Comparison of deep learning convolutional neural network (CNN) architectures for CT lung cancer classification

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Received: 9-October-2020; Revised: 14-January-2021; Accepted: 18-January-2021
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Abstract
Lung cancer has become one of the most common deaths amongst the cancer patients. World Health Organisation states that lung cancer is the second most fatal cancer all over the world in 2014. Alarming, most of the lung cancer patients are diagnosed at the later stages where the cancer has spread. Thus, early screening via Computed Tomography scan particularly among active smokers is encouraged. Manual diagnosis of the cancer is made feasible through the integration of Computer Aided Diagnosis system. For the past few years, deep learning method leads most of the artificial intelligence applications including CAD systems. This paper aims to investigate the performance of five newly established Convolutional Neural Network architectures; GoogleNet, SqueezeNet, DenseNet, ShuffleNet and MobileNetV2 to classify lung tumours into malignant and benign categories using LIDC-IDRI datasets. Their performances are measured in terms of accuracy, sensitivity, specificity and area under the curve of the receiver operating characteristic curve. Experimental results show that GoogleNet is the best CNN architecture for CT lung tumour classification with an accuracy of 94.53%, specificity 99.06%, sensitivity of 65.67% and AUC 86.84%.

Keywords
Computed tomography, Convolution neural network, Deep learning, Lung cancer.

1. Introduction
Lung cancer is the second most fatal cancer worldwide and ranks at the eighth place in the overall reported deaths in Malaysia [1]. Almost 90% of lung cancer patients are both active and passive smokers. As early symptoms of lung cancer are commonly unnoticeable, most of the diagnosis of these patients are found out when they are at stage 3 or 4. Therefore, early screening is highly anticipated. Lung cancer screening can be carried out through computed tomography (CT) scan, sputum cytology and biopsy [2]. Of these three, the non-invasive CT scan is the most viable method to detect lung lesions. However, it is difficult to manually diagnosis small lesions in CT images with high accuracy especially when the surroundings are unclear and noisy.

To assist the physicians on cancer diagnosis, computer aided detection and diagnostic (CAD) system is used. Among the early applications of CADs are on the diagnosis and management of breasts and thoracic cancers [3]. Of recent, the CAD systems are mostly founded based on machine learning methods led by the deep learning method. The main downside of the conventional machine learning methods is the requirement for feature extraction process. The process is omitted when using deep learning. Although deep learning requires heavy computing power and large datasets for
training, the advancement of computational power, top notch graphic processing unit (GPU) and large open online database are constantly and readily available in order to facilitate deep learning implementation. Therefore, improved deep learning neural networks can be created to establish the best CAD system [4–6] to achieve an accurate cancer diagnosis, for instance in lung lesion diagnosis application via CT images.

Up till now, there are several deep learning architectures have been proposed for lung lesion classification. In lieu of this, this paper compares the performance of five different deep learning architectures; GoogleNet, SqueezeNet, DenseNet, ShuffleNet and MobileNetV2 in the application of lung lesion classification into either malignant or benign specifically in CT images.

2. Literature review

There are several studies in the past literature that use deep learning to classify the lung lesions in CT images. Table 1 shows four recent studies for such application. Has compared three deep learning architectures to classify the lung lesions into benign and malignant [4]. They are convolutional neural network (CNN), deep neural network (DNN) and sparse auto encoder (SAE). Their study shows that CNN outperforms the rest of the architectures with the highest accuracy of 84.15%, specificity of 83.96% and sensitivity of 84.32%.

Proposed a two deep 3D customized mixed link network (CMixNet) architecture for detecting and classifying lung nodule [7]. Faster R-CNN is used to learn features from CMixNet and together with U-Net like encoder decoder architecture, nodule detection is achieved. Classification of the lung lesions takes place by learning the features through gradient boosting machine (BGM) from the designed 3D CMixNet structure. Their proposed method achieves an accuracy of 88.79%, specificity of 89.83% and sensitivity of 81.35%.

