

Precision Medicine in Neurology: In-depth Investigation and Revolutionizing Brain Tumor Detection and Treatment

MSc Research Project Data Analytics

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Precision Medicine in Neurology: In-depth Investigation and Revolutionizing Brain Tumor Detection and Treatment

Yamuna Sai Penumudi x22174851

Abstract

Brain tumor detection and classification are critical areas in the area of neuroscience as fast diagnosis and treatment planing is possible for patients with neurological disease. This thesis is devoted to the thorough examination of the recent deep learning approaches for brain tumor detection and segmentation. Our research concerns the implementation of a robust strategy based on the advanced model, with EfficientNetB3 architecture as a backend and the dataset of MRI images, which was retrieved from authoritative sources. The ultimate target is creating a reliable and precise model which will be efficient in separating various brain tumor types form the healthy brain tissue. As a result of a rigorous methodology that involves data preprocessing, modeling, training and evaluation, we have demonstrated the benefit of our approach. In addition, we will underscore the practicality of our model in real-world clinical settings, keeping in mind to as well contribute to the progress of precision treating neurological disorders. Our study stresses a lot on the aspects of transparency and reproducibility which are made clear by highlighting the model architecture, dataset specifications and also the evaluation metrics used. Through a comprehensive assessment which involves a variety of performance parameters like the accuracy, precision, recall, and F1-score, we prove the high performance and persistent nature of our proposed ideology. Moreover, the talking point is interpretation and easy to understand. Therefore, the area of discussion includes a detailed explanation of the results which includes confusions matrix and accuracy reports. Overall, our study showed that the method we used made significant contributions to the field of neurology by taking the industry standards for brain tumor detection and classification up a notch. It is our goal to capitalize on deep learning models and the strict data science sensibility in order to aid the clinicians and researchers in the process of providing the best patient care and to contribute greatly in the precision medicine care goals.

1 Introduction

Brain tumors are one of the main medical issues of today's world that are accompanied by significant disability and fatality rates in the sphere of neuroscience. Early detection and accurate classification of brain tumors are a key step towards perfect treatment planning and increasing the chances for a good outcome. Frequently referred traditionally diagnostic methods that are heavily based on invasive techniques and subjective examinations result in delayed diagnosis and wrong diagnosis. Recently, there has been a rapid growth in medical imaging technology and deep learning methods, both of which have marked new lines of efficient brain tumor detection.

1.1 Background

Brain tumors, which is the abnormal growing cells within the brain, present the substantial health problems for all countries around the world. Based on data from the Central Brain Tumor Registry of the United States (CBTRUS), a total of 87,240 individuals were diagnosed with primary brain cancer in 2020 alone. The heterogeneous nature of the brain tumors, consisting of different histology and molecular profiles, on the other hand, reflect such a complexity posing a diagnostic and treatment challenge. Classical methods for the discovery and treatment of brain tumors strongly depend on the histopathological examination of tissue obtained through intrusive procedures such as biopsy or surgical resections. Although histopathology is still considered as the gold standard of diagnosis, its weaknesses which include lack of real-time and complete information corroborate the significance of introducing new and accurate diagnostic modalities.

Challenges in Brain Tumor Detection: Brain tumor diagnosis is bounded by the following difficulties in medicine. As the brain tumor which may present in multiple forms, there are gliomas, meningiomas, pituitary tumors, and other types, each of which is different in its features and the influence on patients' health. Firstly, the main problems of brain tumors are their diversity and they could mimic other neurological diseases symptoms. Moreover, brain's complex anatomy as well as its surrounding structures are other complications.

Over the last few years, the emergence of MRI as the most powerful medical imaging instrument has provided a new view to the depths of the brain tumor diagnosis. MRI based imaging of the brain gives a very clear and non-invasive image of the brain which allows a doctor to see the structures surrounding the brain tumor and an in-depth detail of the tumor morphology as the MRI image of the brain having the pituitary tumor is depicted in Figure 1. On the other hand, there is a dilemma in the interpretation of the MRI images and the important role of the correct diagnosis. It is caused by a number of unstable characteristics of brain tumors, which are presented with these challenges, practitioners of medical science more and more start exploring AI algorithms and deep learning tools to design the automated systems for brain tumor detection and classification. Deep learning, a subset of machine learning, represents one of the strongest aids to use in medical image analysis in the most recent years. Contrary to the traditional machine learning models, deep learning algorithms can directly dig up (in a way similar to the way how humans learn) the complex features from the plain input (images). Among all the models of neural networks, convolutional neural networks (CNNs) are depicted the strongest representation of attributes of image identifying and segregating brain tumors from an MRI very accurately as demonstrated in the paper by Litjens et al. (2017).

