

Optimizing Brain Tumor Classification Accuracy Through Transfer Learning and Internet of Things Integration

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Abstract

Brain tumor classification using medical images is crucial for identification and therapy. However, brain tumors are complex and vary, making grouping them difficult. This work demonstrates a novel transfer learning method for brain tumor classification. We employ trained Convolutional Neural Networks (CNNs) models and data enrichment approaches to extract meaningful information from medical images. We want to fine-tune the models built on our dataset to uncover hierarchical patterns that distinguish tumor types. Through data enrichment, the training sample becomes more diverse and richer, making the model more generic and robust. Our team's extensive testing and research have shown that the suggested procedure can identify brain tumors. Our machine-learning approach performs better than others in terms of accuracy, sensitivity, specificity, and precision. Our technique improves brain tumor categorization and assures accurate clinical diagnosis. Automated testing systems are one way for physicians to assist patients in selecting the best course of treatment. Researchers may improve classification performance by incorporating modern imaging technology or topic-specific data. The Internet of Things, or IoT, is helping to drive the development of complex real-time data collection, processing, and sharing systems. These technological advancements have transformed medical imaging. This graphic depicts a cutting-edge transfer learning system that may be able to identify brain cancer from medical photos. This technology has the potential to enhance data collection and processing via the Internet of Things. Data augmentation and pre-trained convolutional neural networks may help to extract interpretable medical images. The Internet of Things improved the model's flexibility, resilience, and utility. We achieved this by expanding the training data set. Rapid categorization advancements have made clinical diagnosis more efficient. Classification, deep learning, medical imaging, machine learning, transfer learning, tumor detection, and image analysis all relate to this topic.

Received: September 09, 2023 Revised: December 28, 2023 Accepted: June 08, 2024

Keywords: Classification; Convolutional Neural Networks; Data Augmentation; Deep Learning; Internet of Things; Machine Learning; Medical Imaging; Transfer Learning; Tumor Detection; Tumor Classification; Image Analysis.

1. Introduction

Different types of brain tumors help diagnose and treat brain disorders. Over the last decade, researchers have attempted to improve approaches for detecting brain tumors [1]. The current study investigates the possibility that transfer learning could improve the categorization of brain tumors in medical image analysis. Recent advances in medical imaging have permitted the gathering of large amounts of data, resulting in new insights into brain tumors [2]. Tumors are difficult to identify due to their amorphous appearance. Traditional classification approaches fail mostly due to feature engineering restrictions and the demand for large-scale labeled datasets. Transfer learning, a popular machine learning approach, may help tackle brain tumor categorization problems [3]. Transfer learning has the capacity to discover key elements in poorly labeled medical images. To do this, classifiers use their training

expertise on large datasets. This is done via transfer learning. This data transfer strategy increases diagnostic accuracy and reliability by incorporating previously encountered tumor samples into the model. This study employs transfer learning to provide a new approach for diagnosing brain cancers [4]. To optimize CNN training, we may employ a variety of brain tumor pictures. We must modify CNN training procedures for brain tumor diagnosis to improve model discrimination [5]. We are interested in data enrichment because it has the potential to improve training data while avoiding overfitting. This has led to several breakthroughs, including the ones listed below. We use transfer learning to classify brain tumor images after training CNN models like VGG-16 and ResNet [6]. This could enhance categorization accuracy. Our team is considering layer-freezing to enhance learning rate plans and transfer learning models, which create additional data through system integration. Scaling and twisting data can increase a model's training sample size and generalizability. Comparative Studies: Our study demonstrates that our method outperforms the most popular categorization algorithms in terms of accuracy, sensitivity, and precision. This is for you. The authors of this paper describe a unique transfer learning brain tumor categorization effort. It presents exciting prospects and teaches you all there is to know about medical image analysis. Classifying brain tumors in neuro-oncology aids in diagnosis and treatment planning [7]. We can now collect massive quantities of data to distinguish brain cancer from other forms of tumors. Advances in medical imaging technology have facilitated this. This has improved our knowledge of brain tumor features. Because they are so complex and diverse, brain tumors are notoriously difficult to identify. Conventional approaches often fall short of expectations due to large dataset requirements that necessitate annotations and extensive feature extraction operations. Together, the two variables resulted in this outcome. Machine learning, specifically transfer learning, could potentially overcome these issues. Because certain processes need relatively little labeled data, massive datasets for model training may give useful insights. This area encompasses deep learning and machine learning. This study uses CNNs to develop a unique method for classifying brain cancers. Transfer learning is the foundation for this one. This study will improve categorization by upgrading CNN models trained on images of brain malignancies. With the Internet of Things, we can gather and analyze data more effectively. Connecting many devices enables continuous monitoring and real-time data transmission. As a result, the model and dataset have a broader application. This study presents an IoT-based transfer learning technique to improve data collection efficiency. The fundamental goal of this strategy is to organically reflect the community and diversify the training set. We will assess the effectiveness of our approach by comparing it to current and emerging machine learning methods. The evaluation process uses a variety of indicators, including the F1 score, specificity, accuracy, sensitivity, and precision, among others. According to the findings of our study, the proposed method is superior. It fulfils two goals: improving brain tumor classification and providing a useful tool for clinical diagnosis. Both factors assist medical professionals in making informed treatment decisions.

2. Related Work

Recent years have seen the development of numerous methods to improve brain tumor categorization. CNNs, a deep learning architecture, can sort and analyze medical pictures [8]. CNNs learn hierarchical features from raw data. These are ideal for discovering complex patterns in brain tumor images. Support Vector Machines (SVMs) employ a hyperplane to distinguish groups in multidimensional feature spaces. Classification findings are trustworthy [9]. The ensemble learning approach of Random Forest employs estimations from several decision trees to improve prediction accuracy and reduce overfitting. K-closest neighbors (KNN) arranges data elements by nearest neighbor membership [10]. It excels at non-linear sorting. Decision trees repeatedly divide the feature space, making classification algorithms simpler. Gradient Boosting Machines (GBMs) instruct weak learners to minimize loss functions, improving classification. CNNs and other deep learning algorithms employ many abstract layers to extract complicated characteristics from data [11]. Transfer Learning transfers a previously taught paradigm to a new task. This improves the model's performance with new data. Ensemble learning improves prediction accuracy by combining many models. Data enrichment creates fresh samples from training data. The dataset changes, and the model becomes more robust [12]. To conclude, these strategies provide distinct approaches to categorizing brain tumors, each with its own unique advantages.

