

A modified residual network for detection and classification of Alzheimer's disease

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ABSTRACT

Alzheimer's disease (AD) is a brain disease that significantly declines a person's ability to remember and behave normally. By applying several approaches to distinguish between various stages of AD, neuroimaging data has been used to extract different patterns associated with various phases of AD. However, because the brain patterns of older adults and those in different phases are similar, researchers have had difficulty classifying them. In this paper, the 50-layer residual neural network (ResNet) is modified by adding extra convolution layers to make the extracted features more diverse. Besides, the activation function (ReLU) was replaced with (Leaky ReLU) because ReLU takes the negative parts of its input, drops them to zero, and retains the positive parts. These negative inputs may contain useful feature information that could aid in the development of high-level discriminative features. Thus, Leaky ReLU was used instead of ReLU to prevent any potential loss of input information. In order to train the network from scratch without encountering the issue of overfitting, we added a dropout layer before the fully connected layer. The proposed method successfully classified the four stages of AD with an accuracy of 97.49 % and 98 % for precision, recall, and f1-score.

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1. INTRODUCTION

Alzheimer's disease (AD) is a neurological disease. Its symptoms are a decline in memory, cognitive skill impairment, lack of reasoning and judgment, and difficulties with expression and comprehension [1]. The main reason for Alzheimer's disease is that the brains of patients have higher levels of amyloid [2], [3] and tau protein than the normal brains of older people [4], [5]. Accordingly, the increasing growth of amyloid plaques and tau neurofibrillary tangles leads to a loss of nerve cells' ability to communicate with one another and, eventually, cell death. The first brain area to be negatively impacted by a disease is the hippocampus [6]. Due to the hippocampus's critical role in learning and memory, forgetting people and events is the first sign of AD. The number of people who have Alzheimer's disease Precipitate will quickly grow as life expectancy grows. According to the world Alzheimer's report [7] statistics, an estimated 50 million people had Alzheimer's disease in 2015. By 2050, this number is expected to increase to 131.5 million people worldwide.

Presently, no treatment or drug can delay or stop the growth of Alzheimer's disease, necessitating and requiring effective and precise methods for early detection and preventing the disease from progressing to late stages. The practice of assigning items to a specific set based on their characteristics is called classification [8]. One of the specific goals of artificial intelligence research is to classify diseases [9]. A variety of neuroimaging methods are be of assistance in the classification of AD, for instance, magnetic

resonance imaging (MRI), computerized tomography (CT), functional MRI (fMRI), positron emission tomography (PET), magnetoencephalography (MEG), electroencephalography (EEG). MRI technologies are the most commonly used because images obtained by MRI show the affected cells darker than the healthy regions [10]. Figure 1 depicts several brain MRI images illustrating various stages of Alzheimer's disease. Dechter established the concept of deep learning (DL) in 1986 [11]. Its techniques, especially convolutional neural networks (CNN), have gained traction in a variety of fields, including image processing and analysis, and have achieved good performance in many computer vision tasks and medical imaging applications such as cerebral microbleeds (CMBs) detection [12].

This paper presents a modified version of the 50-layer residual neural network (ResNet) architecture that can solve the classification problem of four stages of Alzheimer's disease. Alzheimer's disease progressively deteriorates brain tissue in a predictable pattern. It reduces the size of the hippocampus and cerebral cortex of the brain while increasing the size of the ventricles [13]. Some outstanding research has been conducted on automated Alzheimer's disease diagnosis. The method of Jongkreangkrai *et al.* [14] consisted of two essential phases: the feature extraction phase, where they used MRI brain images to extract feature sets containing volumes of the hippocampus and amygdala as well as the thickness of the entorhinal cortex. After that, they moved on to the classification stage. They used the support vector machine algorithm to differentiate between Alzheimer's patients and healthy subjects based on their extracted features. Since the amount of medical data available is inadequate in comparison to other fields, as well as the fact that deep neural network training necessitates a large number of computing resources. Transfer learning is a promising alternative methodology developed by researchers. Jain *et al.* [15] used a pre-trained VGG16 network on the ImageNet dataset as a feature extractor for classifying brain MRI images into three categories: CN, MCI, MD, and AD as shown in Figure 1(a) to 1(d). Odusami *et al.* [16] introduced a new brain slice classification approach based on the ResNet18 algorithm. Puente-Castro *et al.* [10] used the first 47 layers of ResNet to extract features from sagittal MRI images. They then added the patient's age and gender to the feature vectors extracted by ResNet.