On the other hand, [8] proposed a hierarchical semantic convolutional neural network (HSCNN) in predicting the malignancy of CT scan pulmonary nodule. The architecture is based on two level output; one is low-level semantic features and another one is high level prediction of malignancy of nodule. The architecture gives an accuracy of 88.9%, specificity of 83.96% and 81.35%.

Compared three CNN architectures; namely Alexnet, VGG16 and their proposed architecture known as SilNet [9]. AlexNet is a very popular CNN architecture for its simplicity whereby it has 5 convolution layers and 3 fully connected layers. AlexNet has three convolution group which are the kernel size of 11x11, 5x5 and 3x3. VGG16 is almost like AlexNet but with the addition of 3x3 filter that replaces large kernel size filter on the AlexNet. It has better accuracy than AlexNet. SilNet consists of a group of layers that makes up a group of convolutions which are convolution layer, batch normalization layer, ReLu and pooling layer. This group of convolutions is used five times in the architecture.

Table 1 Several studies on lung lesions classification using deep learning

| Ref | Architecture | Accuracy | Specificity | Sensitivity |
|-----|--------------|----------|-------------|-------------|
| [4] | CNN          | 84.15    | 83.96       | 84.32       |
|     | DNN          | 82.37    | 80.66       | 83.90       |
|     | SAE          | 82.37    | 83.96       | 81.35       |
| [7] | CMixNet      | 88.79    | 89.83       | 93.97       |
| [8] | HSCNN        | 84.20    | 88.90       | 70.50       |
| [9] | AlexNet      | 79.50    | 76.50       | 83.04       |
|     | VGG16        | 97.50    | 97.60       | 98.26       |
|     | SilNet       | 98.75    | 99.40       | 99.56       |

In addition, [10] proposed 15 layers CNN based architecture to classify long nodules into nodules and nonnodules types, known as LdcNet. It has both feature extraction layer of 3 convolutional blocks and classifier layer. They achieved a classification accuracy of 97.2%. Also applied their novel Gated-Dilated (GD) network that is founded on CNN for benign and malignant lung lesion classification [11]. It achieved an accuracy of 92.75%. Multiple dilated convolutions are used instead of max-poolings while there is a specific network to capture the input features and to determine the suitable dilated convolution.

In this paper, we aim to compare five CNN based deep learning architectures which are GoogleNet, SqueezeNet, DenseNet, ShuffleNet and MobileNetV2 in the application of lung lesion classification into either malignant or benign specifically in CT images.
SqueezeNet, ShuffleNet, DenseNet and MobileNetV2. GoogleNet [12] also known as Inception-v1 is created by Google. This architecture introduces a module called inception module that contains 1x1 convolution used to shrink the dimension to lessen the computational bottleneck. The module also assists in the expansion of the depth and the width of network.

SqueezeNet that is introduced by [13] has the advantage of having 50 times less parameters as compared to AlexNet. This architecture uses 1x1 filter instead of 3x3 filter which makes it 9 times smaller in parameters. It also has a module called fire module that contains a squeeze convolution layer with 1x1 filter. This layer is inputted to the expand layer with both 1x1 and 3x3 convolution filters. To obtain high accuracy in classification, the downsampling is done later in the network so that the activation map is large.

ShuffleNet [14] is created by Megvii Inc. (Face++) that promotes the CNN to achieve not only high accuracy but also high computational efficiency. It uses two components; pointwise group convolution and channel shuffle to realize the computational efficiency. The shuffle channel is where the feature map from the previous group is shuffled and feed to the different group of the next group convolution layer. They also introduce ShuffleNet Unit where pointwise group convolution is implemented before the shuffle operation. To recover the dimension, pointwise group convolution is added twice and 3x3 average pooling is added to increase the dimension of the channel with an adequate computation cost.

DenseNet [15] forms a network that achieves high accuracy with less parameters. This architecture introduces a method in which every layer receives input of feature map from all the previous layers. Hence, this enables the future layer to have bits of information from each layer before it. DenseNet creates a compact and slimmer network as the channel number can be reduced. Other than that, it utilizes computational cost and memory systematically. The last architecture is MobileNetV2 [16] that is created by Google as an improvement of the last MobileNetV1. This architecture introduces a module called inverted residual structure. This module creates a shortcut which is put at the bottleneck layer. It is used as input of low dimension that has been compressed and filtered with simple depthwise convolution. The feature is being casted back to a rendition in low dimension with a linear convolution. This architecture is said to be memory efficient.