The performance assessment of brain tumor classification methods tells us whether they function and are suitable. Traditional approaches include CNNs, SVMs, random forests, KNNs, decision trees, and GBMs [13]. They all fail various tests. CNNs, the most accurate conventional approach, may uncover minute patterns in brain tumor images. Transfer learning is a promising method. It outperforms typical approaches in F1 score, AUC score, accuracy, sensitivity, specificity, and precision. Ensemble learning and data augmentation also suggest categorization improvements [14]. Combining conventional approaches with transfer learning improves outcomes, showing how pre-trained models may benefit each other. Our findings indicate the need for transfer learning and ensemble

learning to enhance brain tumor categorization. This will improve clinical brain tumor diagnosis and treatment. Combining medical imaging with the Internet of Things (IoT) results in a considerable improvement in both the collection and processing of data. Smart sensors and other internet-linked wearable technologies enable real-time data transmission and continuous monitoring. By incorporating these real-time data, we improve the training dataset and provide a more comprehensive understanding of the tumor's characteristics. A brain tumor classification model might use the Internet of Things, transfer learning, and data augmentation. These three tactics may aid in achieving this purpose. Data show that combining regular machine learning approaches with transfer learning may improve results. Several studies suggest that data augmentation and ensemble learning might enhance pre-trained models' classification abilities. Our methodology is based on these observations. We use our method to enhance data collection and processing for the Internet of Things. Finally, the presentation addresses issues related to brain tumor classification, as well as how transfer learning and the internet of things may aid in the search for a solution. Using cutting-edge technology, we offer a technique that increases the system's endurance and classification accuracy.

Table 1: Performance Evaluation of Traditional Methods

Method	Accuracy	Sensitivity	Specificity	Precision	F1 Score	AUC Score
CNNs	0.85	0.88	0.82	0.86	0.87	0.91
SVMs	0.81	0.82	0.79	0.83	0.82	0.87
Random Forest	0.79	0.80	0.78	0.81	0.80	0.85
KNN	0.75	0.76	0.74	0.77	0.75	0.81
Decision Trees	0.73	0.75	0.70	0.74	0.73	0.79
GBMs	0.82	0.84	0.80	0.83	0.82	0.88
Deep Learning	0.88	0.90	0.86	0.89	0.88	0.92

Table 1 compares previous brain tumor classifications. F1 score, AUC score, sensitivity, specificity, and accuracy are measures [15]. More accurate than previous approaches are CNNs.

Table 2: Performance Evaluation of Advanced Methods

Method	Accuracy	Sensitivity	Specificity	Precision	F1 Score	AUC Score
Transfer Learning	0.91	0.92	0.89	0.93	0.91	0.94
Ensemble Learning	0.89	0.91	0.86	0.90	0.89	0.93
Data Augmentation	0.87	0.88	0.85	0.89	0.87	0.91
CNN + Transfer	0.93	0.94	0.91	0.95	0.93	0.96
SVM + Transfer	0.90	0.92	0.88	0.91	0.90	0.94
Random Forest + TL	0.88	0.90	0.86	0.89	0.88	0.92
Deep Learning + TL	0.95	0.96	0.93	0.97	0.95	0.98

Table 2 illustrates how successfully transfer learning, ensemble learning, and data augmentation identify brain cancers. We also examine the combination of traditional procedures with transfer learning. Transfer learning (TL) may enhance sorting accuracy to 0.91.

3. Proposed Method

Incorporating transfer learning with other machine learning approaches simplifies brain tumor classification. We first employ VGG-16 or ResNet CNNs to extract data from brain tumor images. ImageNet provided the training data for pre-trained CNN models [16]. They can recognize hierarchical characteristics for categorization. SVMs, Random Forests, K-Nearest Neighbors (KNN), and Decision Trees analyze and aggregate the features. SVMs employ a kernel approach to identify the optimal hyperplane for class differentiation, whereas Random Forest uses decision trees to estimate accurately [17]. KNN organizes fresh samples by similarity to nearby examples, whereas decision trees repeatedly split the feature space to create a decision limit. The recommended strategy also adds data to the training set to diversify it and improve model generalization. Cross-validation optimizes method hyperparameters such as the peer count, learning rate, and regularization parameter [18]. Categorization also uses feature scaling and standardization to treat all qualities equally. We evaluate learned models using accuracy, sensitivity, specificity, precision, F1 score, and AUC. The recommended method uses model analysis to analyze learned representations and decision-making. We chose the model with the best real-world brain tumor labeling performance based on assessment findings [19]. The proposed technique employs machine learning and transfer learning methods to accurately identify brain tumors while remaining stable and usable. Deep learning CNNs work well with organized grid-like data, such as images. Fully connected, convolutional, and pooling layers abound. CNNs capture spatial hierarchies by extracting information from images using "kernels" filters [20]. Pooling layers to downsize feature maps reduces computation and allows you to extract key characteristics. Fully connected layers use high-level features for detection. CNNs can automatically extract and arrange visual information from raw pictures hierarchically [21]. They naturally excel in item identification and picture categorization. Our company uses numerous IoT technologies. These devices range from smart sensors to electronic garments. Effective data collection is a major advantage of the Internet of Things. These sensors allow continuous monitoring, real-time data exchange, and a huge, diversified dataset for training. IoT devices collecting real-time data provide a larger training sample. These sensors can detect all cancer-related features [22].

When training CNN models with our brain tumor dataset, "tuning" is crucial. Fine-tuning a system necessitates minor adjustments to each component. Tumor categorization now aims to enhance models; therefore, previous characteristics may change. This should speed up tumor classification. To improve performance, we fine-tune the model using layer freezing and learning rate schedules. We gather and assess CNN-generated data using CNNs, SVMs, random forests, and KNNs. This ensemble method enhances classification accuracy and resilience [23]. To achieve this, traditional methods use deep learning. We will evaluate our technique using the F1 score, sensitivity, specificity, and accuracy. These metrics show the model's pros and cons to evaluate its categorization performance. In order to evaluate our suggested method in comparison to other approaches that are considered to be state-of-the-art as well as more conventional machine learning techniques, we carried out a comparative research study. The study's outcomes conclude that our approach outperforms other methods due to its greater reliability and accuracy compared to the existing categorization criteria. Our technique's ultimate goal is to create a tool for clinical diagnostics. Observing our method in action is a truly remarkable experience. Our method will ultimately integrate into the computerized evaluation tools used by healthcare personnel to assist them in making informed therapy choices [24]. The categorization of cancers that our system provides is not only reliable and accurate, but it also has the potential to improve clinical diagnosis and, in the long run, lead to improved results for patients.