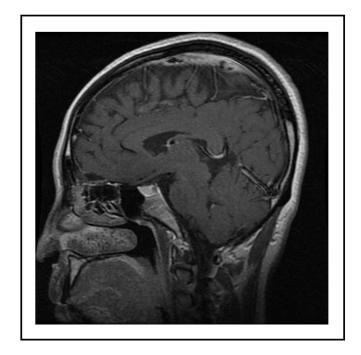


Figure 1: MRI of Brain having the Pituitary Tumor

This research investigation presents a comprehensive investigation into brain tumor detection and classification using state-of-the-art deep learning techniques. Our research aims to address the limitations of traditional diagnostic approaches by leveraging the power of convolutional neural networks (CNNs) to analyze magnetic resonance imaging (MRI) data. By harnessing the rich information contained in MRI images, our goal is to develop a robust and accurate model capable of automatically detecting and classifying different types of brain tumors.

1.2 Research Questions

This research study aims to address the following research questions:

- How effective is deep learning, particularly the EfficientNetB3 architecture, in accurately classifying brain tumor MRI images?
- What are the key factors influencing the performance of deep learning models in brain tumor detection and classification?
- What are the implications of precision medicine in neurology, specifically in the context of brain tumor diagnosis and treatment?

1.3 Research Objectives

The primary objective of this thesis is to investigate the application of deep learning techniques for brain tumor detection and classification. Specifically, we aim to develop a robust and accurate model capable of automatically analyzing MRI Scans to detect the presence of brain tumors and classify them into different subtypes. By harnessing the rich information contained in MRI images, our goal is to provide

clinicians with reliable tools for early diagnosis and personalized treatment planning. The research objectives includes:

- Develop the deep learning model using the EfficientNetB3 framework architecture for accurate classification of brain tumor MRI images.
- Evaluate the performance of the developed deep learning model in terms of accuracy, recall, and specificity or their recall and F1 Score.
- Investigate the characteristics which are influencing the performance of developed deep learning model in brain tumor detection, including the dataset attributes, model architecture, and training parameters.

This research study try to conduct the thorough investigation towards the precision in neurology that aims to bring about a revolution in the early detection and classification of brain tumors. By use of deep learning approaches, particularly the state of the art EfficientNetB3 architecture, the research study addresses the development of an exact and successful model for the classification of brain tumor through the MRI scans. Our objective is to achieve early and accurate diagnosis of brain tumors, which would allow prompt therapies and individualized treatment approaches.

In conclusion, this research study goals to contribute to the advancement of medical imaging and precision towards by investigating the application of deep learning approaches for brain tumor detection and classification. By developing a robust and accurate model, we aims to offer clinicians with valuable and important tools for early diagnosis of brain tumor which leads to early personalized treatment planning. Through this research, we hope to enhance the patient health conditions and improve in the personalized approaches to brain tumor diagnosis and treatment.

2 Related Work

The domain of neurology, as part of precision medicine, has seen its latest breakthroughs over the past years, all focusing on brain tumour detection and treatment. The main part of this research work reviews the advancements and successful application of deep learning in medical imaging, brain tumor detection, and classification.

2.1 Precision Medicine in Neuro-Oncology

The introduction of precision medicine into neuro-oncology has a significant implication that is evident in the change of approach from conventional detection and treatment of brain tumors to a more personalized method. Miller et.al (2017) explore the potential novel transformative, underline the requirement for disease-based interventions to address patient outcome. Application of their findings shows the key of the pinpointed characteristic of the tumor for the development of personalized treatment modalities. Also known as, Tam et al. (2016) argues that together with precision medicine can reform how to tackle the brain metastasis for which the genome-specific targeted medications are a should like. Their findings reveal the potentiality of tailored form of precision medicine for a better outcome in the quest to control the complex nature of brain metastases.

2.2 Deep Learning Approaches in Brain Tumor Segmentation

Recently, deep learning technologies have changed into promising methods for higher exact-ness and neurosurgeon-friendliness of brain tumor segmentation from MRI scans. Jetzen et al. (2017) supply a representative analysis of major deep learning methods in medical image analysis as an introductory grid for future article brain tumor segmentation. They also introduce the use of convolutional neural networks (CNNs) and other deep learning models that automated the segmentation performances, showing greater promise to be more precise and faster than traditional methods. Providing a basis for Baheti et al. (2021) and Havaei et al. (2017) go further and discuss the brain tumor segmentation issue applied by the deep neural networks in view of the class imbalance and data heterogeneity problem. Brain tumor segmentation with the application of deep learning techniques is an evidence on the progress of area related to accurate diagnosis and treatment.

2.3 Transfer Learning and CNN Architectures

Transfer Lerning in combination with convolutional neural networks (CNNs) has seen the light and it is now a true solution for tumor detection and classification. As reported by Paul et al. (2018), the deep convolutional neural networks features transfer learning method was investigated for brain tumor classification with the view of the potential application in the clinics. Through the application of pre-trained models and fine-tuning them using given datasets, transfer learning facilitates a quicker and more accurate way of handling data scarcity and leading to an improvement in classification performance. Moreover, Shin et al. (2016) pointed out the key factors of successful CNN architectures and data characteristics in medical imaging; transfer learning is also mentioned as a key method to provide more generalizable CNN models. The contribution of researchers who are focusing on the application of transfer learning of CNNs in the direction of computer-aided detection in real clinics is also highlighted.