3. Methodology
The compared architectures are tested to classify lung lesions in CT images in which the datasets are taken from the Lung Image Database Consortium image collection (LIDC-IDRI) [17]. As many as 1646 datasets are collected from this database that comprise of malignant and benign lesions. For each dataset, 4 annotations done by the radiologists are recorded and the lesion is labelled into 5 categories:
1. likely to be benign
2. fairly likely to be benign
3. intermediately likely to be benign and malignant
4. fairly likely to be malignant
5. likely to be malignant.

We extracted this information from the attached extensible markup language (XML) file of the datasets and classify the data into malignant if the average label value is more than 2.5 and vice versa. From the collected data, 1423 malignant cases and 223 benign cases are obtained. These data then are resized to 244x244 and are saved in PNG file. These data are segregated into 70% training and 30% testing. Figure 1 shows the example of two datasets after preprocessing.

![Figure 1 Example of datasets of the CT lung lesions after pre-processing](image)

MATLAB R2020a is used to train and test the deep learning architectures in classifying the lung lesions. Five architectures are tested and compared. They are GoogleLeNet, SqueezeNet, ShuffleNet, DenseNet and MobileNetV2. These networks are trained using NVIDIA GTX-1060 6 GB GPU. Table 2 shows the parameters used in the training process for each architecture. The parameters are common for all architectures for comparable comparison purpose.
Table 2 Training parameters

| Parameters            | Value |
|-----------------------|-------|
| Initial learn rate    | 0.01  |
| Validation frequency  | 50    |
| Max Epochs            | 20    |
| Mini batch size       | 64    |
| Learn rate drop factor| 0.2   |
| Learn rate drop period| 5     |
| Momentum              | 0.9   |

We evaluate the performance of the five networks based on classification metrics such as accuracy, specificity, sensitivity and area under the curve of the receiver operating characteristic curve (AUC). Accuracy is the measurement of the correct data predicted by the network divided by the whole data. This can be calculated as below:

\[
\text{Accuracy} = \frac{TP + TN}{TP + FP + FN + TN} \quad (1)
\]

Specificity is the number of data correctly labelled as negative over the whole negative data. The formula is as stated:

\[
\text{Specificity} = \frac{TN}{FP + TN} \quad (2)
\]

Sensitivity is the number of data correctly labelled as positive over the whole positive data. It is calculated as:

\[
\text{Sensitivity} = \frac{TP}{TP + FN} \quad (3)
\]

AUC is also being analyzed through the Receiver Operating Characteristics (ROC). This parameter is used to analyze the performance of the network to classify the data into their respective classes. AUC of 1 represents a perfect classification while 0 indicates a definite classification failure. Figure 2 shows the overall block diagram of the implemented classification using the CNN based networks.

Figure 2 Block diagram of the classification method using deep learning networks

4. Result

Table 3 tabulates the performance of the networks in terms of accuracy, specificity, sensitivity and AUC while Figure 3 shows the graphical comparison of the performance for each network. GoogLeNet architecture outperforms all the other architectures with an accuracy of 94.53%, specificity of 99.06% and sensitivity of 65.67%. GoogLeNet able to perform the best in accordance to the way the architecture is designed that is to limit the computational bottleneck. Therefore, there is an ease to run the architecture smoothly. The second best performed architecture is SqueezeNet with an accuracy of 94.13%, specificity of 99.06% and sensitivity of 62.69%. Both architectures implement a...
1x1 filter in their novel modules. It shows that by integrating 1x1 filter, it can improve the architecture performance as it lessens the burden of facing computational bottleneck.

