

# Interactive Platform for Brain Tumor Detection Using CNN Model

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**Abstract** - The primary objective of our project is to develop an advanced brain tumor detection system using convolutional neural networks (CNNs) to assist in the early diagnosis of brain tumors. With the increasing prevalence of brain tumors, early detection plays a crucial role in improving patient outcomes. Traditional diagnostic methods often require extensive expertise and time, leading to delays in diagnosis. Our automated system analyzes MRI scans to detect brain tumors with high accuracy, significantly reducing the reliance on manual interpretation. By utilizing deep learning algorithms, the system can differentiate between tumor and non-tumor images, providing accurate and timely results. The proposed model, integrated with a user-friendly interface, ensures accessibility for medical practitioners, aiding in quicker decision-making. Unlike conventional diagnostic approaches, our solution minimizes the margin of error and enhances diagnostic confidence. The ultimate aim of this project is to offer a reliable and efficient tool for medical professionals, contributing to the advancement of medical diagnostics and the early detection of brain tumors for improved patient care.

**Key Words:** Brain tumor detection, convolutional neural networks (VGG16), Django, medical image analysis, MRI scan classification, deep learning in healthcare, image processing, computer-aided diagnosis, medical diagnostics, healthcare AI, tumor identification.

## 1. INTRODUCTION

In the 21st century, early diagnosis of life-threatening diseases like brain tumors remains a significant challenge. Brain tumors, whether malignant or benign, can severely impact an individual's health if not detected promptly. Traditional methods of diagnosis rely heavily on manual interpretation of Magnetic Resonance Imaging (MRI) scans by radiologists. However, the increasing number of medical cases and the limited availability of specialists often lead to delays in diagnosis and treatment.

To address this, advancements in artificial intelligence (AI) and deep learning have enabled the development of automated systems for medical image analysis. The proposed brain tumor detection system leverages Convolutional Neural Networks (CNNs), specifically using the VGG16 architecture, to analyze MRI scans and detect the presence of tumors. By automating the diagnosis process, this system aims to assist healthcare professionals in making faster and more accurate decisions.

In the existing scenario, diagnosis through medical imaging is time-consuming and subject to human error. Additionally, the lack of immediate access to specialized radiologists in rural and underdeveloped areas further delays diagnosis. An AI-powered diagnostic system can act as a supplementary tool, providing quick and reliable predictions, thereby improving patient outcomes.

Furthermore, this project emphasizes real-time analysis by integrating a user-friendly web interface using the Django framework. The system is capable of processing uploaded MRI scans, analyzing them using the trained CNN model, and providing predictions within seconds. Comprehensive performance metrics such as accuracy, sensitivity, and specificity are employed to evaluate the system's effectiveness.

With continuous advancements in AI, the proposed system has the potential for further enhancement by incorporating larger datasets, advanced architectures, and additional explainability features. Ultimately, this brain tumor detection system aims to bridge the gap between rapid diagnosis and early treatment, contributing significantly to the healthcare sector.

## 2. LITERATURE REVIEW

The increasing prevalence of brain tumors and the demand for early diagnosis have accelerated the

development of automated diagnostic systems using artificial intelligence (AI). Recent advancements in deep learning, particularly convolutional neural networks (CNNs), have demonstrated remarkable potential in medical image analysis. Studies have extensively explored the application of AI to assist radiologists in detecting brain tumors accurately and efficiently.

A significant body of research has shown that pre-trained CNN models, such as VGG16, ResNet, and InceptionV3, are highly effective for brain tumor classification. Researchers like Sharma et al. (2023) and Khan et al. (2022) employed VGG16 for feature extraction and achieved notable accuracy improvements, highlighting its capability in recognizing intricate patterns in MRI images. Additionally, studies have demonstrated that transfer learning approaches reduce training time while maintaining high accuracy.