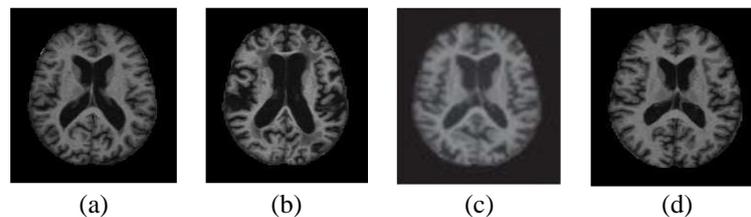


Figure 1. Brain MRI images for various stages of Alzheimer's disease: (a) cognitively normal (CN), (b) mild demented (MCI), (c) moderate demented (MD), and (d) very mild demented (AD)

The support vector machine (SVM) technique is then used as a classifier on these extracted feature vectors to determine whether the patient is in any stage of Alzheimer's disease or CN. Rabeh *et al.* [17] proposed an application for the early detection of Alzheimer's disease. The application framework has two steps: segmentation using region of interest (ROI) to isolate three critical regions: hippocampus, corpus callosum, and cortex. Following that, there is a classification step for components by using SVM and applying a decision tree to make the final decision. Ji *et al.* [18] developed a method for early diagnosis of Alzheimer's disease using MRI images of human brains and ResNet50, NASNet and MobileNet as base classifiers, which were trained in an end-to-end process

2. THE PROPOSED METHOD

Accuracy in medical diagnosis is more important than anything else, even more, important than speed of diagnosis. After all, the wrong diagnosis of an ordinary person as a patient causes severe consequences and psychological pressure, as well as diagnosing a patient, as usual, leads to the development of the disease because the wrong diagnosis, in this case, will delay treatment. Therefore, building an automated medical diagnostic system must be at a high level of accuracy.

2.1. Convolutional neural networks (CNN)

Convolutional neural networks are used in various applications, including image classification, segmentation, and pattern recognition [19], [20]. Due to its autonomous nature has developed into a critical tool for machine vision and artificial intelligence. CNN is a particular neural network that directly applies

image processing to the pixels without prior treatments [21]. The CNN architecture consists of three primary layers: the convolution layer, the pooling layer, and the fully connected layer. The convolution layer is the building block of the CNN algorithm. It is in charge of extracting the essential and beneficial features from the input images using a set of trainable filters, forming a feature map [22]. Between successive convolutions is a pooling layer used to reduce the feature mapping dimensions in computational space [23], thereby lowering the computational cost of the subsequent convolution layer, which contributes to the acceleration of training and the enhancement of generalization ability [24]. Typically, fully connected layers are inserted at the end of the CNN structure, utilized for recognition and classification. Every node in the fully connected layers is linked with trainable weights in adjacent fully connected layers. The fully connected layer produces a final output equal to the number of classes. CNN is a stacked version of all the layers that make up the CNN architecture. Each CNN, with a few exceptions, uses the same architecture.

Fully training a new CNN from scratch is not without its challenges. Firstly, CNN needs large amounts of labeled data for the training process that may be difficult to obtain, especially in medical imaging. In addition, to train a CNN, need to put in lots of computing and memory resources. Otherwise, the training process would take a long time without these resources. Tuning hyperparameters is time-consuming and complicated, and it can result in overfitting or underfitting, which leads to poor model performance. Researchers have demonstrated a promising alternative method known as transfer learning to overcome these obstacles. Transfer learning means improving the learning of a new task through transferring knowledge from a previously learned task [25]. The fundamental goal of this research is to compare the performance of pre-trained ResNet50 against the modified residual network in detecting and automatically classifying Alzheimer's disease using MRI scans.

2.2. Proposed framework

This section presents the general architecture of the proposed framework. The proposed framework consists of the following stages: data collection, data preparation, model selection, training stage to build the model that helps in diagnosis, validation, and evaluation. Each stage is independent of the other and is responsible for implementing a specific function. At the same time, these stages can communicate with each other since the result of one stage will be the input to the different stages. Figure 2 describes the proposed framework.