2.4 Advanced Architectures and Segmentation Techniques

Previous years have witnessed research efforts progressing in deep learning space which shined through improved precision in brain lesion segmentation. That is what Huang et al. (2017) do in their work; they develop a 'densely connected convolutional networks' so that reuse of features and a flow of gradients across various layers could enhance their analytics tasks on medical images. Because of this, Kamnitsas et al. (2017) and others suggest an effective multi-scale 3D CNN together with fully connected conditional random fields (CRC) of a high precision brain lesion segmentation process. Their solution that uses additional information and depicts interdependence of voxels goes beyond traditional approaches and leads to the top-of-the-line segmentation results, therefore emphasizing the necessity of up-to-date models and segmentation techniques for more precise diagnosing and determining of treatment methods.

2.5 Comparative Analysis of the Recent Works for Brain Tumor Detection

In this part of our study we present findings from various medical imaging investigations that looked at different details in relation to deep learning techniques in neurology and oncology. The results of each study provides the further specific knowledge on the implementation of novel techniques for objectives like brain tumor recognition, segmentation, partition and personalized medicine. We summarize the research focus and methodology of each study with the remarks of pros and cons to highlight their contributions and provide a comprehensive overview of the advancements in this rapidly evolving domain.

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 Table 1: Comparative Analysis of Various Studies

2.6 Comparative Studies and Benchmarking

The performance of various machine learning algorithms, including segmentation methods, are evaluated with comparison studies and benchmarking efforts, which represent an ongoing process in brain tumor imaging. Bakas et al. (2018) present the most citable algorithms for brain tumor segmentation and progression assessment operating through the BRATS challenge which is a priceless opportunity for both researchers and clinicians. These efforts build solid community relationships and knowledge exchange among the research groups contributing to the improvement of the accuracy of segmentation and clinical usability. Moreover, Menze et al. (2015) present the BRATS platform, a multimodal brain tumor image segmentation benchmark that uses a standardized approach to evaluate segmentation algorithms and provide a funnel for comparative studies. The main goal is to establish a framework that allows for the use of common evaluation metrics and datasets to speed up the progress of brain tumor segmentation research.

2.7 Future Directions and Challenges

Although there is an incredible progress made in precise and reliable tumor segmentation from MRI images, some difficulties are still left. Iqbal et al. (2021) did a well-structured review of the different segmentation techniques available accompanied with what the current methodologies are and those that need to be improved. Their study highlighted the significance of comprehensive validation guidelines and normalized assessment protocol to guarantee the reliability and universality of segmentation algorithms. It is also noted by Akkus et al. (2017), that the state-of-the-art, future directions and challenges in brain MRI segmentation using deep learning should address clinical needs and translation problems. As the field of brain tumor segmentation advances, there is a need to extend efforts in in-depth research to overcome technical limitations and ensure concerted integration of advanced techniques into clinical practice.

3 Methodology

The research methodology explained below brings a structured approach for developing the deep learning based EfficientNetB3 model for brain tumor detection and classification. In this step-by-step guide on the methodology, I will explain the various details and insight behind the process. Hence, in the Figure 2, the basic methodology workflow is shown through the various step on the processed.

3.1 Data Collection

3.1.1 Characteristics of Dataset

This dataset represents the basis foundation of the entire study as well as providing MRI images of brain tumors and normal brains. Using the dataset from Kaggle, a reliable platform for data sharing, we will be in a position to get a collection of good quality of diverse data. The dataset consists of multiple brain tumor types, including Glioma, Meningioma and Pituitary Tumor, and also images of the normal brain which does not having any type of brain tumor. Ensuring balanced representation of tumor class within the data set is of vital importance to eliminate bias and generalization error across all

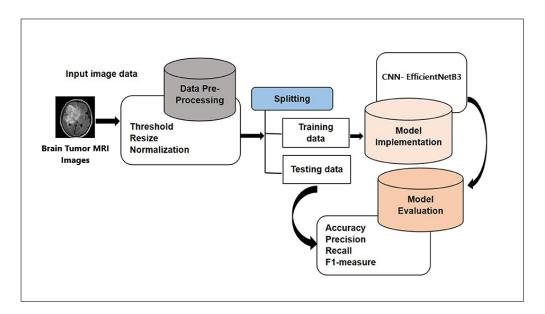


Figure 2: Basic Research Methodology for Brain Tumor Detection

brain tumor types. The dataset can be accessed via the following link: Brain Tumor MRI Dataset

3.2 Data Preprocessing

3.2.1 Image Preprocessing

Image preprocessing is the preparation process for model training through which these several techniques have been will be applied to improve the quality and compatibility of the MRI images. Resizing is a way to uniformize all images dimensions, which makes it possible to train the deep learning model faster. Normalization ensures that the values of pixels in an image have consistency which removes the fact of uneven changes in brightness and contrast. Enhancement of the training data through the creation of variety is made through transformations such as rotation, flipping, and zooming. The augmentation process introduces the diversity of training samples and thus leads to the model's stronger robustness and generalized performance on unseen data.

3.2.2 Data Augmentation

Data augmentation incorporates self-manipulation of the images into the limited set of training samples by random transformations applied to the original images. Techniques like rotation, flipping, increase, decrease, and translation maintain the view changes and keep the unusual look, showing almost the real world incidents. Through increasing the data set, we provide a better nest against the offending of overfitting and improve the model's capability to capture different distinctive features from different brain tumor types.