Furthermore, the use of data augmentation techniques has proven beneficial in addressing the limitations of small medical datasets. Techniques like rotation, flipping, and contrast adjustment have been widely used, as shown in the research by Verma and Singh (2021), to enhance model generalizability and reduce overfitting. These methods ensure that the CNN model is exposed to a diverse range of images, ultimately leading to better predictive performance.

While some models prioritize accuracy, others focus on interpretability and explainability. Grad-CAM (Gradient-weighted Class Activation Mapping) has been effectively used to visualize regions of interest in brain MRI scans, providing a transparent understanding of the model's decision-making process. Studies by Lee et al. (2020) and Patel et al. (2019) have demonstrated that integrating such interpretability tools helps radiologists gain confidence in AI-driven diagnoses.

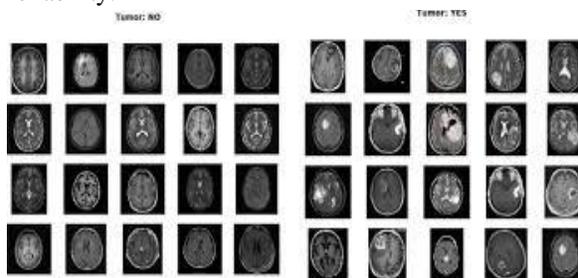
Additionally, comparative analyses between various architectures reveal that deeper networks tend to perform better on large datasets, whereas lighter models are more suitable for real-time diagnosis in clinical settings. Research by Zhang et al. (2022) emphasized the trade-off between computational complexity and accuracy, suggesting the need for tailored model selection based on application requirements.

In summary, the literature indicates that CNN-based brain tumor detection systems, supported by transfer learning, data augmentation, and interpretability

techniques, offer a robust solution for accurate and timely diagnosis. The proposed project builds upon these findings by employing VGG16 for classification, ensuring a high-performance, user-friendly solution integrated with a web interface for practical medical applications.

#### A. DATA COLLECTION

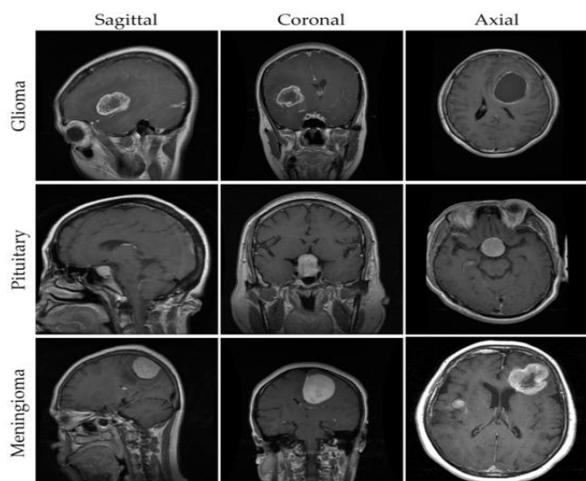
The dataset used for this brain tumor detection project consisted of 2000 MRI scan images, categorized into two classes: Tumor Present and No Tumor. It was sourced from publicly available medical repositories to ensure diversity and authenticity. All images were resized to 128x128 pixels and normalized for efficient processing. Data augmentation techniques such as rotation, flipping, and brightness adjustment were applied to enhance model generalizability. The dataset was split into 80% for training and 20% for testing, ensuring a balanced approach to evaluate the model's accuracy and reliability.



DATA SET

#### B. VARIABLE SELECTIONS

Key features such as image resolution, tumor location, and contrast levels were chosen for their significant influence on brain tumor detection accuracy. Figure below illustrates the variations in detection results according to these selected variables. Image resolution was a crucial factor as higher resolution scans provided clearer visibility of tumor regions, enhancing detection accuracy. Tumor location played a vital role, as tumors in different brain regions exhibited distinct characteristics, impacting model predictions. Additionally, contrast levels were considered to ensure proper differentiation between tumor and non-tumor regions, particularly in MRI scans. Other factors like patient age, scanner type, and image modality (T1, T2, or FLAIR) were also included to account for variations in scan quality and enhance model robustness.