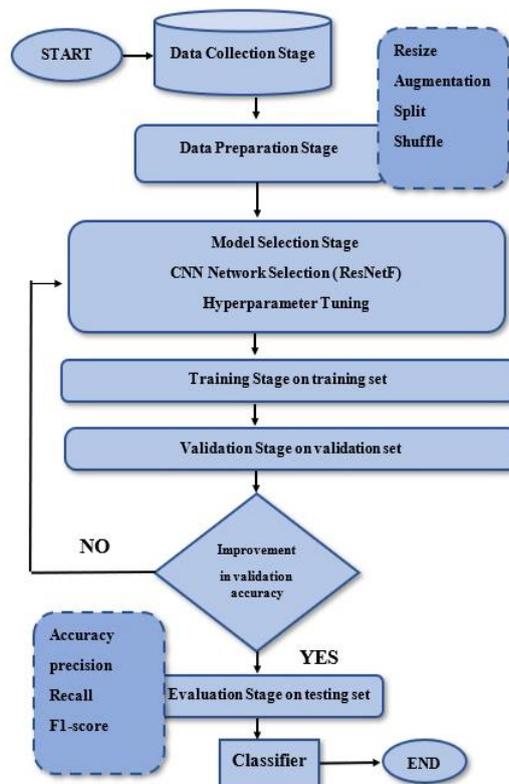


Figure 2. Flowchart of the proposed framework for early diagnosis of Alzheimer's disease

2.2.1. Data preparation

Data collection is the first step in the machine learning pipeline for training the selected model. The accuracy of ML systems' predictions is only good when the data used to train them is good. So, the first stage in the framework of the proposed work in this work is to collect data and obtain it from data sources related to this work in order to solve the research problem and evaluate the results. In this study, the Alzheimer's brain MRI dataset was obtained from the open access of the Kaggle website. The dataset contains 6400 images with a size of 176×208 pixels. It has four classes (NonDemented, MildDemented, ModerateDemented, VeryMildDemented) with a non-uniform distribution of the images per class.

The second step is changing size of image to (125×125) in order to decrease the time of training the neural network by lowering the number of pixels in an image because the more pixels in an image, the more input nodes there are, which raises the model's complexity. After that and because of the non-uniform distribution of images in each class, we generated new training examples using one of the data augmentation techniques only on the training set to improve deep neural network generalization capabilities and prevent overfitting. Horizontal flipping is the augmentation technique used in this step. This technique works by shifting all of the pixels in an image in the horizontal direction, or other words, by reversing the entire rows and columns of image pixels horizontally.

The data set is separated into two independent sets; the training set which has (5,121) images and the testing set which has (1,279) images. The training set was then partitioned into into a 20% validation dataset and an 80% new training dataset. The proposed model is trained using the "new training set". In contrast, the validation set periodically evaluates the model's performance during the training phase to avoid overfitting problems. The testing set is later used to evaluate how well the model generalizes to unseen data. Shuffling is the last step in data preparation. It regulates the weights, which means getting lower weights closer to zero. The most critical aspect is preventing the model from learning the training order. This step eventually helps the training converge quickly so that the network can provide better generalizations. During the validation phase and testing phase, there is no shuffling process for the model's parameters. During the validation and testing phases, we calculate accuracy and loss. Their calculation method is not sensitive to the order of samples, so shuffling does not affect the testing and validation data.

2.2.2. Network architecture

The residual neural network, known as ResNet, is a deep neural network that uses shortcuts, called "skip connections", to jump over some layers [26], [27]. ResNet has demonstrated outstanding performance in computer vision, so that it will be used in current research. He *et al.* [28] invented the ResNet in 2016, and it earned first place in the ILSVRC 2015 classification competition with a 3.57 error rate.