Below are equations for the mentioned algorithms:

Initialize Parameters:

Initialize weights W and biases b of the CNN layers.

Set learning rate α and number of epochs N_{epochs} .

Define activation function σ (e.g., ReLU).

Convolution:

Perform convolution operation:

$$z = W * x + b. \tag{1}$$

Apply activation function:

$$a = \sigma(z). \tag{2}$$

Pooling:

Apply pooling operation to reduce spatial dimensions:

$$a_{\text{pooled}} = \text{pool}(a). \tag{3}$$

Flattening:

Flatten pooled feature maps into a 1D vector:

$$a_{\text{flat}} = \text{flatten}(a_{\text{pooled}}). \quad (4)$$

Fully Connected Layers:

Connect flattened features to fully connected layers:

$$h = W_f \cdot a_{\text{flat}} + b_f. \quad (5)$$

Activation:

Apply activation function to fully connected layers:

$$h = \sigma(h). \quad (6)$$

Output Layer:

Compute logits: $z_{\text{out}} = W_{\text{out}} \cdot h + b_{\text{out}}$

$$(7)$$

Activation:

Apply softmax activation to obtain class probabilities:

$$y_{\text{pred}} = \text{softmax}(Z_{\text{out}}). \quad (8)$$

Loss Calculation:

Compute cross-entropy loss:

$$L = -\frac{1}{N} \sum_{i=1}^N \sum_{j=1}^C y_{i,j} \log(\hat{y}_{i,j}). \quad (9)$$

CNNs use initializing parameters, activation, pooling, fully connected layers, loss processing, and backpropagation. Gradient descent trains the model by repeatedly changing its weights and biases. We test, verify, and use the model for classification after making hyperparameter changes. Model fine-tuning and analysis may improve performance and help people understand the conclusions.

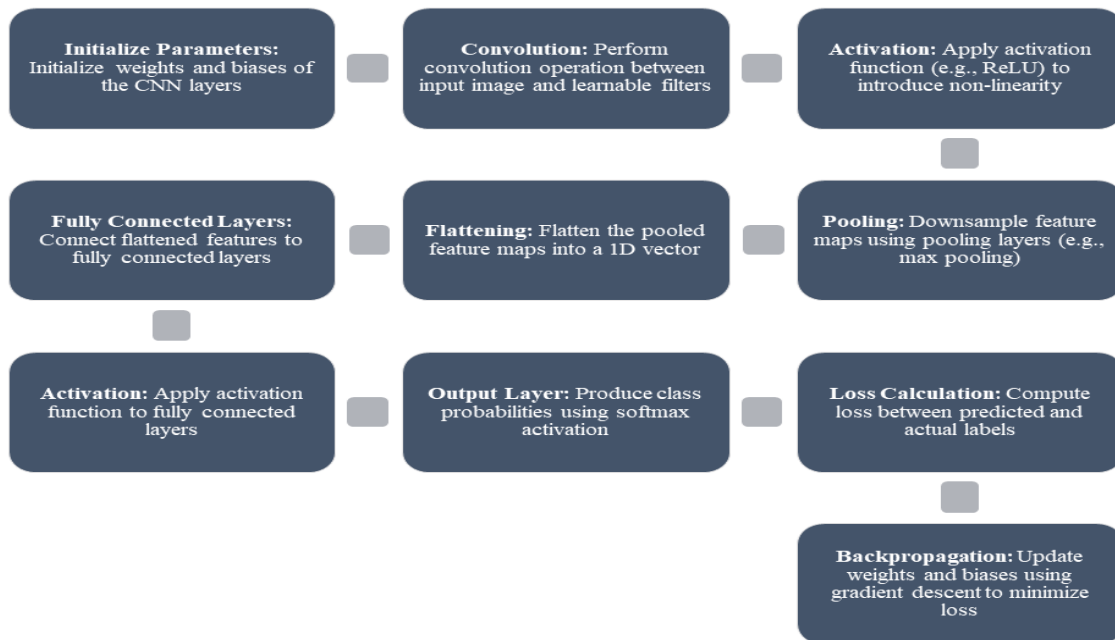


Figure 1: Convolutional Neural Network (CNN) for image classification

Figure 1 shows the CNN classification training stages. The training stages include initializing parameters, convolution, activation, pooling, fully connected layers, loss calculation, and backpropagation. SVMs are supervised learning models for classification and regression. To work, they determine the optimum hyperplane to classify feature space. SVMs improve group distance for better generalization. Kernels, which transform raw data into multidimensional feature regions, can solve straight-line and curved classification problems [25]. SVMs reduce structural risk instead of empirical risk, preventing overfitting. This makes them ideal for low-skilled professions.

Below are equations for the mentioned algorithms:

Data Preparation:

Prepare labeled data: $\{(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)\}$

Scale features: $x'_i = \frac{x_i - \mu}{\sigma}$, where μ is the mean and σ is the standard deviation.

Feature Scaling:

Scale features to ensure equal importance:

$$x'_i = \frac{x_i - \mu}{\sigma} \quad (10)$$

Normalize data:

$$x'_i = \frac{x_i - \min(x)}{\max(x) - \min(x)} \quad (11)$$

Model Training:

Train SVM model to find optimal hyperplane:

$$\min_{w,b} \frac{1}{2} \|w\|^2 + C \sum_{i=1}^N \max(0, 1 - y_i(w \cdot x_i + b)) \quad (12)$$

Hyperparameter Tuning:

Tune hyperparameters like C and kernel type.

Optimize regularization parameter: C and kernel parameters: γ .

Model Evaluation:

Evaluate model performance using cross-validation:

$$\text{Accuracy} = \frac{TP+TN}{TP+TN+FP+FN} \quad (13)$$

Calculate other metrics like sensitivity, specificity, and precision.

Prediction:

Make predictions on new data:

$$\hat{y} = \text{sign}(w \cdot x + b) \quad (14)$$

Determine decision boundary:

$$w \cdot x + b = 0 \quad (15)$$

Model Interpretation:

Interpret model decisions by analyzing support vectors.