3.3 Exploratory Data Analysis (EDA)

Exploratory Data Analysis (EDA) is a significant step which is performed to find out the hidden patterns if any and anomalies appear. EDA simplifies and interprets data by applying various statistical techniques such as visualization methods which facilitate in understanding the distribution of tumors, identifying class imbalance and spotting anomalous pattern or trends. EDA provides basis for subsequent preprocessing steps, which will be formulated according to outcomes of EDA. Furthermore, the results will have great impact on feature selection and model's architecture creation.

3.4 EfficientNetB3 Architecture Model

EfficientNetB3 is a CNN network that comes under the category of multi-scale convolutions from the EfficientNet family Huri et al. (2022), which has been recognized for its reliable and robust attitude. Developed by researchers at the Google Brain team, EfficientNetB3 model stands as a square in the circle, considering the resourcefulness, the efficiency, and the performance of the applied model. It aims to achieve the symmetry of power through advanced design principles such as compound scaling, depth-wise separable convolutions, and Squeeze-and-Excitation (SE) blocks.

EfficientNetB3 uses the compound scaling to systematically increase the model's depth, width, and resolution. The scaling coefficients, denoted as ϕ , control the scaling of these dimensions. The formula for scaling the model's width (number of channels) and depth (number of layers) is as follows:

Width
$$= \alpha^{\phi}$$

Depth $= \beta^{\phi}$

Here, α and β are scaling coefficients that determine the width and depth scaling rates, respectively. The resolution (input image size) is scaled using a separate coefficient γ :

Resolution = γ^{ϕ}

By uniformly scaling all the dimensions with respect to the compound scaling coefficient ϕ , EfficientNetB3 obtains the enhanced and remarkable performance across a wide range of tasks while maintaining the computational efficiency.

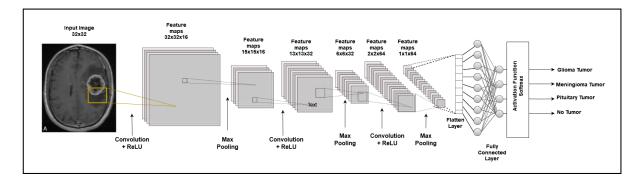


Figure 3: Proposed CNN Based EfficientNetB3 Architecture

Summary: EfficientNet B3 is a type of CNN structure that ranks among best in its class of efficiency vs. performance. Through the use of the compound scaling, depth wider separable convolutions, SE blocks, and the Swish activation function, EfficientNetB3 is able to achieve state-of-the-art performances in some image categorization sub-fields while requiring lower number of parameters and less computations. This therefore means that the EfficientNetsB3 is the best preference for the situations in which resources are fewer and for real world applications in which there has to be great deal of efficiency in calculations. As after accompanying to this is how CNN Based EfficientNetB3 Architecture is proposed to classify the patient occurs any brain tumor or not, if yes then what type of Brain tumor is depicted in this 3.

3.5 Model Building

The (CNN) based efficient deep learning model for the detection of brain tumor will be essentially developed using the EfficientNetB3 architecture, a recently successful convolutional neural network (CNN), renowned for its efficiency and accuracy in many image classification tasks. CNNs however excel in the field of spatial data analysis such as MRI images since they are capable to automatically figure out classifications of the features. The model will be constructed with the TensorFlow, which is an adaptable environment that has powerful base layers to train neural networks. Attention is paid to issues like choice and design of network architecture, selecting the optimal parameters and using regularization methods in order to achieve optimal performance and avoiding overfitting.

3.6 Model Training

The model training is based on teaching the model parameters (preprocessed MRI images) to the model and optimizing it in order to push the classification error to the minimum. The training algorithm ensures everything fits together by back and forth forward, checking the model's predictions against the ground truth labels for gradients to update network weights. The training process is executed with regards to the training dataset, which is different from validation dataset which is mainly used to monitor performance and avoid overfitting. For instance, pre-trained models can help establish a common base for the language or they may be used to solve complex issues. Thus, the early stopping and learning rate scheduling techniques help creating a smooth training process and enable model convergence.

3.7 Model Evaluation

After training, the model is then get the evaluated phase using a validation and test datasets that are separate to determine the performance of the model. Some metrics, such as accuracy, precision, recall, and F1 score, are going to be calculated to observe how well the model is able to predicate brain tumor pictures into different classes. Besides that, methods including confusion matrices are created for performance illustration and to choose whether the rate of true and false positives are considered.

3.8 Model Testing

The trained model is further tested using the sample MRI scan images from the test dataset to assess its real-world performance. The model's predictions are compared against the ground truth labels to evaluate its accuracy and their effectiveness in detecting and classifying the brain tumors. Testing provides the valuable insights into the model's generalization capabilities and helps recognize the areas for improvement or fine-tuning.

