, and 90 are used to generate the results. The findings are then compared with one another. Finally, the results from the three selected epochs demonstrate that VGG16, VGG19, and some AlexNet layers outperform ResNet and ResNet-Inception-v2. Their respective rates of accuracy were 98.71, 98.55, and 98.55 percent. After 50 iterations, AlexNet and VGG16 both reached 98.55% precision. Nonetheless, VGG16 was slower than AlexNet and took more time to train.

In [1], the author explains how neural networks have changed over time. Using examples of classification, detection, and segmentation on pulmonary medical images, this paper provides an introduction to the application of deep learning to medical image analysis. The steps involved in using deep learning to identify the pulmonary nodule are also detailed. Lung conditions such as pulmonary nodule disease, pulmonary embolism, pneumonia, and interstitial lung disease are all treated using these strategies.

In order to categorize pneumonia, a chest X-ray dataset is used in [2]. They used three different methods on this data set, the first of which was a linear support vector machine model with local rotation and orientation free features. Second, we use two different convolutional neural network models to perform transfer learning. The capsule network is the third method. Since the dataset used was insufficient, various forms of augmentation are applied to the same data in an effort to boost accuracy. Results from all three experiments pointed to CNN-based transfer learning as the most effective method. Superior efficiency is achieved by capsule networks compared to ORB and SVM classifiers.

### III. Proposed Model

The architecture of convolutional neural networks is optimized to perform image classification with a small number of parameters. A convolutional neural network is built up from several interconnected layers, each of which performs a specific

architecture of a ConvNet is a neuronal connectivity pattern that is reminiscent of the Visual Cortex. When data is "augmented," its volume and complexity grow. Instead of converting the existing data, we will collect new data. Since we need large amounts of data for deep learning, and in some cases it's not really feasible to gather millions of images, data augmentation is a crucial step. These data optimize the dataset magnitude and add ambiguity inside the patients, here are the references.

### IV. Preliminaries

Several machine learning and deep learning methods on diverse datasets may be used to diagnose lung cancer. Various studies on lung cancer detection have already been done by researchers on different datasets. Brief explanations of the studies reported in publications about methods for detecting lung cancer are provided below. The UCI machine learning repository and data world have yielded varying outcomes for the various classifiers used to detect lung cancer. Classifiers such as DT, LR, NB, and SVM were used to put the findings into practice. It has been shown that SVM outperformed other classifiers on the dataset [3]. In this research, the author conducted experiments using the Lung Imaging Database Consortium image collection (LIDC-IDRI) using a deep neural network and auto encoder to identify lung cancer. Accuracy levels of 79%, 81%, and 79% were attained by them, respectively [4]. They used Random Forest, an ensemble of U-Net, XGBoost, and ResNet, to obtain an accuracy of 84% on the Lung Imaging Database Consortium image collection (LIDC-IDRI), proposing this way of creating deep residual networks and classifiers [5]. This research proposes a CNN architecture for identifying lung cancer from whole-section histopathology pictures. They performed a VGG and a ResNet. Receiver Operating Characteristic (ROC) plots were used to evaluate VGG and ResNet's final results. In terms of patch-level accuracy, the VGG16 model scored 75.41% and the ResNet101 model scored 72.05%. The author of this paper [8]

describe how theyported a CNN network over to the LC25000. Using a combination of lung and colon histopathological images, the author strained a CNN model to distinguish between three distinct types of lung cancer. The constructed CNN model has training and validation accuracies of 96.11 and 97.2 percent, respectively.

