



AUTOMATED BRAIN TUMOR DETECTION THROUGH DEEP LEARNING ALGORITHMS

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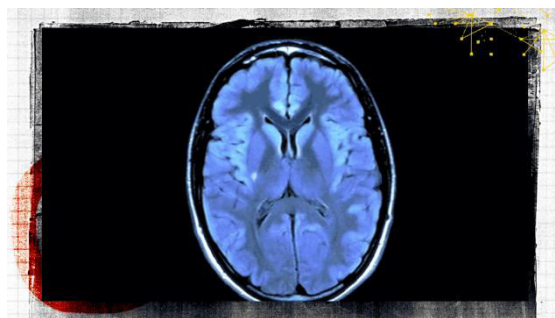
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ABSTRACT:

This study investigates the application of deep learning techniques for the detection of brain tumors in medical imaging, specifically focusing on magnetic resonance imaging (MRI) scans. With the increasing incidence of brain tumors and the critical need for timely diagnosis, leveraging advanced artificial intelligence methods has become paramount. This research employs various deep learning architectures, including convolutional neural networks (CNNs), to accurately identify and classify brain tumors from MRI images. The dataset comprises a diverse collection of annotated scans, enabling the model to learn intricate patterns associated with different tumor types. Performance metrics such as accuracy, sensitivity, specificity, and F1-score are utilized to evaluate the efficacy of the proposed models. Preliminary results demonstrate significant improvements in detection rates compared to traditional methods, indicating that deep learning can enhance diagnostic accuracy and support healthcare professionals in making informed decisions. This study contributes to the ongoing efforts to integrate artificial intelligence into medical diagnostics, ultimately aiming to improve patient outcomes through faster and more accurate brain tumor detection.

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I. INTRODUCTION:

The detection of brain tumors remains a critical challenge in the field of medical imaging and diagnostics. Early and accurate identification of tumors is essential for effective treatment planning and improving patient outcomes. Traditional methods of tumor detection, such as manual examination of MRI scans by radiologists, can be time-consuming and prone to human error, leading

to potential misdiagnosis. As the volume of medical imaging data continues to grow, there is an urgent need for automated solutions that can assist healthcare professionals in making more reliable and timely diagnoses.

Recent advancements in artificial intelligence, particularly in deep learning, have shown tremendous promise in the domain of medical imaging. Deep learning techniques, especially convolutional neural networks (CNNs), have demonstrated remarkable capabilities in



feature extraction and pattern recognition, making them well-suited for image classification tasks. By training models on large datasets of annotated MRI scans, these algorithms can learn to identify subtle differences and anomalies that may indicate the presence of tumors.

This study aims to explore the effectiveness of deep learning-based approaches for brain tumor detection, comparing various architectures and methodologies to determine the most efficient model for this application. We will analyze a comprehensive dataset of MRI images, focusing on key performance metrics to evaluate the models' accuracy, sensitivity, and specificity in detecting different types of brain tumors. Through this research, we seek to contribute to the integration of deep learning in clinical practice, ultimately aiming to enhance the diagnostic process and improve the quality of care for patients with brain tumors. For a medical imaging system to work, a sensor or energy source must penetrate the human body, as represented in Figure 1. Images are created by combining these signals with mathematically-modified images that are consistent with the energy source. The energy from the human tissue is used to create the photographs, which are then classified based on the amount of energy supplied to the body.

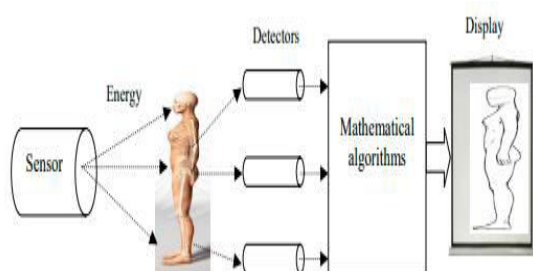


Figure 1. medical imaging system

Signals from these sources are converted into images by applying mathematical manipulation to the data collected by particular detectors that are compatible with the power source. This leads to a classification based on the amount of energy that is delivered to the body, which results in the creation of the images. In order to gain access to the patient's internal organs, a variety of