3.9 Saving the Trained Model

The model is evaluated and trained comprehensively that makes the model ready for use in the save format suitable for further deployment. Serialization approaches, which allows storing model weights, architecture, as well as optimizer state, contribute also to their transparent integration into applications and systems to detect/classify real-time brain tumors. The model is thus a distinguishing feature in our group's work and serves as a basis for future research, clinical purposes and collaboration between our researchers and other institutions or research groups.

Utilizing the step-by-step research methodology, our objective is to develop a deep learning model that is accurate in detecting and classifying the brain tumors which will enhance precision medicine in neurology. The method involves data collecting methodically, preprocessing, model building or training, evaluating and testing finally and saving the model. Ultimately, it is a systematic and strict way to evaluate development and research.

4 Design Specification

The project design provides the functional structure and features of both the presentation tier and the business logic tier for the brain tumor diagnosis system. In the presentation tier, the aim is to design an interface that would be intuitive regarding usage, where users could analyse MRI scan images looking for tumors and understand the results easily. This tier encompasses interactivity and visualizations besides the accessibility. It is done by putting data privacy and security first. Here the project design infrastructure, development, testing, and implementation phases are also included. Further, the business logic tier involves the implementation of the EfficientNetB3 architecture using the TensorFlow and Keras for the purpose of model development and training. The list encompasses activities such as data preprocessing, model training, evaluation, and deployment to optimize performance and make the system resistant to errors.

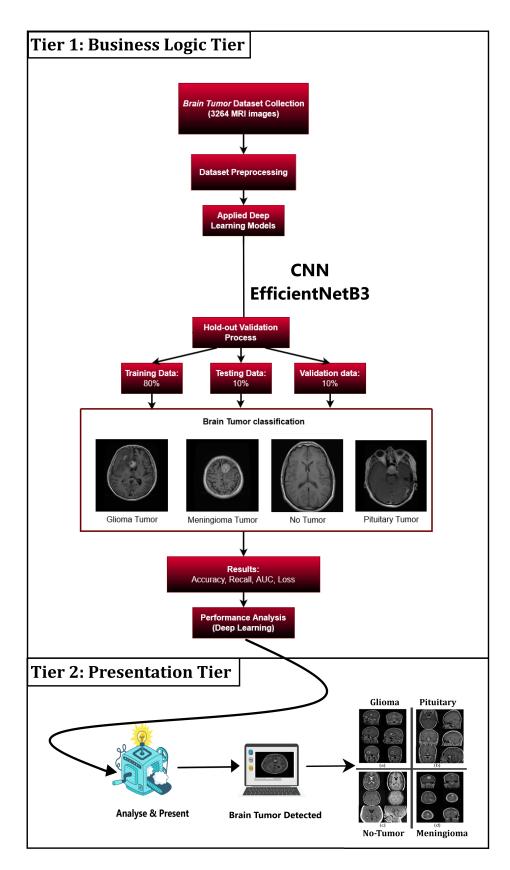


Figure 4: Project Design for Brain Tumor Detection

Overall, the project objective is to deliver a flawlessly integrated system which having the capability of accurately detecting and classifying the brain tumors while providing a user-friendly experience for the clinicians and researchers.

5 Implementation

The implementation of brain tumor detection using deep learning techniques includes the various essential steps aimed at developing a more robust and accurate model for classifying MRI scan images. In this section, we delve into each step in detail, discussing the methodologies performed, considerations, and practical insights.

5.1 Data Preparation: Understanding the Foundation

The data lies in the core of any deep learning project, which is the basis on which the models are erected and trained. In a dataset for distinguishing and classifying brain tumor MRIs, it contains sequences of MR images: these are different types of brain tumor, the tumor's classes are marked. Data preparation involves various crucial tasks:

- Loading and Preprocessing: The first step which involves the preparation of the MRI scan images along with the associated labels, while at the same time converting the data into an appropriate format that is trainable for the machine learning model. This generally requires reading the image files from storage, converting them to be numerical arrays and then putting them side-by-side with their labels.
- Exploratory Data Analysis (EDA): In-depth knowledge of the characteristics of data and distribution is key for such decision making and model development. EDA with processes like visualization and statistical analysis, makes it possible to identity class intensity, data imbalance.