ResNet's hypothesis is that deeper networks are more difficult to optimize, as the deeper model should be capable of performing as well as the shallower model by copying the shallower model's learned parameters and setting additional layers for identity mapping. To aid in the optimization of deeper models, residual blocks are designed to fit a residual mapping $F(x)$ rather than the desired underlying mapping $H(x)$ to assist in the optimization of deeper models, and entire ResNet architectures are built by stacking residual blocks. Figure 3 illustrates the concept of a residual block. If we assume that the input is x , the convolution layer's output is $F(x)$, which is added to x as the mapping input, and the resulting output $H(x) = F(x) + x$ is passed to the next layer. This is significantly easier than matching an identity map through a collection of nonlinear layers, and it does not require the network to include additional parameters and calculations. Simultaneously, it can significantly increase the training speed and effectiveness of the model as the number of layers increases. This residual block structure can effectively solve the problem of gradient vanishing in deep networks [28]. There are two types of residual blocks in ResNet. While the first type is suitable for training shallow networks, the second type (bottleneck) is recommended for more than 50 layers. Additionally, the two types share a similar level of time complexity.

In this paper, an improved residual neural network based on ResNet50 (named ResNetF) has proposed. The number of convolution layers increased to 58 layers. More generally, as the number of network layers increased, the features that are extracted from the different layers become more diverse and richer. Additionally, the more deeply embedded the network is, the more abstract the features that are extracted. As a result of improving the network's feature extraction abilities, its effectiveness in AD diagnosis has improved. To ensure that overfitting is effectively avoided. We added a dropout layer before the fully connected layer in this architecture and set the dropout ratio to 50%. In ResNet50 ReLU is commonly used as an activation function. Basically, in CNN, ReLU takes the negative parts of its input and drops them to zero, and retains the positive parts. However, these negative inputs may contain useful feature information that could be used to aid in the development of discriminative high-level features [29].

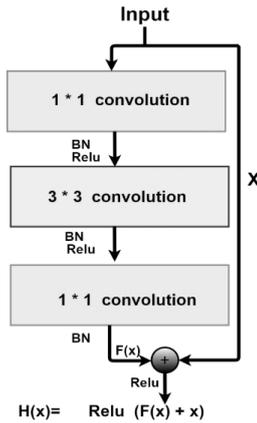


Figure 3. Structure of bottleneck block (for ResNet 50, 101,152)

If a neuron's output is 0, its gradient will never update the neuron's weight, resulting in the neuron never being activated. When the network contains a high number of inactive neurons, the convergence of the model becomes extremely difficult. Accordingly, this is referred to as the “dying ReLU” problem. This may prevent the network from learning and result in underperformance. To address this problem, Leaky ReLU is used instead of ReLU to prevent any potential loss of input information. Leaky Rectified Linear Activation (LReLU) has added an alpha parameter to the semi-axis of ReLU, resulting in a small gradient but not zero. Nodes that were previously inactive with ReLU will now have their weight-adjusted. Figure 4 shows our enhanced network. ReLU is given by (1), (2):

$$f(x_i) = \max(0, x_i) = f(x) = \begin{cases} x_i & x_i > 0 \\ 0 & x_i \leq 0 \end{cases} \quad (1)$$

$$f'(x) = \begin{cases} 1 & x_i > 0 \\ 0 & x_i \leq 0 \end{cases} \quad (2)$$

Leaky ReLU is defined as (3), (4):

$$f(x_i) = \begin{cases} x_i & x_i > 0 \\ a_i x_i & x_i \leq 0 \end{cases} \quad (3)$$

$$f'(x) = \begin{cases} 1 & x_i > 0 \\ a_i & x_i \leq 0 \end{cases} \quad (4)$$

where a_i is a predefined parameter that falls in the range of (0,1). It is usually 0.01, x_i is the activation function's input.