The next performed classifier is DenseNet with accuracy of 93.52%, specificity of 89.83% and sensitivity of 59.7%. The propagation of input layer to the subsequent layers proves to have beneficial attributes as the front layers able to learn more features and increasing its accuracy. However, the 121 layers that it consists may not be suitable to be used on a simple single GPU computer as it can slow down the running time. ShuffleNet performance is not well received compared to the rest as it tackles the accuracy of 92.91%, specificity of 98.83% and sensitivity of 55.22%. The last out of all the architectures is MobileNetV2 with an accuracy of 92.91%, specificity of 97.65% and sensitivity of 62.64%. However, it still manages to achieve the accuracy of higher than 90%. This show that all the architectures perform greatly with their novelty modules that are designed for high accuracy while maintaining a good number of parameters.

*Figure 4* shows the ROC curve for each architecture. Based on the ROC curves, GoogleNet has the highest area under the ROC curve (AUC) of 86.84%. DenseNet achieves AUC of 86.12%, followed by SqueezeNet (85.04%), ShuffleNet (82.34%) and lastly MobileNetV2 (82.11%).

*Figure 5 to Figure 9* show the confusion matrix for each architecture in the classification task. The confusion matrix is used to recognise the correctly labelled data to their respected classes and the incorrectly labelled data to their respected classes. This subsequently be calculated in the form of specificity and sensitivity.

The next simulation result is an example of prediction by each architecture on a specific dataset. In this case, the dataset is malignant. *Figure 10 to Figure 14* show the prediction made by each architecture together with the accuracy values. It is seen that GoogLeNet, SqueezeNet and DenseNet predicted the data correctly while ShuffleNet and MobileNet missed the prediction. GoogleNet gives the highest prediction which is 0.99 while MobileNetV2 wrongly classifies the data into benign with an accuracy of 0.72.

**Table 3 Accuracy, specificity, sensitivity, AUC performance**

| Architecture   | Accuracy (%) | Specificity (%) | Sensitivity (%) | AUC (%) |
|----------------|--------------|-----------------|-----------------|---------|
| GoogleNet      | 94.53        | 99.06           | 65.67           | 86.84   |
| SqueezeNet     | 94.13        | 99.06           | 62.69           | 85.04   |
| ShuffleNet     | 92.91        | 98.83           | 55.22           | 82.34   |
| DenseNet       | 93.52        | 98.83           | 59.7            | 86.12   |
| MobileNetV2    | 92.91        | 97.65           | 62.64           | 82.11   |

*Figure 3* Comparison of accuracy, specificity, sensitivity and AUC of each network
Figure 4 Comparison of ROC curves between GoogleNet, DenseNet, SqueezeNet, ShuffleNet and MobileNet

Figure 5 GoogLeNet confusion matrix

Figure 6 SqueezeNet confusion matrix

Figure 7 ShuffleNet confusion matrix

Figure 8 DenseNet confusion matrix
In overall, it is observed that GoogLeNet architecture is the best deep learning classifier to classify the lung lesions in CT images into benign or malignant lesions followed by SqueezeNet, DenseNet, ShuffleNet and lastly, MobileNetV2. Here, our results are only based on LIDC-IDRI database. However, the results are reliable as LIDC-IDRI is the biggest online lung CT dataset available of all.
5. Conclusion
In this paper, the performance of five CNN architectures, namely GoogleNet, SqueezeNet, DenseNet, ShuffleNet and MobileNetV2 to classify the lung lesions in CT images into benign and malignant lesions have been compared and analyzed. The experiment results show that the GoogleNet architecture is the best deep learning classifier to classify the lung lesions in CT images into benign or malignant with an accuracy of 94.53%. The least performed architecture of all is MobileNetV2 with an accuracy of 92.91%. Therefore, further study on the GoogleNet network is required in order to improve the classification accuracy of lung lesions in CT images.

Acknowledgment
The authors would like to acknowledge the Center of Research and Instrumentation Management (CRIM), Universiti Kebangsaan Malaysia, Selangor, Malaysia and the Ministry of Education, Malaysia (MOE) for the Fundamental Research Grant Scheme (FRGS) with project code: FRGS/1/2019/TK04/UKM/02/5 to support this project.

Conflicts of interest
The authors have no conflicts of interest to declare.

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