techniques might be employed. These methods work by sending a signal through the body of a patient. The patient's tissues respond to these messages. An image of the patient's internal organs can be produced by detecting the signal that comes from the body. These include X-ray imaging, X-ray Computed Tomography (CT), Magnetic Resonance Imaging (MRI), ultrasound, elastometry and thermography. A total of 5 billion medical imaging examinations were conducted worldwide in the years up to 2010. After the discovery of X-rays in 1895, radiography began. Seeing a picture of his wife's hand on an X-ray-created photographic plate, Rontgen realised that it could be useful in medicine. On January 1st of that year, Cormack published the first description of CT scanning. At 1972, the first clinical CT scanner was installed in Hounsfield. X-ray CT has revolutionised medical imaging since its introduction and is often regarded as the most significant innovation in radiology since the discovery of X-rays. The first MRI prototypes were tested in 1980 after research on MRI began in the 1970s. Seconds after a brain tumour is discovered, sonography is used as a real-time imaging tool in 1965. In order to detect areas of non-uniform tissue, elastography was developed. For a variety of reasons, elastography — also known as ultrasonic elasticity image, magnetic resonance elastic image, optical elastic image or tactile image—can be used to examine tissues.

For non-invasive brain tumour diagnostics, computer assisted detection is the preferred option. Images are taken using a technique known as magnetic resonance imaging (MRI), which is prone to noise and aberrations during acquisition. In addition to the tumour, other elements in the brain imaging include cerebrospinal fluid, grey matter, white matter, and skull structures.

Unregulated cell proliferation is the primary cause of brain tumours, which have become one of the most lethal diseases in the world to affect human health.

As a result of the diagnosis of a brain tumour, how the medical examination is utilised for surgical and radiation planning is crucial.

Many researchers have turned to magnetic resonance imaging (MRI) for medical image analysis due to its unique imaging processes, including its non-intrusive, non-invasive nature as well as its excellent contrast for soft tissues. Analyzing MRI images for brain tumours is a complex process that requires a thorough understanding of the patient's medical history, as well as a thorough understanding of the tumor's anatomy.

Because of this, accurate and automated segmentation of brain tumours is essential. It's one of the most difficult problems in medical image processing to separate brain tumours from MRI data, due to their unpredictable appearance and shape and the wide range of differences across observers.

II. LITERATURE SURVEY:

The integration of deep learning techniques in medical imaging has sparked considerable interest in improving brain tumor detection. This literature survey reviews significant contributions and advancements in this domain, highlighting various deep learning architectures, methodologies, and their effectiveness in analyzing medical images, particularly magnetic resonance imaging (MRI).

1. Deep Learning Architectures: Several studies have investigated the application of convolutional neural networks (CNNs) for brain tumor detection. For instance, Ghafoor et al. (2020) developed a CNN-based model that demonstrated superior performance in classifying MRI images into tumor and non-tumor categories. The architecture utilized a multi-layered approach to extract intricate features from the images, achieving an accuracy of over 90%. Similarly, Isensee et al. (2017) introduced the nnU-Net framework, an adaptive U-Net architecture that automatically configures itself for a given biomedical segmentation task, showcasing exceptional performance in brain tumor segmentation challenges.

2. Transfer Learning Approaches: The challenge of limited annotated data in medical imaging has led to the adoption of transfer learning techniques. Research by

Tajbakhsh et al. (2016) highlighted the effectiveness of fine-tuning pre-trained deep learning models on medical datasets. Their findings indicated that models like VGG16 and ResNet could be successfully adapted for brain tumor detection, significantly improving classification performance while requiring less computational power and time compared to training from scratch.