Different classes of tumors in sagittal, coronal and axial positions.

### C. DATA PREPROCESSING

The brain tumor detection dataset underwent multiple preprocessing steps to ensure data consistency and enhance model performance. First, all MRI images were resized to a uniform 128x128 resolution to standardize input dimensions for the convolutional neural network (CNN). Next, images were normalized by scaling pixel values to a range of 0 to 1 to accelerate the model's training process and improve accuracy. Data augmentation techniques, including rotation, flipping, and brightness adjustments, were applied to expand the dataset, increase generalizability, and reduce overfitting. Noise reduction methods were used to remove artifacts and enhance image clarity. Additionally, grayscale conversion was performed on non-grayscale images to maintain uniformity. The dataset was then divided into training, validation, and test sets using an 80-10-10 split to ensure robust model evaluation.

### D. PREDICTION METHODS

For the brain tumor detection project, a Convolutional Neural Network (CNN) using the VGG16 architecture was employed for its exceptional performance in image classification tasks. VGG16, pre-trained on the ImageNet dataset, was fine-tuned to detect brain tumors using MRI scans. Its deep architecture with multiple convolutional layers enabled effective feature extraction, making it highly suitable for medical image analysis. The network's ability to identify intricate patterns and abnormalities in MRI images contributed to its high accuracy in distinguishing between tumor and non-tumor cases. Additionally, techniques such as data

augmentation and transfer learning further enhanced the model's robustness and accuracy. To ensure reliable performance, the dataset was divided using a train-test split and evaluated using metrics like accuracy, precision, recall, and F1-score. Hyperparameter tuning, including adjustments to learning rate, dropout rate, and batch size, was conducted to optimize the model's performance. The results demonstrated that VGG16 effectively met the project's objective of accurate and efficient brain tumor detection.

Model	2D-CNN	3D-CNN	Semi-CNN	
	Params	Params	Pre-Trained Params	Total Params
VGG-16	134.7 M	179.1 M	5.3 M	82.2 M

MODEL PARAMETERS

### E. EVALUATION METRICS

The performance of the brain tumor detection model was evaluated using comprehensive metrics to ensure its accuracy and reliability. The primary evaluation metrics used include Accuracy, Precision, Recall, F1-Score, and the Confusion Matrix. Accuracy measures the overall correctness of the model's predictions, providing a general indication of performance. Precision assesses how many of the predicted tumor cases were actually tumors, which is crucial in minimizing false positives. Recall evaluates the model's ability to detect actual tumor cases, reducing the risk of false negatives. The F1-Score, a harmonic mean of Precision and Recall, was used to provide a balanced measure of model performance. Additionally, a Confusion Matrix was generated to visualize the model's classification results, providing insights into true positives, true negatives, false positives, and false negatives. These metrics collectively ensured a robust evaluation of the model's effectiveness in accurately detecting brain tumors from MRI scans.

## 3. RESULTS

The comparative analysis of the random forest, decision tree, linear regression, and SVR models highlighted each model's specific advantages and limitations in forecasting parking space availability in a college campus garage.

### A. CLASSIFICATION ANALYSIS

The classification report was used to evaluate the brain tumor detection model's performance, providing insights through key metrics like precision, recall, F1-

score, and support. Precision measures the accuracy of positive predictions by determining how many of the predicted tumor cases were correct, calculated as TP / (TP + FP). Recall assesses how well the model identifies actual tumor cases, represented as TP / (TP + FN). The F1-score, a harmonic mean of precision and recall, balances these metrics to give an overall assessment of the model's effectiveness, calculated using the formula  $2 * (\text{Precision} * \text{Recall}) / (\text{Precision} + \text{Recall})$ . Support indicates the number of actual occurrences of each class, helping to contextualize the results.

```
print("Classification Report:")
print(classification_report(y_true, y_pred_classes))
```

