Brain Tumor Classification with Hybrid Deep Learning Models from MRI Images

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Abstract: Brain tumor classification using MRI images plays a crucial role in medical diagnostics, enabling early detection and improving treatment planning. Traditional diagnostic methods are often subjective and time-intensive, emphasizing the need for automated and precise solutions. This study explores hybrid deep learning models alongside fine-tuned pre-trained architectures for classifying brain tumors into four categories: glioma, meningioma, pituitary tumor, and healthy brain tissue. The proposed approach incorporates both hybrid ensemble models—Ens-VGG16-FT-InceptionV3, Ens-ViT-FT-InceptionV3, and Ens-CNN-ViT—and fine-tuned architectures—FT-VGG16, FT-VGG19, and FT-InceptionV3. To enhance model robustness and generalization, data augmentation techniques such as rotation and scaling were applied. Among these models, the hybrid ensemble Ens-VGG16-FT-InceptionV3 achieved the highest accuracy and F1-score of 99%, outperforming both standalone models and other hybrid configurations. These findings demonstrate the effectiveness of integrating complementary architectures for improved brain tumor classification. Ultimately, this study highlights the potential of hybrid ensemble learning to advance brain tumor diagnostics, providing more accurate, reliable, and scalable medical imaging solutions.

SCIENCE AND TECHNOLOGY PUBLICATIONS

1 INTRODUCTION

Brain tumors are among the most serious and lifethreatening medical conditions, contributing significantly to global morbidity and mortality. Recognized by the World Health Organization (WHO) as a significant public health concern (Arabahmadi et al., 2022), the increasing prevalence of brain tumors underscores the urgent need for improved diagnostic techniques. Early and precise diagnosis is essential for effective treatment planning and better patient outcomes. Magnetic Resonance Imaging (MRI) is widely used for detecting and analyzing brain abnormalities, offering detailed insights into tumor size, location, and type (Lee et al., 2020). However, manual interpretation of MRI scans by radiologists is time-consuming and subject to variability, particularly in distinguishing subtle tumor variations (Rivkin and Kanoff, 2013). Recent advancements in artificial intelligence (AI) and deep learning have demonstrated significant potential in automating this process (Bharadiya, 2023), leading to faster and more accurate brain tumor classification (Pham et al., 2023).

Despite progress in medical imaging, brain tumor classification remains a challenging task. Traditional diagnostic methods depend on manual evaluations, which are inherently subjective and require extensive expertise (Sanvito et al., 2021). While machine learning has shown promise in automating classification, existing studies often face limitations such as small datasets, simplistic architectures, or inadequate feature extraction (Iqbal et al., 2023), (Mokni and Haoues, 2022). Additionally, many models struggle to generalize across diverse patient populations, emphasizing the need for robust, scalable solutions capable of accurately classifying tumors into glioma, meningioma, pituitary tumors, and healthy brain tissue (Farmanfarma et al., 2019).

This study aims to address the challenges of brain tumor classification from MRI scans by leveraging advanced deep learning techniques. The primary goal is to explore and compare the performance of various hybrid deep learning models in detecting and catego-

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rizing brain tumors into four classes: glioma, meningioma, pituitary tumor, and no tumor. To achieve this, the research focuses on:

- Applying Advanced Deep Learning Models: Utilizing Convolutional Neural Networks (CNNs), Vision Transformers (ViTs), and transfer learning with fine-tuned pre-trained models (InceptionV3, VGG16, and VGG19) to improve classification accuracy and performance in brain tumor detection. The fine-tuned models are referred to as FT-InceptionV3, FT-VGG16, and FT-VGG19.
- Exploring Hybrid Architectures and Ensemble Learning: Developing and evaluating innovative hybrid architectures by combining multiple models, such as CNN and ViT (Ens-CNN-VIT), ViT and FT-InceptionV3 (Ens-VIT-FT-InceptionV3), and FT-VGG16 and FT-InceptionV3 (Ens-FT-VGG16-FT-InceptionV3). Ensemble learning enhances diagnostic precision by leveraging the complementary strengths of different models, mitigating their weaknesses, and improving overall robustness and generalization.