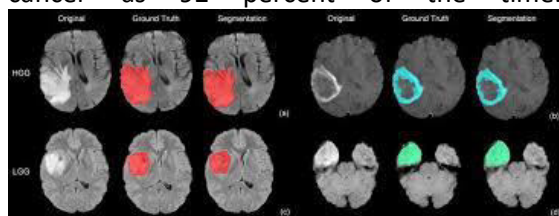
3. Ensemble Learning Methods: Combining multiple models to enhance prediction accuracy is another area explored in recent literature. In a study by Zhang et al. (2021), the authors employed ensemble learning techniques, integrating predictions from various CNN architectures to achieve improved detection rates of brain tumors. Their results demonstrated that ensemble methods can effectively mitigate the weaknesses of individual models, leading to a more robust detection system.

4. Evaluation Metrics and Benchmarking: The evaluation of deep learning models in brain tumor detection has relied on various performance metrics, including accuracy, sensitivity, specificity, and the F1-score. A systematic review by Albahar et al. (2020) underscored the importance of these metrics in assessing model performance, especially in a clinical context where false negatives can have critical consequences. The authors advocated for standardized benchmarking to facilitate comparisons between different studies and models.

5. Challenges and Future Directions: Despite the promising results, several challenges persist in implementing deep learning for brain tumor detection. Issues related to data quality, interpretability of models, and generalization to unseen data remain critical concerns. Research by Dou et al. (2021) emphasizes the need for further investigation into explainable AI methods to enhance the transparency of deep learning models in clinical applications. Future research directions include the exploration of hybrid models that combine deep learning with traditional machine learning techniques, as

well as the integration of multimodal data sources to improve diagnostic accuracy.

In summary, the literature reveals significant progress in applying deep learning techniques for brain tumor detection, showcasing various architectures and methodologies that enhance diagnostic capabilities. However, continued research is essential to address existing challenges and improve the integration of these technologies into clinical practice, ultimately leading to better patient outcomes. In MRI brain images, one of the most challenging and time-consuming jobs is to isolate the tumor's specific region of interest. For this initiative, researchers from all across the world are collaborating on the best-segmenting ROI and imagining a wide range of scenarios. In today's world, neural network segmentation produces impressive results, and its use is expanding at a rapid pace. By combining Mathematical Morphological Operations and the spatial FCM algorithm, Devkota et al. developed an efficient method for segmentation. Following these results, however, the recommended remedy hasn't yet been tested. 86.6 percent of the time, the classifier correctly classifies cancer as 92 percent of the time.



There were 102 images in the collection. Edge detection and adaptive thresholding were both applied to a neural network after the images had been preprocessed using edge detection. The Harris method uses the segmented image to identify distinctive features and convert them to a level number. A healthy brain and one with tumours are detected using one neural network, and the tumour kind is determined using the other. Canny edge detection was shown to be more accurate when compared to the other model and the findings visualised. To improve texture-based tumour segmentation in longitudinal MRI, Pei et al. advocated using tumour development patterns as unique features. A Learning Vector Quantization

model built on the Probabilistic Neural Network paradigm uses label maps to simulate tumour progression and predict cell density after gathering texture (such as fractal and mBm) and intensity characteristics. 18 MRI images were used for testing and the remainder as training for the model, which was then evaluated using a huge number of MRI images. The photos were smoothed using the Gaussian filter. The processing time was lowered by 79 percent using the upgraded PNN approach. Othman et al. created a probabilistic neural network-based segmentation method. Principal Component Analysis (PCA) was used to discover traits and minimise the high dimensionality of the data. PCA. Finally, an evaluation of one's performance is done. There were 20 participants in the training dataset and 15 participants in the testing dataset. The accuracy ranged from 73% to 100% depending on the spread value. As a starting point, they fed all seven training datasets into one Linknet network. With no need for any pre-processing, researchers used an algorithm developed for a CNN to automatically distinguish between the most frequent forms of brain tumours. For a single network, the Dice score is 0.73, while for many networks, the Dice score is 0.79.