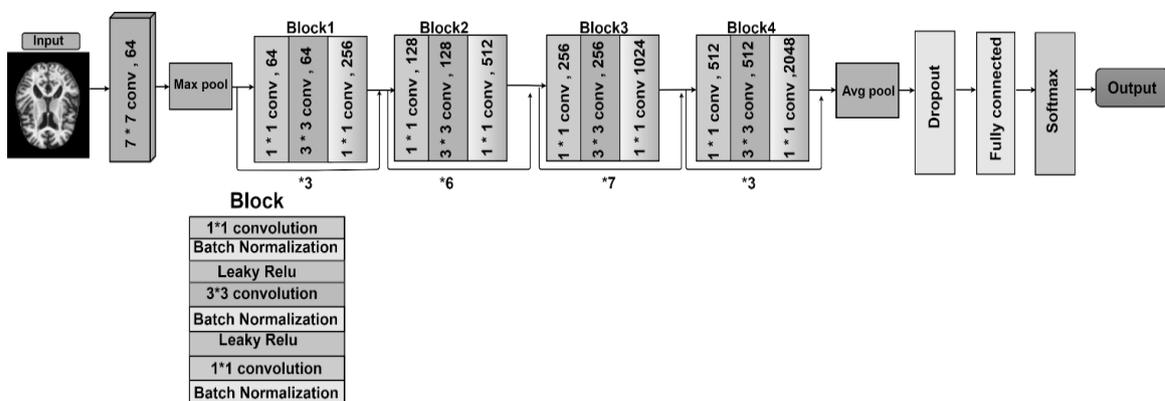


Figure 4. Modified residual neural network (ResNetF) structure

2.2.3 Training stage

In this stage, we used training set and the backpropagation algorithm to train our proposed network (ResNetF). We also used Adam as the optimizer with a learning rate of 1e-5, cross-entropy as the loss function. After each epoch during the training phase we assessed the model's performance by computing the validation accuracy and validation loss using our validation set.

2.2.4. Evaluation Stage

In this stage, popular criteria are used to assess the output of machine learning models to measure the classification model's ability to classify the testing dataset accurately. Accuracy, precision, recall, and F1-score are the common measurement tools, as mentioned in (5) to (8), respectively. True positive, false positive, and false negative predictions for a given class label are represented by the variables TP, FP, and FN, respectively [30], [31].

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

$$Precision = \frac{TP}{TP+FP} \quad (6)$$

$$Recall = \frac{TP}{TP+FN} \quad (7)$$

$$F1 - score = 2 \times \frac{Precision \times Recall}{Precision + Recall} \quad (8)$$

3. RESULTS AND DISCUSSION

This section explains the results of the research and, at the same time, gives a comprehensive discussion. The proposed network was trained for 13 hours, 54 minutes, 53 seconds. The Pytorch framework and Python 3.7.10 were used to run experiments on an Nvidia Tesla P100 GPU with 25 GB of memory.

To evaluate the classification model using testing data, various criteria are used, including precision, recall, the f1-measure, and accuracy to assess all stages of Alzheimer's disease. According to the results of our experiment, the training accuracy was 99%, and the validation accuracy was 97%. The results of the evaluation are summarized in Table 1.

The proposed model has been compared to a previous study [32] that addressed a similar issue and used the same data samples. We noticed that the model achieves encouraging results and surpasses the previous state-of-the-art in all criteria. The success can be attributed to expanding the number of convolutional layers, adding a dropout layer, and substituting an activation function (Relu) for an activated function (leaky Rule) in our modified network.

Table 1. Performance of the ResNetF model

Class	Model	Accuracy	Precision	Recall	F1-score
Non Demented	Yildirim and Cinar [32]	90	90	96.42	93.09
	ResNetF	96.56	99	97	98
Mild Demented	Yildirim and Cinar [32]	96.6	96.6	90.62	93.51
	ResNetF	96.64	99	97	98
Moderate Demented	Yildirim and Cinar [32]	70	70	70	70
	ResNetF	100	100	100	100
Very Mild Demented	Yildirim and Cinar [32]	90	90	90	90
	ResNetF	99	95	99	97

4. CONCLUSION

An accurate diagnosis of Alzheimer's disease allows the patient to receive the most appropriate treatment. This challenging task focuses on many researchers, who have built up many computer-aided diagnosis (CAD) systems to diagnose AD. This paper presented an enhanced residual neural network to classify four stages of Alzheimer's disease. By increasing the number of convolution layers, the network can effectively capture as many AD biomarkers as possible. In addition, substituting Relu for the activation function (Leaky Relu) can solve losing valuable features that could assist in the construction of high-level discriminative features. To avoid overfitting, we added a dropout layer before the fully connected layers to train all our architecture layers from scratch. The findings of the experiments demonstrate that the enhanced residual neural network is suitable for Alzheimer's disease diagnosis.

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