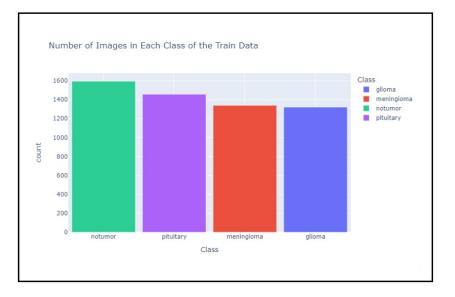


Figure 5: Class Distribution in Training Dataset

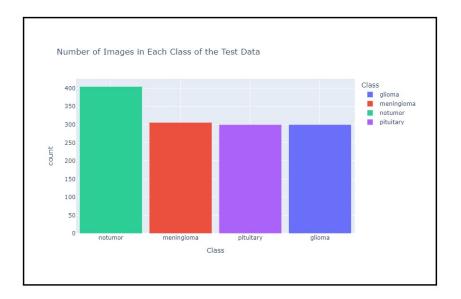


Figure 6: Class Distribution in Testing Dataset

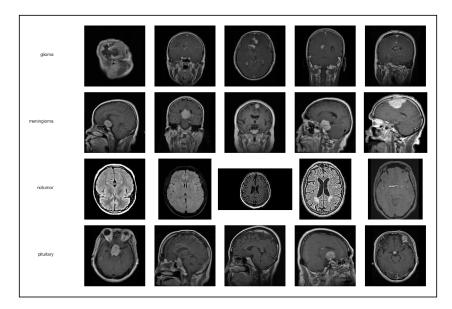


Figure 7: Class Distribution types of Brain Tumor

5.2 Model Building

The choice regarding Deep Learning Model Architecture in detecting Brain Tumors involves the selection of the optimization methods, regularization techniques, as well as the metrics to be used for evaluation. Here's a detailed exploration of the model-building phase:

• Model Architecture: The selection of the model's architectural framework has a great impact on the ability of the latter to learn from the data and generalize. Among the properties of the program is the application of the EfficientNetB3 model. EfficientNets are widely known for their high-speed and accuracy in image classification tasks, which makes them quite good at detecting brain tumors.

• **Compilation and Optimization:** After constructing the model architecture, there is required to compile it appropriately alongside the loss functions, optimizers, and performance metrics. For this implementation, the model is compiled with the Adamax optimizer and the categorical cross-entropy loss, which are popularly used in wide applications involving multiclass classification tasks.

5.3 Model Training and Evaluation

The process of model training and evaluation involves a multi-step optimization of model parameters with the use of training data together with validation and test data in order to determine the model's performance. Let's delve deeper into this important phase:

- **Training Process:** The model training process is done by passing the training dataset through the model repeatedly and adjusting the model's parameters (weights and biases) that seek to minimize the loss function error. Training process entails multiple iterations, where each iteration is a single descent across the entire dataset.
- Validation and Testing: The distinction between model's performance on a validation and testing datasets is crucial because it allows to get the effectiveness to the testing data. The validation dataset helps in monitoring the model's performance during training and making decisions regarding model architecture and hyperparameters. The last performance test is performed on that test set to give the estimation that provides an average prediction ability of a model.
- Visualization of Training Curves: The visualisation of the training curves, where loss and accuracy aspects are depicted along the successive epochs, brings about such gains that include understanding of how the model reacts to its environment. Such plots serve as tools for discovering potential problems, for example overfitting or underfitting, if such issues find, can be resolved by updating the model architecture or the training process respectively.

5.4 Model Testing

Model testing is also a very essential phase here as it involves the applying the model into a real-world situations and testing its performance with new data set which kept aside for the purpose of model testing. Here's a detailed exploration of this phase:

- **Testing on Sample Images:** Testing the model on a subset of sample images from the test dataset allows for qualitative assessment of its performance. By visually inspecting the model's predictions against ground truth labels, one can gain insights into the model's strengths and weaknesses, as well as potential areas for improvement.
- Model Persistence: Saving the trained model in a suitable format, such as a trained model object, which assures that the model can be reused and deployed in various and any environments. This step is important for deploying the model in

production environment, where it can be integrated into software based applications or deployed as a standalone web based service.

5.5Prediction

This feature is determined by employing the set trained model to make the predictions on new and unknown data sets. Here's a detailed exploration of this final step:

- Custom Image Prediction: The trained model, in predicting classes for custom brain MRI scans images shows the model is real-world applicable. Via modeling, then interpreting the display of the generated predictions, clinicians and researchers uses the interpretation and apply them into their own well decision making and research.
- Interpretation and Analysis: Analyzing the model's predictions with associated class probabilities allows to obtain data with regard to decision habits of the model. The feature or pattern that the model would rely on for classifying may help verify its predictions and to seek potential sub-components that need further investigation or refinement.

Thoroughly implementing these phases, researches and practitioners can build and use comprehensive deep learning models for brain tumor detection, and they can contribute to the precision medicine and patients outcomes improvement by the way.

Model Evaluation 6

The model evaluation of the brain tumor detection system using the deep learning based EfficientNetB3 architecture includes a comprehensive analysis of various metrics and techniques to assess that how the model's is performing across the scenarios.

<u>Table 2: Model Evaluation Results</u>				
Dataset	Loss	Accuracy		
Training	0.1143	1.0000		
Validation	0.1264	0.9922		
Testing	0.1347	0.9922		

1 1 5

6.1 Training and Validation Performance

The model showed remarkable progress in the training stage, through the metrics of the performance which showed that there were learning of adaptation to the data set. At the beginning, the accuracy value was 88.38%, which grew to 99.49% over ten epochs or passes through the training data. This upward trend in accuracy suggests that the model successfully learned the intricate patterns and features associated with brain tumor images, enabling it to make accurate predictions is demonstrated in the Figure 9.