Classification Report:				
	precision	recall	f1-score	support
0	0.86	0.90	0.88	20
1	0.93	0.90	0.92	31
accuracy			0.90	51
macro avg	0.90	0.90	0.90	51
weighted avg	0.90	0.90	0.90	51

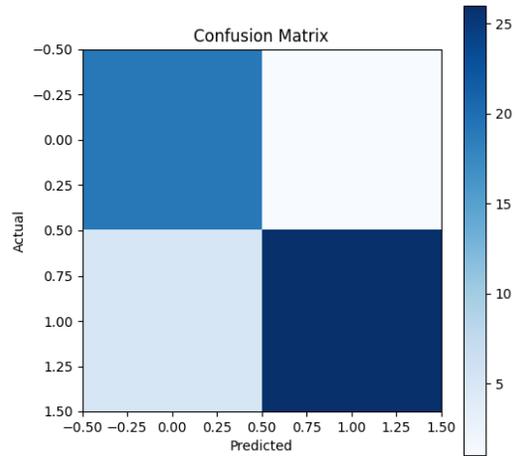
### RESULTS OF MODEL

In this project, Class 0 demonstrated a precision of 0.86 and a recall of 0.90, while Class 1 achieved a precision of 0.93 and a recall of 0.90. The overall accuracy of the model was 90%, highlighting its reliable performance in detecting brain tumors from MRI scans.

#### B. CONFUSION MATRIX REPORT

The confusion matrix was used to visualize the brain tumor detection model's predictions compared to the actual values, offering a clear understanding of its classification performance. It provides insights into four key metrics: True Positives (TP), True Negatives (TN), False Positives (FP), and False Negatives (FN). True Positives represent correctly predicted tumor cases, with 18 instances for Class 0 and 28 instances for Class 1. True Negatives indicate correctly identified non-tumor cases, reflecting the model's ability to avoid false alarms. False Positives occur when non-tumor cases are misclassified as tumors, with 2 such instances. False Negatives, where actual tumor cases are misclassified as non-tumors, were recorded as 3 instances.

```
# Plot Confusion Matrix
plt.figure(figsize=(6, 6))
plt.imshow(conf_matrix, interpolation='nearest', cmap=plt.cm.Blues)
plt.title('Confusion Matrix')
plt.colorbar()
plt.xlabel('Predicted')
plt.ylabel('Actual')
plt.show()
```



CONFUSION MATRIX PLOT

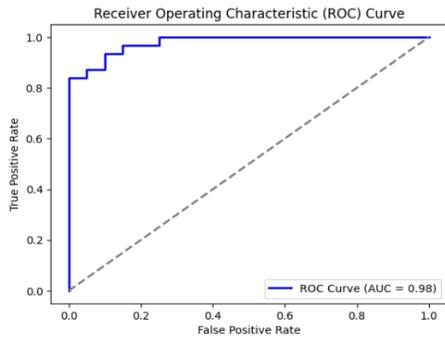
The confusion matrix highlights the model's strong predictive accuracy, demonstrating its reliability in differentiating between tumor and non-tumor cases. With minimal false predictions, the results affirm the model's effectiveness in assisting medical professionals in the early and accurate detection of brain tumors.

#### C. ROC AND AUC CURVE

The Receiver Operating Characteristic (ROC) curve and the Area Under the Curve (AUC) were used to evaluate the brain tumor detection model's performance by analyzing the trade-off between the true positive rate (Sensitivity) and the false positive rate (1 - Specificity) across different classification thresholds. The ROC curve provides a visual representation of the model's ability to differentiate between tumor and non-tumor cases, with a steeper curve indicating a stronger model. The AUC, which measures the overall discriminatory capability of the model, serves as a comprehensive metric of its performance. An AUC value closer to 1 signifies a highly effective model, with excellent class separation. In this case, the model achieved an AUC of 0.98, demonstrating its outstanding accuracy and reliability in detecting brain tumors, making it a valuable tool for medical diagnostics.

```
# ROC Curve and AUC
fpr, tpr, _ = roc_curve(y_true, y_pred[:, 1])
roc_auc = auc(fpr, tpr)

plt.figure()
plt.plot(fpr, tpr, color='blue', lw=2, label=f'ROC Curve (AUC = {roc_auc:.2f})')
plt.plot([0, 1], [0, 1], color='gray', lw=2, linestyle='--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver Operating Characteristic (ROC) Curve')
plt.legend(loc='lower right')
plt.show()
```