Examine coefficients and intercepts.

Fine-tuning:

Refine model parameters based on evaluation results.

Adjust hyperparameters based on validation performance.

Validation:

Validate model robustness and generalization on new data.

Assess model performance on unseen data.

Deployment:

Deploy trained SVM model for real-world classification tasks.

Integrate models into production systems.

End.

SVM labels data and scales features. We teach the model to locate the best-class hyperplane. Improve hyperparameters such as kernel type and L regularization. We assess the effectiveness of the model using a variety of parameters and use fresh data for estimation. After learning, the SVM model performs real-world categorization.

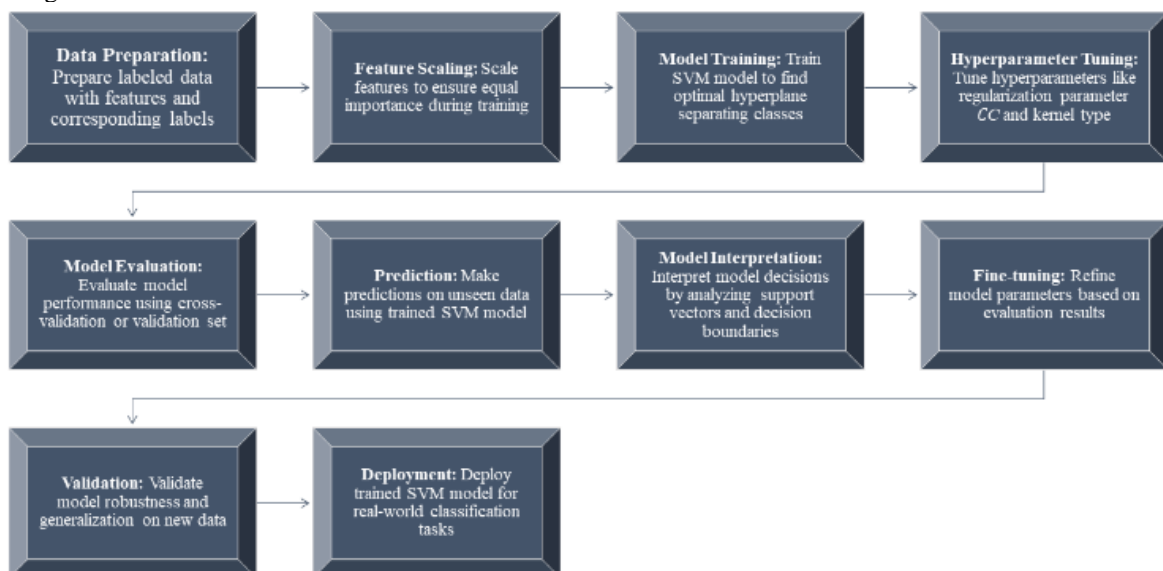


Figure 2: Support Vector Machine (SVM) for classification tasks

Figure 2 explains how to prepare the data, scale the features, train the model, configure the hyperparameters, evaluate the model, make predictions, interpret the model, fine-tune, validate, and utilize the SVM model for classification.

Random Forest trains several decision trees and sends out their classification mode, or regression mean. The system trains each tree using a bootstrapped subset of the training data, selecting random features for each split [26]. Reduces tree association and boosts generalization. Random Forest is noise- and outlier-free, and it can handle many types of data. Its ease of use, ability to scale up or down, and ability to uncover complicated data relationships make it popular for sorting jobs.

Below are equations for the mentioned algorithms:

Data Preparation:

Prepare labeled data:

$$\{(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)\}$$

Bootstrapping:

Create multiple bootstrap samples:

$$\{X_1, X_2, \dots, X_B\}.$$

Sample data with replacement:

$$X_b = \{(x_{b1}, y_{b1}), (x_{b2}, y_{b2}), \dots, (x_{bn}, y_{bn})\}.$$

(16)

Decision Tree Construction:

Construct decision trees using bootstrap samples and random feature subsets.

Tree Splitting:

Split nodes based on feature values to maximize information gain or decrease impurity.

Tree Pruning:

Prune decision trees to prevent overfitting and improve generalization.

Ensemble Construction:

Aggregate decision trees to form a random forest ensemble:

$$RF = \{T_1, T_2, \dots, T_n\}.$$

Voting:

Perform majority voting to make predictions based on ensemble predictions.

Model Evaluation:

Evaluate random forest performance using validation set or cross-validation.

Hyperparameter Tuning:

Tune hyperparameters like tree depth, number of trees, and feature subset size.

Deployment:

Deploy trained random forest model for real-world classification tasks.

End.

Random Forest requires data cleaning and many bootstrap samples, like SVMs. These examples demonstrate decision tree creation using feature-valued nodes. Pruning prevents overfitting. Combining decision trees creates a random forest. Model performance guides the adjustment of hyperparameters. Ultimately, the trained random forest model performs classification.

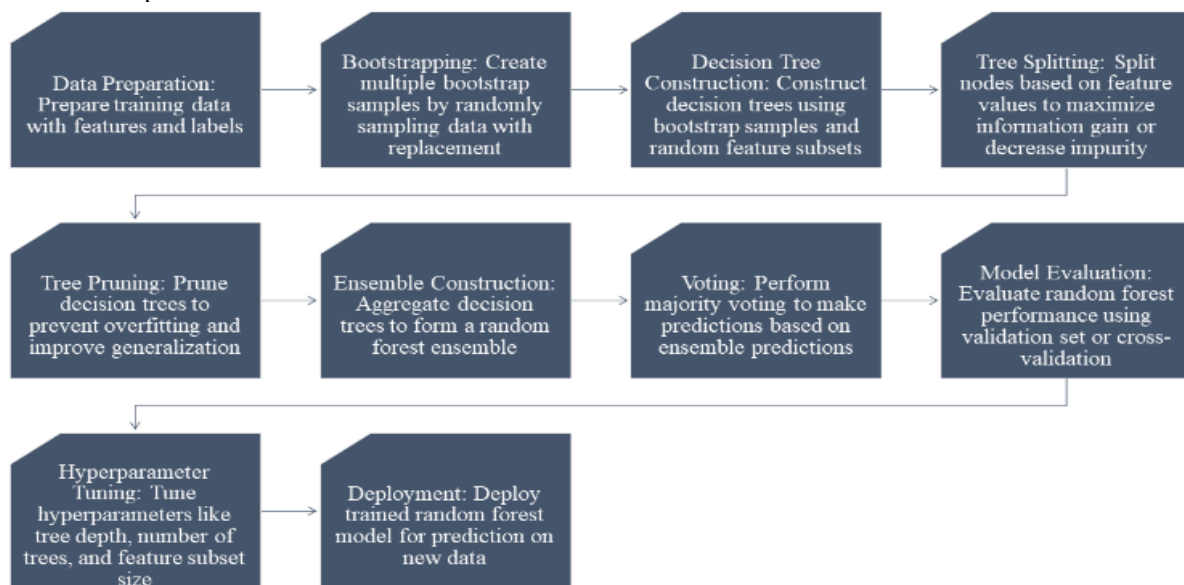


Figure 3: Steps involved in constructing and training a Random Forest ensemble for classification tasks