### V. Operational Training using CNN

The Convolutional Neural Networks (CNNs or ConvNets) utilized for the categorization and recognition of images were built using a linear stack of layers. Convolutional layers with kernel filters, max pooling, and fully linked layers were used on both the training and testing pictures. The item was put through a classification process using the softmax algorithm. The model was developed and tested using hardware with the GPU moniker provided by the Google Colaboratory. For this assignment, we used a neural network with three hidden layers, one input layer, and one fully connected layer. To facilitate both training and evaluation, images are divided 90:10. The input layer received images of size (180, 180) pixels. Each convolutional layer used a kernel matrix of size (3, 3) with the activation function ( $\text{ReLU}(x) = \max(0, x)$ ). To lessen the load on the subsequent convolutional layer's computing parameters, a max pooling size of (2, 2) was introduced. The model used a dropout value of 0.1, which is rather little. The class probabilities for the final output classes were calculated using a dense value of three and the sigmoid activation function. Adam, an optimizer based on adaptive moment estimation, was used to determine the rates of parameter learning. To determine whether the model could correctly identify malignant lesions on radiographs, a detection performance test was run on a per-lesion basis using the test dataset. A chest radiograph lesion's likelihood of being

malignant was converted to a number between 0 and 255 by the model. True positives were determined by whether or not the median of the model's output fell within the range of the gold standard (TP). The remaining outcomes were all FPs. When the model suggested many TPs for the same underlying fact, we aggregated them into a single TP. For every missing result for a single ground truth, the model returned one FN. A.S. and D.U., two radiologists, looked at the radiograph and CT after the fact to see what structures were picked up by the FP output. Segmentation accuracy was also measured using the dice coefficient.

There are two primary components to the method suggested for detecting lung cancer. As a preliminary step, we do preprocessing on the photos before feeding them to 3D convolutional neural networks. In order to identify CT images as positive or negative for lung cancer, we first had to discover the nodule candidate that would be utilized to train 3D convolutional neural networks.

#### 1) Preprocessing and Segmentation

For each patient, we first convert the pixel values in each image to Hounsfield

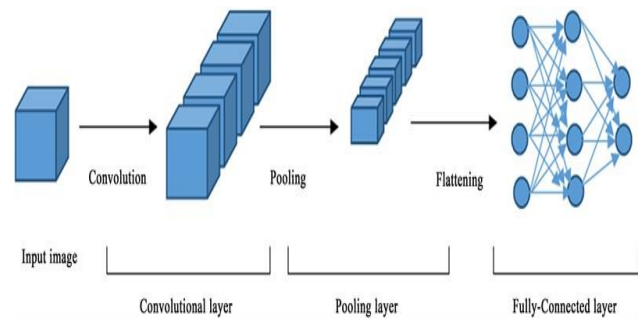


Fig1. Preprocessing of Images in Convolutional Neural Network

### VI. Description of dataset

Kaggle dataset is used in this study. There are a total of 250,000 images in this dataset, with 5,000 images from each of five categories. Three forms of lung cancer and two of colon cancer are present. Histopathological images are the focus of this dataset. All of the JPEG files are 768 pixels on the longest side. Histopathology is the study of disease through microscopic analysis of tissue. In histopathology and clinical medicine, it refers specifically to the examination of a processed and histologically sectioned surgical or biopsy specimen on glass slides by a pathologist. Histopathologists are specialists in cancer diagnosis; they are the ones who examine the tissues and cells taken from suspected "lumps and bumps," determine whether or not they are malignant, and then report back to the treating physician with details about the cancer's type, grade, and response to treatment. The pancreas and retro peritoneum, for example, were previously inaccessible for biopsy tissue retrieval, but modern imaging techniques have made this possible. After the tissue has been collected, it is usually processed overnight before being examined microscopically (Figure 1).

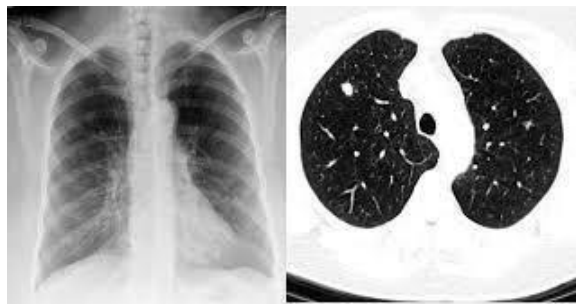


Fig2. Image of Lungs

**Image resizing:** In computer vision, resizing pictures is a crucial part of the preprocessing phase. Our machine learning algorithms, in general, benefit most from training on lower resolution photos. In addition, our raw gathered photos may not