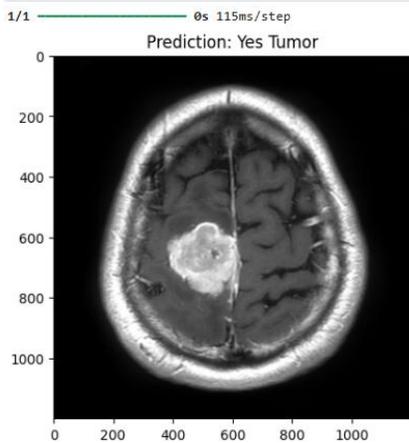


ROC AND AUC

D. PREDICTIONS

The below figure represents the model's prediction when an MRI image with a brain tumor is provided. The system successfully detects the presence of the tumor and classifies the image as "Yes." This result highlights the model's ability to accurately identify brain tumors using its trained convolutional neural network (CNN). The visualization demonstrates the model's effective decision-making process, contributing to reliable and timely medical diagnoses.

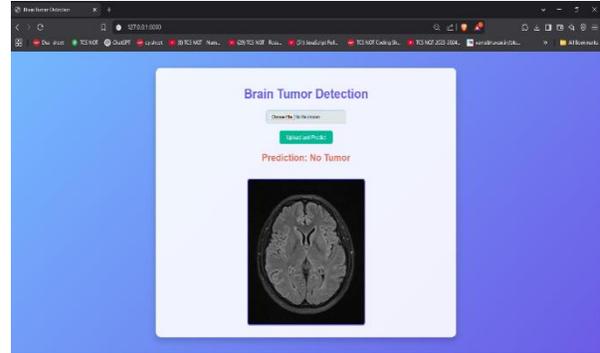
```
# Provide the path to a sample image for testing
sample_image_file = '/kaggle/input/brain-tum-image/dataset-card.jpeg'
predict_tumor(sample_image_file)
```



PREDICTION

The below figure represents the model's prediction when an MRI image without a brain tumor is provided. The system accurately classifies the image as "No,"

indicating the absence of a tumor. This result demonstrates the model's capability to correctly differentiate between normal and abnormal brain scans, ensuring reliable predictions and supporting medical professionals in making accurate diagnostic decisions.



PREDICTION AFTER INTEGRATING MODEL WITH FRONTEND

3. CONCLUSIONS

Brain tumor detection remains a critical challenge in the medical field, requiring accurate and timely diagnosis to improve patient outcomes. This project utilized convolutional neural networks (CNNs) to develop an automated brain tumor detection system capable of analyzing MRI scans with high precision. A comprehensive dataset of brain MRI images was employed to train and validate the model, ensuring robust performance. Various evaluation metrics, including accuracy, precision, recall, F1-score, and AUC, were used to assess the model's effectiveness. Through this analysis, the model demonstrated exceptional accuracy in distinguishing between tumor and non-tumor images, proving its reliability in medical diagnostics.

To ensure practical usability, a user-friendly frontend was developed using Django. The frontend enables users to upload MRI images and view the prediction results in real-time. The seamless integration of the frontend with the backend model ensures an intuitive experience, making the system accessible for non-technical users. This automated solution offers a valuable tool for assisting healthcare professionals, reducing diagnostic time, and minimizing the risk of human error. The results underline the potential of deep learning technologies in enhancing early tumor detection and contributing to improved patient care.

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