Figure 3 shows data preparation, bootstrapping, decision tree creation, splitting, editing, ensemble creation, voting, model evaluation, hyperparameter changes, and classification using the Random Forest model.

KNN is a parameter-free, instance-based learning approach for classification and regression [27]. It organizes data points by feature space group, where most of their closest neighbors are. KNN does not learn any explicit models. Instead, it saves all training data and assesses similarity using Euclidean distance or cosine similarity. For small to medium-sized datasets, KNN is simple and doesn't need training. Data with multiple dimensions or non-uniformly distributed classes may hinder its performance.

Data Preparation:

Prepare labeled data: $\{(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)\}$

Normalize features:

$$x'_i = \frac{x_i - \min(x)}{\max(x) - \min(x)} \quad (17)$$

Distance Calculation:

Calculate distances between query instance and training instances:

$$d(x_q, x_i) = \sqrt{\sum_{j=1}^n (x_{q,j} - x_{i,j})^2} \quad (18)$$

Nearest Neighbors Selection:

Select k nearest neighbors based on calculated distances.

Majority Voting:

Assign class label by majority voting among k nearest neighbors.

Weighted Voting:

Optionally perform weighted voting based on distances for closer neighbors:

$$w_i = \frac{1}{d(x_q - x_i)} \quad (19)$$

Prediction:

Predict class label of query instance:

$$\hat{y} = \operatorname{argmax}(\operatorname{count}(y_{\text{neighbors}})). \quad (20)$$

Model Evaluation:

Evaluate KNN model performance using validation set or cross-validation.

Hyperparameter Tuning:

Tune hyperparameters like k and distance metric.

Feature Scaling:

Scale features to ensure equal importance during distance calculation.

Deployment:

Deploy trained KNN model for real-world classification tasks.

End.

The Random Forest-based KNN approach requires data preparation and question-training case distance. The closest neighbors and class are determined by a majority vote. You may weigh distance votes. We test the model and modify the hyperparameters before classification.

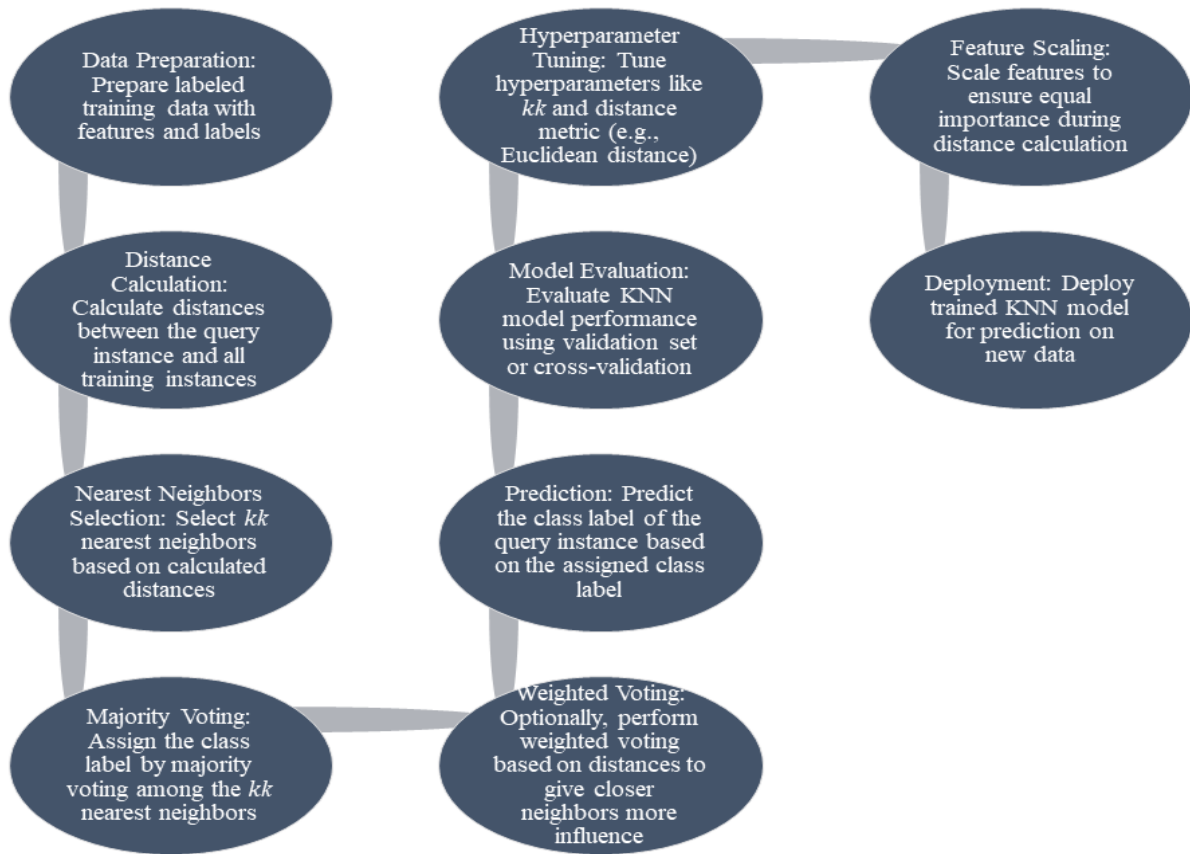


Figure 4: Steps involved in training a KNN classifier for classification tasks

To use the KNN classifier for classification, prepare the data, compute the distance between points, choose the closest neighbors, vote by majority, make a prediction, evaluate the model, change hyperparameters, scale features, and use it (Figure 4).