Through these objectives, the study aims to contribute to the growing field of AI-driven medical diagnostics, ultimately enabling early, accurate detection and improved treatment of brain tumors, while reducing the global burden of these life-threatening conditions.

The remainder of this paper is organized as follows: Section 2 reviews related work on brain tumor classification and the application of deep learning in medical imaging. Section 3 outlines the proposed hybrid models. Section 4 presents the experimental results, followed by a comparative evaluation with previous studies in Section 5. Finally, Section 6 concludes the paper and outlines future research directions.

2 RELATED WORK

In this section, we review the literature and findings from research studies that focused on brain tumor detection.

Gómez-Guzmán *et al.*, (Gómez-Guzmán et al., 2023) evaluated seven deep CNN models for brain tumor classification using MRI scans. Their study compared a generic CNN with six pre-trained models, including InceptionV3, ResNet50, and Xception. Among these, InceptionV3 achieved the highest accuracy of 97.12%, demonstrating its ability to extract robust and distinctive features for classification. Their findings emphasized the effectiveness of CNNs

in identifying brain tumors by capturing hierarchical patterns in MRI images. This research contributed to the advancement of automated brain tumor classification, reinforcing the suitability of CNN-based architectures for medical imaging applications.

Khaliki and Basarslan(Khaliki and Başarslan, 2024) advanced this research by incorporating transfer learning techniques, leveraging pre-trained models like VGG16, VGG19, EfficientNetB4, and InceptionV3. These architectures, initially trained on large-scale datasets, were fine-tuned for brain tumor classification tasks, achieving impressive accuracies of 98% and 97% with VGG16 and Efficient-NetB4, respectively. The use of transfer learning significantly reduced training time while enhancing the models generalization capabilities on smaller medical datasets. Their hybrid classification system demonstrated that leveraging pre-trained networks could bridge the gap between limited medical data availability and high-precision diagnostics.

Balamurugan and Gnanamanoharan (Balamurugan and Gnanamanoharan, 2023) and Fki *et al.*, (Fki et al., 2024)introduced a hybrid approach that combined a Deep Convolutional Neural Network (DCNN) with the enhanced LuNet algorithm for both segmentation and classification of brain tumors. Their model achieved unprecedented accuracies of 99.4% in segmentation and 99.5% in classification, emphasizing the efficacy of integrating advanced feature extraction techniques with deep learning models.

Dhakshnamurthy *et al.*, (Dhakshnamurthy et al., 2024) proposed a hybrid model combining VGG16 and ResNet-50 architectures, achieving a ground-breaking accuracy of 99.98%. This model outperformed standalone architectures such as AlexNet, VGG16, and ResNet-50, emphasizing the effectiveness of combining complementary strengths to enhance diagnostic precision. The integration of ResNet skip connections with VGG16 deep feature extraction layers facilitated improved gradient flow during training, addressing challenges associated with vanishing gradients in deep networks.

Mahmud and Muntasir *et al.*, (Mahmud et al., 2023) further contributed to this field by designing a CNN-based approach for early detection of brain tumors, achieving an accuracy of 93.3%. Their model incorporated extensive preprocessing and data augmentation techniques, including rotation, flipping, and zooming, to enhance model robustness. By comparing their CNN model with architectures such as ResNet-50 and InceptionV3, they demonstrated that carefully designed CNN architectures could outperform more complex models in tasks requiring precision and computational efficiency.

Asiri *et al.*, (Asiri et al., 2023) explored the capabilities of Vision Transformers (ViTs) in medical imaging by developing a fine-tuned Vision Transformer (FT-ViT) for brain tumor classification. The FT-ViT achieved a classification accuracy of 98.13% on the BraTS dataset, showcasing exceptional precision and recall across various tumor classes. This approach marked a paradigm shift from convolutionbased methods to transformer-based architectures, which rely on self-attention mechanisms to capture global and local relationships in visual data. The findings demonstrated the ViTs ability to process MRI images as sequences of image patches, enabling a more nuanced understanding of tumor morphology.