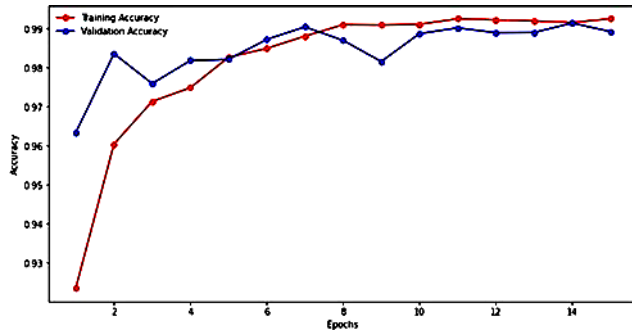
all be the same size, yet this is a requirement of many deep learning model designs. In order for the CNN to function properly, all photos must be scaled to the same dimensions before being fed into it. In the dataset, the original image size is a sizable 768 pixels by 768 pixels. Computer vision is used to resize the photos to 224 by 224 pixels so that the model may be trained more quickly and at a lower computational cost.

**Dataset splitting:** Separating your data into a train, validation, and test set is a must if you want to succeed in machine learning. The rationale for this practice is elementary. The model should be tested using the same data it has already seen if we did not separate the data into various sets. Hence, issues like overfitting may arise throughout training without our awareness. In this case, the Effective Net-B0 model was used. Pre-trained on ImageNet, this model was fine-tuned by swapping out the last layer along with a series of linear and dropout ones. In this case, we use a ReLU activation function in 2 dense layers. Dropout layer with a 0.2 percent rate is used in ReLU ( $f(x) = \max(0, x)$ ). Dropout should be utilized to prevent the model from being overfit. Once a last dense output layer was placed, the process was considered complete.

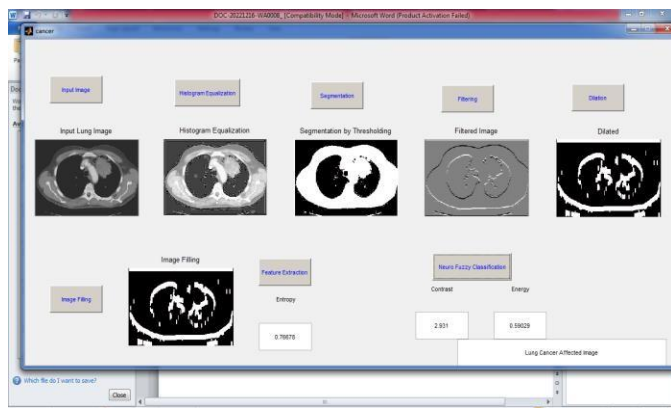
Table 1: Details of Specification

Model	MATLAB R2014a
Used Software	224*224
Image Size	32
Batch Size	1.00E-03
Learning Rate	15
Epochs	5
Additional Layers	0.2
Dropout Rate	Adam
Optimizer	CrossEntropy
Loss Function	Loss





**Figure3:** Model Training and validation accuracy vs. epoch



**Fig3** Overall system GUI

## VII. RESULT AND DISCUSSION

Above, Fig. 2 and Fig. 3 shows the plot of model accuracy vs. epoch and model loss vs. epoch for training and validation images. The images are trained for 20 epochs with batch size 65 with 212 steps in each epoch. The model achieved a training accuracy of 97.11% and a validation accuracy of 98.20% in the final approach.

## VIII. CONCLUSION

Using histopathological scans, this study demonstrates how lung cancer may be diagnosed. One picture was trained using a CNN to determine if it belonged to the benign, Adenocarcinoma, or squamous cell carcinoma categories. The model's training accuracy was 97.11 percent, and its validation accuracy was 98.20 percent. Model performance was evaluated using a

number of metrics,

including precision, f1-score, recall, and a confusion matrix plot.

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