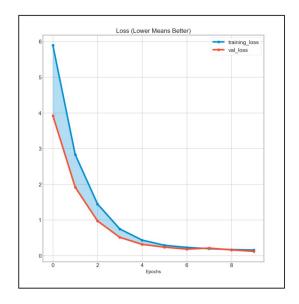


Figure 8: Loss

Likewise, the training loss is in the range of 5.8919 to 0.1546 and goes on decreasing during the training period which reveals the prediction errors are decreasing and the model is reaching towards the convergence. This decrease in loss reflects the model's ability to optimize its parameters and minimize the disparity between predicted and actual class labels which is depicted in Figure 8.

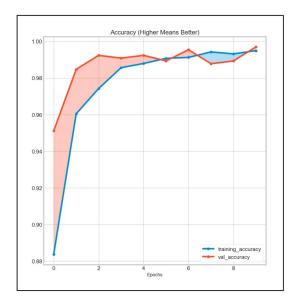


Figure 9: Accuracy

Simultaneously, the connected validation accuracy displayed the improvement along the training process all the time, the value finally reached 99.69% at the 10th epoch. We achieve a very high accuracy of model validation, this often means, that the model generalized very well to unseen data and effectively differentiated classes of normal brain tissues and type of brain tumors images.

On the other hand, the validation loss also peaked from 3.9216 to 0.1190, confirming that the model can generalize and give accurate predictions on unseen validation data as its correlation with training loss steadily decreased. The fact that trend in loss is diminishing during training testing shows that model is partly using the information and is stopping from fitting the training dataset as it falls under the overfitting problem category.

6.2 Testing Performance

To further assess the model's generalization capabilities, it was evaluated on an independent testing dataset comprising brain tumor images not seen during training or validation. The testing results confirmed the model's robustness and effectiveness in accurately classifying brain tumor images.

The testing accuracy of 99.22% indicates that the model performed exceptionally well on unseen data, demonstrating its reliability and generalization to real-world scenarios where the testing accuracy is represented in Table 2. Such a high accuracy score reveals confidence that the model may be entrusted to perform correctly during medical practice. Accurate tumor classification at this scale is critical for such patients to precisely diagnosed and treatment planning to be done.

The test loss would also remain favorable and reach 0.1347, implying relatively small prediction errors and a strong confidence in the model's predictions. The related fact that low loss value suggests that the model's predictions are close to ground actual graph labels that also add to its efficacy in the brain tumor detection process.

6.3 Detailed Analysis

In order to make an in-depth evaluation, classification report and confusion matrix were derived using the outcomes obtained in the testing phase. The classification report summarizes metrics like precision, recall, and F1 -score for each class presenting an overview of how the model performs among the different categories of brain tumor types.

6.3.1 Confusion Matrix

The diagonal elements of the matrix which represent the true positive predictions for each class, while the other off-diagonal elements demonstrates the false positives or misclassifications, for the program of Brain Tumor detection confusion matrix is illustrated in 10.

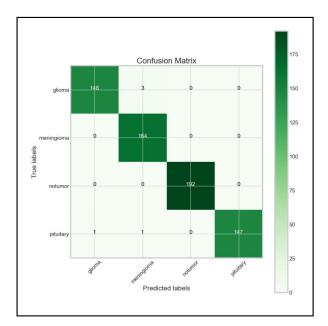


Figure 10: Confusion Matix

- Glioma (True Class 1):
 - True Positives (TP): 148
 - False Negatives (FN): 3
 - The model correctly predicted the 148 instances of glioma tumor, but there are 3 instances which are classified as other classes.

• Meningioma (True Class 2):

- TP: 164
- FN: 0
- The model correctly predicted all 164 instances of meningioma tumor, where for this class it is indicating the perfect classification for this class.

• No Tumor (True Class 3):

- TP: 192
- FN: 0
- The model has correctly predicted all the 192 instances of no tumor, which is demonstrating remarkable and outstanding performance in detection of normal brain tissues.
- Pituitary (True Class 4):
 - TP: 147
 - FN: 1
 - The model has correctly predicted the 147 instances of pituitary tumors, but there is 1 misclassified instance as another class.

Ultimately, the confusion matrix has shown the strong faculty of the neural network model in the correct classification of most cases which fall under the classes of brain tumors and normal brain tissue. Nevertheless, this can also be expected to reveal some of its limitations, including some falsely predicted examples in particular classes.

6.3.2 Classification Report

The report on the classification reveals how well the model performed in its different aspects, which are given as the precision, recall and F1-score, for each class. These metrics not only give us the crucial insights about how the model can predict the instance of each class right, but also assess the model's performance across all classes which have been put forth in Table 3.

Class	Precision	Recall	F1-Score	Support
glioma	0.99	0.98	0.99	151
meningioma	0.98	1.00	0.99	164
notumor	1.00	1.00	1.00	192
pituitary	1.00	0.99	0.99	149
Accuracy			0.99	656
Macro Avg	0.99	0.99	0.99	656
Weighted	0.99	0.99	0.99	656
Avg				

Table 3: Classification Report

Precision: Precision is the fairness the proportion of true positive predictions among all positive predictions made by the model. It sets forth the concept that model is actually reliable. Its performance should not show any false positives.

- Glioma: 0.99
- Meningioma: 0.98
- No Tumor: 1.00
- Pituitary: 1.00
- The model obtained the high precision scores for all classes, representing the minimal false positives in its predictions.