Table 1: Summary of previous studies on brain tumor classification.

| Proposal | Dataset | Method | Result |
|------------------|----------|-------------------|--------|
| Gómez- | Brain | InceptionV3 | 97.12% |
| Guzmán et | Tumor | ResNet50 | 96.97% |
| al.,(Gómez- | Classi- | InceptionResNetV2 | 96.78% |
| Guzmán et al., | fication | Xception | 95.67% |
| 2023) | (MRI) | MobileNetV2 | 95.45% |
| | | EfficientNetB0 | 90.88% |
| | | Generic CNN | 81.08% |
| Khaliki and | Brain | EfficientNetB4 | 97% |
| Basarslan | Tumor | InceptionV3 | 95% |
| (Khaliki and | Classi- | VGG19 | 96% |
| Başarslan, | fication | VGG16 | 98% |
| 2024) | (MRI) | CNN | 91% |
| Balamurugan | BRATS | DCNN-LuNet | 99.50% |
| and Gnana- | | | |
| manoha- | | | |
| ran(Balamurugan | CE. | AND TE | |
| and Gnana- | | | |
| manoharan, | | | |
| 2023) | | | |
| Fki et al., (Fki | BRATS | DCNN-LuNet | 99.50% |
| et al., 2024) | | | |
| Dhakshnamurthy | Brain | AlexNet | 95.60% |
| et | Tumor | VGG16 | 97.66% |
| al.,(Balamurugan | Classi- | ResNet50 | 96.90% |
| and Gnana- | fication | VGG16-ResNet50 | 98.80% |
| manoharan, | (MRI) | | |
| 2023) | | | |
| Mahmud et | Brain | VGG16 | 71.60% |
| al.,(Mahmud | Tumor | ResNet-50 | 81.10% |
| et al., 2023) | Classi- | CNN | 93.30% |
| | fication | Inception V3 | 80.00% |
| | (MRI) | * | |
| Asiri (Asiri | Brain | FT-ViT | 98.13% |
| et al., 2023) | Tumor | | |
| | MRI | | |
| | Dataset | | |

3 PROPOSED METHODOLOGY

This section presents a hybrid approach for brain tumor classification using ensemble learning to improve diagnostic accuracy. It combines the outputs of complementary deep learning models, including (CNN and ViT), (ViT and FT-InceptionV3), and (FT-VGG16 and FT-InceptionV3), to leverage their strengths in pattern detection, feature extraction, and representation learning. This ensemble learning method reduces overfitting, mitigates biases, and enhances the robustness and generalization of the classification system. The integrated features are processed through a dense layer to finalize tumor classification. In fact, we developed three hybrid architectures: Ens-CNN-ViT, Ens-FT-VGG16-FT-InceptionV3, and Ens-ViT-FT-InceptionV3. These models demonstrated remarkable accuracy in tumor classification, offering a promising new approach for early and accurate brain tumor diagnosis. Figure 1 represents an overview of our proposed models for brain tumor classification. This approach consists of six key steps: (1) Dataset collection, (2) Data preprocessing, (3) Data augmentation, (4) Deep learning-based model development, (5) Ensemble learning-based Hybrid model, and (6) Classification to classify brain tumors into four stages.

3.1 Dataset Description

We used a publicly available "Brain Tumor Classification MRI" dataset from Kaggle (Bhuvaji, 2024) . This dataset contains approximately 3,264 MRI images, classified into four categories: glioma, meningioma, pituitary tumor, and no tumor, as illustrated in Figure 2.

3.2 Dataset Preprocessing

This section describes the basic preprocessing steps taken to prepare the data for model training. The data preparation process involves loading brain tumor images from the specified folders (training and testing), resizing them to 150 * 150 pixels, and then normalizing the image pixel values to the range [0, 1] by dividing them by 255.0, which contributes to faster convergence during training. Finally, we used the function 'Train test split' which allows us to split the data into training and testing sets, with 80% used for training and 20% reserved for testing, allowing for a proper evaluation of the model's performance on unknown data.

3.3 Dataset Augmentation

In our study, we applied data augmentation strategies to enhance model performance by expanding the dataset. The objective is to improve accuracy and opDATA 2025 - 14th International Conference on Data Science, Technology and Applications



Figure 1: The proposed approach process for brain tumor classification.