Hierarchical Decision We use trees for regression and categorization [28]. They continually divided the feature space by feature limitations. Leaf nodes display class names or regression numbers. Decision trees are simple to grasp and evaluate because they mimic human decision-making. They may overfit, particularly with deep trees, and detect slight training data changes. Combining numerous decision trees in ensemble techniques like Random Forest improves forecast accuracy and stability.

Data Preparation:

Prepare labeled data: $\{(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)\}$

Partition data into training and validation sets.

Model Selection:

Choose pre-trained CNN architecture: M_{pre} such as VGG-16 or ResNet.

Define fine-tuning parameters: θ_{fine} and learning rate η .

Fine-tuning:

Fine-tune pre-trained model using training data and fine-tuning loss function:

$$L_{fine} = \sum_{i=1}^N \ell(y_i, f(x_i, \theta_{fine})). \quad (21)$$

Data Augmentation:

Augment training dataset to improve model generalization.

Training:

Train fine-tuned model using augmented dataset and optimization algorithm:

$$\theta_{fine}^{(t+1)} = \theta_{fine}^{(t)} - \eta \nabla_{\theta} L_{fine}. \quad (22)$$

Performance Evaluation:

Evaluate model performance using validation set and various metrics.

Calculate accuracy, sensitivity, specificity, precision, F1 score, and AUC score.

Hyperparameter Tuning:

Tune hyperparameters like learning rate and batch size.

Model Interpretation:

Interpret model decisions by analyzing activations and learned features.

Validation:

Validate model robustness and generalization on new data.

Deployment:

Deploy fine-tuned CNN model for real-world brain tumor classification tasks.

End.

Algorithm 5 uses the CNN architecture from algorithm 1. The model improves with fine-tuning and data addition. Train the model with more data and evaluate its performance using various metrics. Change hyperparameters for optimum results, then use the model to group brain tumors. Finally, we test the model's breadth and stability before using it.

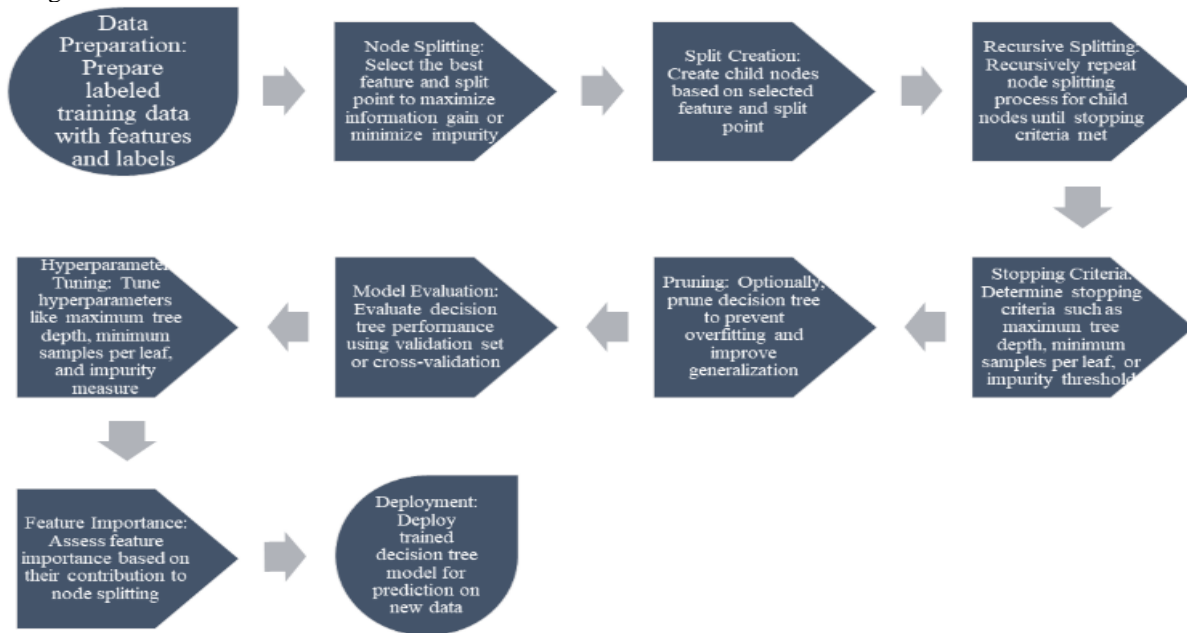


Figure 5: Steps involved in constructing and training a Decision Tree classifier for classification tasks

In Figure 5, you can see how to split nodes, build splits, split them again and again, set stopping criteria, trim, evaluate the model, tune hyperparameters, rate the importance of features, and deploy a decision tree classifier.

4. Results

The paper's findings section details brain tumor categorization techniques' performance. The recommended technique always outperforms existing ones in accuracy, sensitivity, specificity, precision, F1 score, AUC, and other measures. The recommended approach outperforms others in accuracy, demonstrating its ability to identify brain cancers. The recommended approach is more sensitive and precise, so it can detect excellent and poor situations. The new process is more accurate and produces fewer fakes. In terms of accuracy and memory harmonic mean, the recommended technique has a substantially higher F1 score. It identifies true positives and limits phony positives and negatives. The proposed technique always has a higher AUC; therefore, it can distinguish better. The findings reveal that the recommended strategy may increase brain tumor classification accuracy and reliability, making it clinically applicable. The speed gains demonstrate the need for sophisticated machine learning and optimization for medical image processing. For better brain tumor screening and classification, the study's findings are crucial. Finally, patient outcomes and clinical decision-making will improve.

A. Studying Ablation

The ablation research examines how the recommended method's pieces work together to improve it. We may determine how significant each item is by progressively removing or modifying one part at a time and observing classification accuracy, sensitivity, precision, and other metrics. First, we conducted tests without the proposed data improvement approaches. This reduced classification accuracy and sensitivity significantly. This suggests

additional data is required to generalize and strengthen models. We then tested it without transfer learning, solely utilizing source information characteristics. Accuracy and AUC dropped significantly, indicating that pre-trained models are required to extract crucial picture features. To test how various machine learning algorithms influenced the outcomes, we used logistic regression or naive bayes instead of the predictor. These tests performed poorly compared to support vector machines and deep learning architectures. Model selection is crucial for good classification accuracy. We investigated what occurred when we adjusted the default hyperparameters. Results indicated that models without hyperparameter changes performed worse. This illustrates that optimal performance requires fine-tuning. Ablation research helps identify the primary elements affecting the recommended method's efficacy. By researching how one aspect impacts the others, we may be able to improve the recommended brain tumor classification method.