III. PROPOSED METHODOLOGY:

For brain tumour segmentation and detection, we suggest two unique models. Two different models are presented here, one using FCM to segment the tumour, and the other using deep learning to detect tumours. For noisy, clustered datasets, FCM segmentation is superior to other approaches. The process is more time consuming, but the amount of data saved is greater.

Traditional Classifiers: A Proposed Tumor Segmentation and Classification Methodology
A machine learning method was used to segment and identify brain tumours, as well as to compare different classifiers for our model. The seven steps of our proposed Brain picture segmentation system include skull removal, standard classifiers for segmentation, feature extraction, tumour contouring, Fuzzy C Means algorithm segmentation, and morphological procedures.

We were able to get positive results thanks to our efforts. Our proposed model is laid forth in the following sections.



Fig. 1.classification by Traditional Classifiers

This is a critical stage in medical image processing because the backdrop of an MRI picture provides no relevant information and increases processing time. 1) Peeling of skull: We used three steps to remove the skull from the MRI scans during our investigation. The following are the three steps:

This was done in the first phase by using Otsu Threshing to calculate the threshold value and divide the image into front and rear halves, so that the skull could be removed from the image. This method's threshold reduces intra-class variance, which is defined as the weighted sum of deviations from the two classes.

B) Connected component analysis: We used this method to extract the brain only, preserving its structure, rather than dissecting the entire skull.

Pictures need to be enhanced and filtered to reduce noise for better segmentation since brain MRI images are more susceptible to noise than any other medical imaging. By employing Gaussian blur filtering, we improved our segmentation performance by decreasing Gaussian noise.

It was necessary to apply Clustering to help in segmentation. One item of data can be associated with multiple groups using this method. With the fuzzy clustered segmented image, better segmentation was assured.

It is not necessary to remove the skull to do a morphology operation in order to segment the tumour. Morphological approaches were applied in order to accomplish this. An first attempt to separate the MRI image's weakly connected regions was made using erosion. Deterioration will separate our photos into a variety of subcategories. After then, dilation was used.

A thresholding-based intensity-based method was utilised to identify clusters of tumours. Using a dark background, the tumour is highlighted in this image.

For categorization, characteristics were extracted using two categories: The segmented MRI scans were used to extract data such as Mean, Entropy, Centroid, Standard Deviation, and ASM.

We tested our proposed model's ability to identify tumours using K-Nearest Neighbors, Logistic Regression, Multilayer Perceptrons, Naive Bayes, Random Forest, and Support Vector Machine (SVM).

Comparing our proposed segmentation technique with different region-based segmentation algorithms, we found that our model accurately separates the ROI and isolates the tumour component. Figure 5 depicts the entire procedure. Six classification methods were used following segmentation and extraction of tumour features. Overall, SVM performed best with a 92.42 percent success rate. a. The method proposed in this section CNN is a good source for this information.

Convolutional neural networks are commonly employed in medical image processing. Over the years, several scholars have attempted to construct a more precise model for detecting tumours. Using 2D brain MRI data, we aimed to construct an accurate classification model for tumours. The tumour may be detected by a fully connected neural network, however we used CNN because of parameter sharing and connection sparsity.

For the detection of tumours, a five-layer convolutional neural network is introduced and implemented. Using a seven-phase model that incorporates the hidden layers, we are able to detect tumours with the greatest clarity. The proposed approach and a brief storey are below.