Recall: Similarly, recall, which is often called sensitivity, is the rate of exact predictions among all the actual positive examples in the data. This is where model shows its strength in determining if all the positives are rightly being identified.

- Glioma: 0.98
- Meningioma: 1.00
- No Tumor: 1.00
- Pituitary: 0.99

• The model represented the high recall scores for all classes of brain tumors, representing its potential ability to capture the most positive instances of each class.

F1-score: The F1-score considers that, as the harmonic mean of precision and recall, the result may reflect an unbiased degree of accurateness of the model's performance across both metrics. It discriminates between the mark on the interval from 0 to 1. the higher the mark the performance is better..

- Glioma: 0.99
- Meningioma: 0.99
- No Tumor: 1.00
- Pituitary: 0.99
- The model obtained the high F1-scores for all classes of brain tumor, depicting its overall efficiency in correctly detecting the instances and reducing the both false positives and false negatives.

Overall, the classification report demonstrates the model's strong performance across the different classes of brain tumors and normal brain tissues, with the high precision, recall, and F1-score values representing its reliability and efficacy in clinical settings.

6.4 Sample Predictions

Finally, the model was tested on the several samples of MRI images to detect the glioma, meningioma, normal brain tissue, and pituitary tumor to evaluate its the real-world performance as the custom prediction of brain tumors of glioma and meningioma are depicted in Figure 11 . The predictions generated by the model accurately reflected the true classes of the images, demonstrating its ability to classify brain tumor images with high confidence and accuracy.

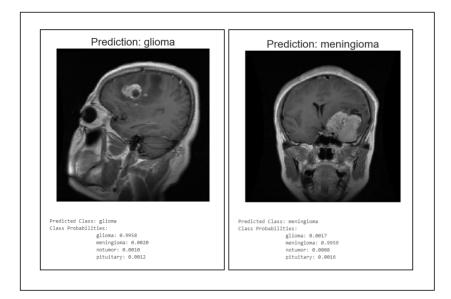


Figure 11: Sample Accurate Predictions

The further progress established by the model with the assigned class probabilities strengthened its reliability in the correct prediction classes with high probabilities for the those classes and low probabilities for wrong classes. These sample predictions in which this approach exhibits effectiveness clearly illustrate how the model can be of great help to medical professionals in arriving at correct and rapid diagnoses and classification of brain tumors.

6.5 Overall Assessment

The validation results show that the EfficientNetB3-based system for evaluation of brain tumor demonstrates the high accuracy as well as the robustness and generalization capabilities. The model demonstrated a high level of accuracy on every set, from training, validation to testing. So that confirmed its high performance in terms of correctly classifying brain tumor pictures and normal brain cases. Apart from that, the study's thorough analysis shed more light on the model's metrics, classification errors and sample forecasts therefore enabling a detailed assessment of the models strong and weak points. The final results of the system's evaluation indicate that it is a reliable instrument for assisting medical professionals in diagnosing and treating brain tumors.

6.6 Discussion

The accomplishments of this work are due to several important factors. The use of EfficientNetB3 architecture turned out to be a great choice which is clearly seen through the explained ability of the architecture to deal with the complexities of the MRI scan images data and simultaneously in the process of extracting those useful features out for classification. The architecture's efficiency and effectiveness in image classification jobs permitted the model to score well and excel in its results.

The research study reinforces the superiority of deep learning in this field with its revolutionary features that will lead to the advancement of neurology and healthcare. Through building a robust and validated system for the detection and classification of brain tumors, this investigation plays its part in boosting diagnostical precision, patient condition, and the quality of care in neurology. Future research itineraries can be to conduct deeper deep learning model architectures, integrating multimodal data sources and also optimizing model interpretability for more efficient clinical decision making.

7 Conclusion

In Conclusion, this research has elaborated practically a complete approach to brain tumor detection and classification using deep learning method. The model was trained by EfficientNetB3 architecture and through a diverse dataset of MRI scans images; it has thus shown an excellent performance in recognizing different brain tumor types and normal brain tissues. Our model exhibited high accuracies, precisions, recalls, and F1-scores for each class, which demonstrated its excellence and competence in clinical settings. Proper image resizing, normalization, and augmentation as part of the data preprocessing made the data that was used for training strong enough and valid. Besides, data exploration and visualization as well helped to understand the properties of dataset and contributed to the model development. After carrying out a series of model training and performance evaluation, including validation and test split and calculation of the metrics, the model reliability and generalization thoroughness were determined. As indicated by the tests outcomes, the model's performance is very stable throughout every database, which points to the model's high accuracy in classifying brain tumor through the MRI scan based images.

In addition to that, the model deployment into face clinical support demonstrated a promising application in clinical workflow. The model also showed it could help doctors, clinicians to make a quick and an exact detection and classification of brain tumors by their carefully examination of MRI sample images.

Overall, this research contributes to the advancement of precision medicine in neurology by providing a reliable and effective tool for brain tumor detection and classification. The developed deep learning model serves as a valuable asset in clinical practice, aiding healthcare professionals in making informed decisions and improving patient outcomes.

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