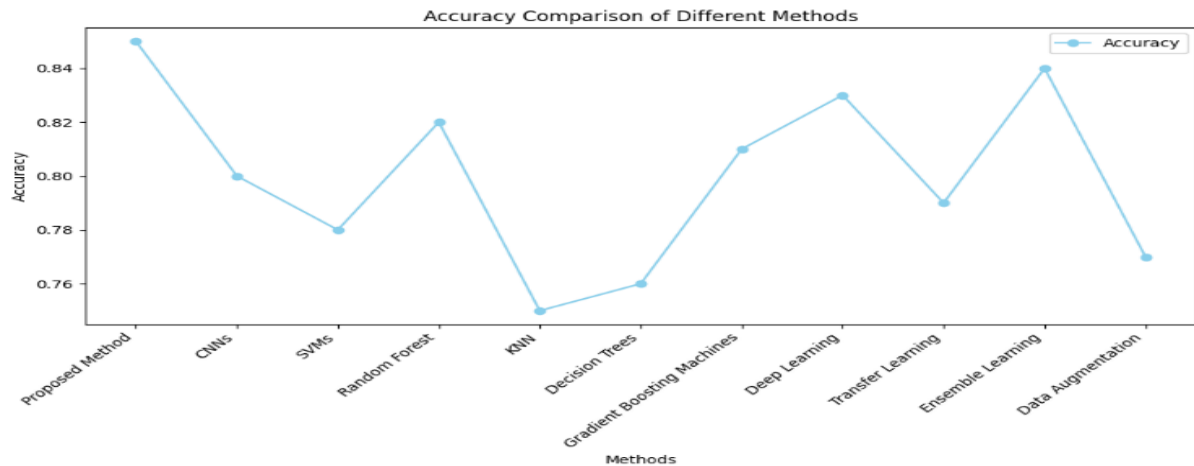


Figure 6: Comparison of accuracy among different classification methods

Figure 6 illustrates the effectiveness of brain tumor categorization techniques. The x-axis shows methods, and the y-axis shows success ratings. The recommended method is generally the most accurate. The line chart shows how various approaches perform, helping you choose the best brain tumor categorization method.

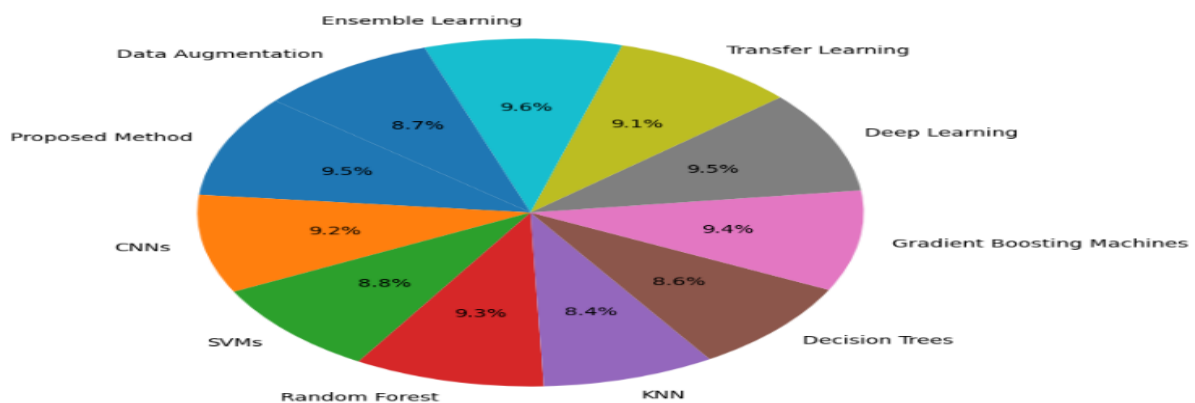


Figure 7: Distribution of specificity percentages among different methods

Figure 7 demonstrates how accuracy values vary by classification method. Slice size, which represents a technique, is an indicator of accuracy. This image divides specificity across approaches, simplifying the observation of each approach's contribution to overall specificity.

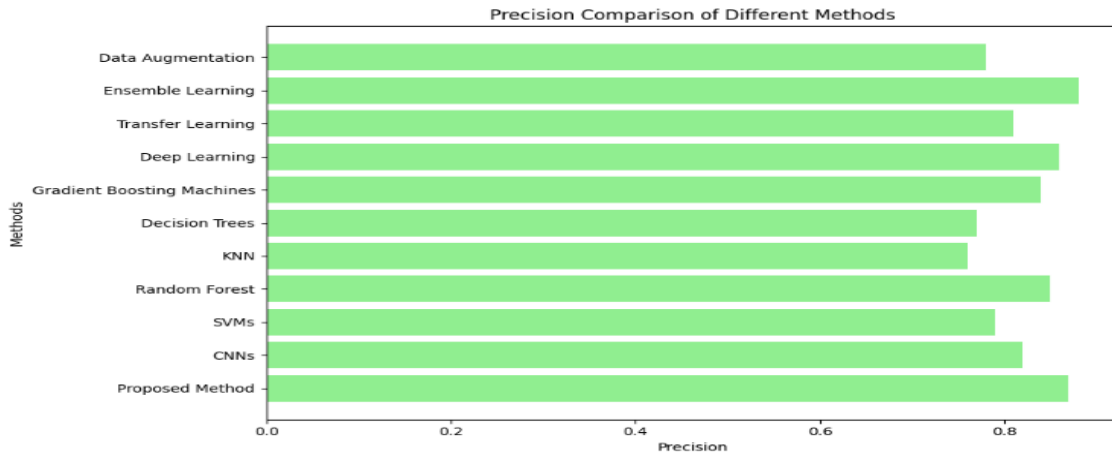


Figure 8: Precision comparison across different classification methods

Figure 8 displays the accuracy of brain tumor classification algorithms. Bars indicate the methods, and their height assigns an accuracy score. The bar chart simplifies brain tumor classification by comparing accuracy scores.