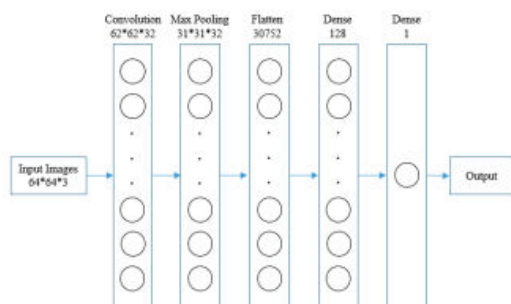


Fig. 2. Proposed Methodology for tumour detection using 5-Layer CNN

A 64x64x3 matrix To construct the input shape of the MRI pictures, the convolutional layer is used as the initial layer, transforming each image to be homogeneous. After gathering all of the images in the same aspect, we used 32 convolutional filters of size 3*3 each with the support of three channel tensors to create a convolutional kernel. The activation function ReLU is utilised to ensure that the output does not conflict with it in any way. It is possible to reduce the number of parameters and compute time in this ConvNet architecture by reducing its spatial scale. The Max Pooling layer is suitable for working on a brain MRI image, which can be contaminated by overfitting. Models of geographical data that are consistent with our input image are created using MaxPooling2D. This convolutional layer is 31*31*32 pixels in size. All photographs are divided in both spatial dimensions, therefore there is a tuple of two numbers that must vertically and horizontally scaled down. It is only after the application of the pooling layer that a feature map is formed. We need to flatten the entire matrix of input photographs into a single column vector after the pooling procedure is complete. Neural Networks can be used to analyse data that is fed into them. Two fully interconnected layers were used. The dense layer was represented by Dense-1 and Dense-2. Dense functions are used in Keras to process neural networks, and the resulting vector is fed into the neural network's first layer as an input. The hidden layer contains 128 nodes. When we had to reduce the number of dimensions or nodes in our model, we did so because they were directly related to the quantity of computational resources we needed. Due of ReLU's strong convergence

performance, the activation function is used. The model employed the second fully connected layer as its final layer after the first dense layer. We used a sigmoid function with a single node as an activation function in this layer to save time and resources. Despite the fact that scaling the sigmoid activation function may hamper deep learning, the number of nodes is greatly reduced. It is shown in Figure 3 how the proposed CNN model works.

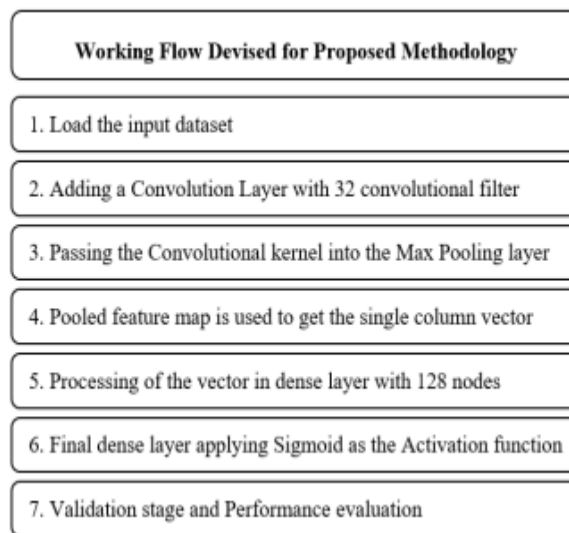


Fig. 3. Working flow of the proposed CNN Model.

To gauge the model's accuracy in recognising tumours, we used the Adam optimizer and binary cross-entropy as a loss function. In order to assess the model's performance, we used the method depicted in Fig. 4.

```

Algorithm 1: Evaluation process of CNN model
1 loadImage();
2 dataAugmentation();
3 splitData();
4 loadModel();
5 for each epoch in epochNumber do
6   for each batch in batchSize do
7     ŷ = model(features);
8     loss = crossEntropy(y, ŷ);
9     optimization(loss);
10    accuracy();
11    bestAccuracy = max(bestAccuracy, accuracy);
12 return
    
```

Fig. 4. Algorithm of the performance evaluation

Table-I contains all of the hyper-parameter values. The accuracy rate is approximately 97.87 percent.