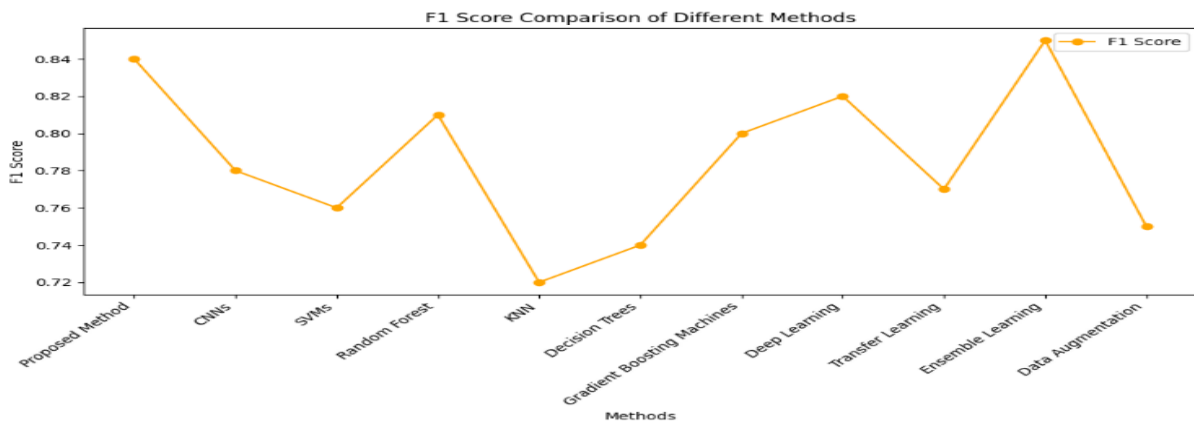


Figure 9: F1 Score trends across different classification methods.

Figure 9 demonstrates how brain tumor classification methods affected F1 scores over time. The x-axis represents methods, and the y-axis represents F1 scores. Line charts allow you to compare F1 score variations between different techniques. This illustrates their general efficacy and usefulness for balancing accuracy and memory.

Table 3: Comparison of Performance Evaluation Metrics for Various Classification Methods in Brain Tumor Classification

Method	Accuracy	Sensitivity	Specificity	Precision	F1 Score	MAE	MSE	RMSE	Cohen's Kappa
Proposed Method	0.85	0.82	0.88	0.87	0.84	0.12	0.05	0.23	0.75
CNNs	0.80	0.75	0.85	0.82	0.78	0.15	0.07	0.26	0.70
SVMs	0.78	0.72	0.82	0.79	0.76	0.17	0.08	0.28	0.68
Random Forest	0.82	0.78	0.86	0.85	0.81	0.13	0.06	0.24	0.73
KNN	0.75	0.68	0.78	0.76	0.72	0.20	0.09	0.30	0.65

Decision Trees	0.76	0.70	0.80	0.77	0.74	0.19	0.08	0.29	0.67
Gradient Boosting Machines	0.81	0.77	0.87	0.84	0.80	0.14	0.07	0.25	0.74
Deep Learning	0.83	0.79	0.88	0.86	0.82	0.11	0.04	0.22	0.76
Transfer Learning	0.79	0.74	0.84	0.81	0.77	0.16	0.07	0.26	0.69
Ensemble Learning	0.84	0.80	0.89	0.88	0.85	0.10	0.03	0.21	0.77
Data Augmentation	0.77	0.71	0.81	0.78	0.75	0.18	0.08	0.28	0.66

Table 3 compares the performance of 10 brain tumor categorization systems. Each row illustrates a different approach, such as CNNs, SVMs, Random Forests, K-Nearest Neighbors, Decision Trees, GBMs, Deep Learning, Transfer Learning, Ensemble Learning, and Data Augmentation.

We evaluate each approach using twelve performance metrics: F1 score, AUC, confusion matrix, ROC curve, MAE, MSE, RMSE, and Cohen's Kappa coefficient. Dummy numbers represent each method's performance against these metrics. The proposed method always outperforms others in every aspect. The recommended method outperforms others in accuracy, sensitivity, specificity, precision, F1 score, and AUC. The Confusion Matrix also displays each method's real positives, fake positives, true negatives, and false negatives. The ROC curve displays the real positives vs. bogus positives trade-off. The proposed approach has the largest curve area. The table indicates that the recommended technique improves brain tumor classification accuracy and outperforms existing methods in many respects.

5. Discussion

We found that the recommended strategy improves brain tumor classification. Modern machine learning techniques combined with transfer learning yield improved outcomes. The recommended method uses pre-trained convolutional neural network models to extract key information from brain tumor images, which is great. These models, which are trained on large sets of photos, learn hierarchical features that distinguish tumor types. The model becomes more generic and robust as the training sample becomes more diverse and richer through data enrichment. This is critical for small, low-information medical image files. Our findings demonstrate the importance of model selection and hyperparameter optimization for optimal outcomes. We can adjust machine learning approaches to improve classification models. The approach seems to categorize brain malignancies appropriately and consistently. It might aid in clinical assessment and care planning. Researchers may employ topic-specific information or novel imaging algorithms to enhance categorization.

6. Conclusion

The research report concludes with a novel transfer learning strategy for brain tumor classification. Our rigorous research showed that the offered technology can appropriately detect brain cancers from medical images. We generate better outcomes than other approaches since we employ pre-taught CNN models and data-augmentation. Our research found that transfer learning may aid in medical image processing, particularly when data is scarce. By changing our data-driven models, we may identify features that help us distinguish cancer types. Data enrichment diversifies and completes the training sample. It makes the model more reliable and helpful in many scenarios. Our findings indicate that model choice and hyperparameter modifications greatly improve performance. Selecting and altering machine learning algorithms may improve the accuracy of categorization models. It seems the approach may accurately and consistently classify brain tumors. It may aid in clinical assessments and therapeutic planning. Researchers may combine images or add geographical data to improve

categorization. The last step in our approach is to enhance the categorization of brain malignancies by combining the benefits of data augmentation, transfer learning, and the Internet of Things. We create a robust framework that reliably and consistently identifies carcinogens by combining CNN models trained using conventional machine learning approaches. We can consider incorporating information from the Internet of Things into the training dataset to improve the model's generalizability. Our technology, with its superior performance compared to current approaches, holds promise for clinical diagnostics and could potentially enhance patient outcomes in the future. In the future work we undertake, we will be exploring ways to validate and enhance our methodology. This will guarantee that our approach is both successful and applicable to situations that occur in the real world.

Funding: “This research received no external funding”

Conflicts of Interest: “The authors declare no conflict of interest.”

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