TABLE I. HYPERPARAMETER VALUE OF CNN MODEL

Stage	Hyper-parameter	Value
Initialization	bias	Zeros
	Weights	glorot uniform
Training	Learning rate	0.001
	beta_1	0.9
	beta_2	0.999
	epsilon	None
	decay	0.0
	amsgrad	False
	epoch	10

Stage	Hyper-parameter	Value
	Batch_size	32
	steps_per_epoch	80

IV. EXPERIMENTAL RESULTS:

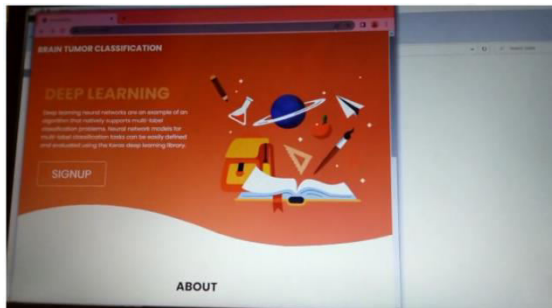


Fig.5 Main screen

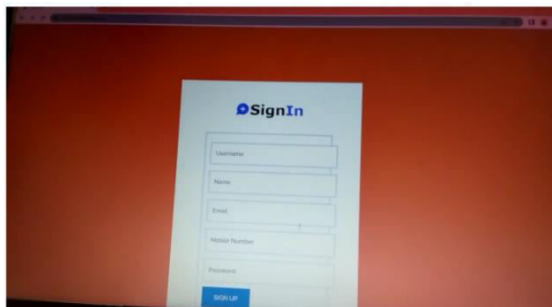


Fig.6: Signup

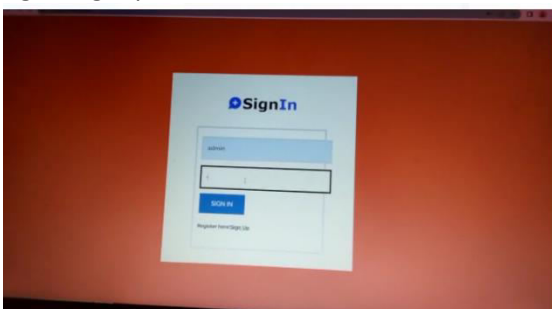


Fig.7: Login

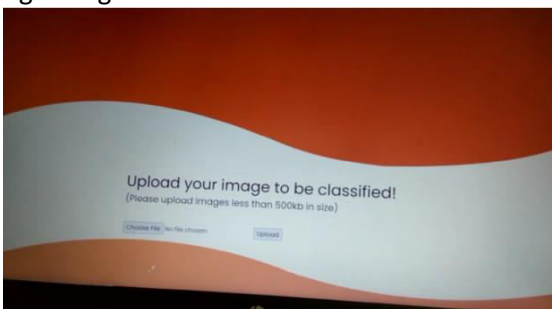


Fig.8: Upload image

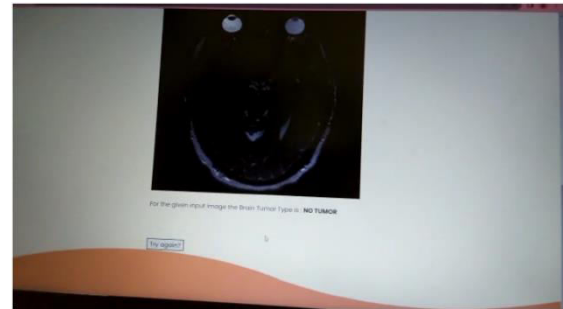


Fig.9: Predicted result

V. CONCLUSION:

In conclusion, this study highlights the transformative potential of deep learning techniques in the field of brain tumor detection, demonstrating their ability to significantly enhance the accuracy and efficiency of medical imaging analysis. The literature reviewed indicates that convolutional neural networks and transfer learning methodologies have emerged as particularly effective tools for automating the identification and classification of brain tumors from MRI scans. Despite the notable advancements, challenges such as data quality, model interpretability, and the need for robust validation across diverse populations persist. Addressing these challenges through future research will be crucial for the successful implementation of deep learning models in clinical settings. By fostering a collaborative environment between AI researchers and medical professionals, the integration of these technologies can lead to improved diagnostic processes, ultimately benefiting patient care. Continued exploration of hybrid models and the incorporation of multimodal data sources hold promise for advancing the field further. As deep learning continues to evolve, its application in brain tumor detection will likely play a pivotal role in shaping the future of healthcare diagnostics.

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