Figure 2: Sample MRI images.

timize the model's ability to classify brain tumor images. The key techniques used include:

1. The 'ImageDataGenerator' class of Keras: apply alternative transformations to images during training including:

- Rotation: Images can be rotated up to 40 degrees.
- **Translation:** Horizontally and vertically deform images (up to 20% of the large size and height).
- **Shearing:** Apply a cisaillement transformation up to 20%.
- Zoom: Zoom in before or after 20%.
- Horizontal Flip: Images can be returned horizontally.
- Filling: Use appropriate values to represent the new pixels created during the transformations

Figure 3 presents the different Data augmentation techniques.

2. Normalization: images are normalized to avoid pixel values between 0 and 1, which helps the model to approximate faster through training.

3. SMOTE (Synthetic Minority Oversampling Technique): To generate synthetic examples from the minority class to equalize the data game. This helps to mitigate model bias in case of class imbalance.

3.4 Model Architecture Development

The methodology involves the application of both individual deep learning models and ensemble learningbased hybrid architectures for the classification of brain tumors.



Figure 3: Dataset Augmentation.

3.4.1 Convolutional Neural Networks (CNNs)

The proposed CNN for brain tumor classification follows a structured architecture designed for effective feature extraction and classification. It begins with multiple Conv2D layers, applying various filters to capture hierarchical features from input images. These layers are followed by BatchNormalization to stabilize learning and MaxPooling2D to reduce spatial dimensions while preserving key information. The model then flattens the 2D feature map, converting it into a vector that passes through a Dense layer with neurons, followed by a Dropout layer to prevent overfitting. The final classification is handled by a Dense output layer with 4 units and a Softmax activation function, categorizing brain tumors into glioma, meningioma, non-tumor, and pituitary tumor. Figure 4 presents this structured approach.



Figure 4: Architecture of the Proposed CNN Model.

3.4.2 Vision Transformer (ViT)

This study utilizes ViT to classify brain tumors from medical images.

Instead of processing entire images, ViT divides them into patches, which are flattened and treated as token sequences, enabling the model to capture spatial relationships and contextual information effectively. The architecture incorporates multi-head selfattention mechanisms, residual connections, and layer normalization to enhance feature extraction and stabilize training. Position embeddings preserve spatial awareness, while feed-forward networks with ReLU activations, dense layers, and dropout regularization ensure generalization and prevent overfitting. The final classification is performed through a dense layer. ViT's scalability and ability to learn complex patterns make it a strong complement to CNNs, significantly improving diagnostic accuracy in brain tumor classification. Figure 5 provides a detailed overview of the proposed ViT architecture.

3.4.3 Transfer Learning with Pre-Trained Models

In this paper, we explore the use of transfer learning and fine-tuning on pre-trained models (VGG16, VGG19, and InceptionV3) to develop three enhanced architectures: FT-VGG16, FT-VGG19, and FT-InceptionV3. Initially, these models, pre-trained on the ImageNet dataset, leverage their learned feature extraction capabilities, enabling faster convergence and improved performance. Their pre-trained weights remain frozen during the initial training phase to preserve the integrity of the learned features. Subsequently, we apply a transfer learning approach by modifying the architecture: first, the output is flattened before passing through a single dense layer. This structure is then refined by replacing the dense layer with a GlobalAveragePooling2D layer, which reduces the feature map size while retaining important information. Additionally, two fully connected dense layers with ReLU activation and Dropout regularization (0.5) are incorporated to enhance generalization and mitigate overfitting. Finally, a softmax layer classifies images into four tumor categories: glioma, meningioma, pituitary tumor, and no tumor. FT-InceptionV3 integrates these improvements within the InceptionV3 architecture, while FT-VGG16 and FT-VGG19 apply similar modifications to their respective VGG architectures, ensuring optimized accuracy and robustness. Figures 6, 7, and 8 present the architectures of FT-InceptionV3, FT-VGG16, and FT-VGG19, models, respectively.

3.4.4 Hybrid Ensemble Learning

We aim to enhance brain tumor classification performance through a hybrid ensemble learning approach, leveraging the strengths of multiple deep learning models, such as CNNs, ViT, and pre-trained models like FT-VGG16 and FT-InceptionV3 for improved accuracy and robustness. By combining these models, each excelling in different aspects of image processing, our approach aims to capture both local and global features of brain tumor images. CNNs specialize in detecting fine-grained spatial patterns, while ViT excels at modeling long-range dependencies. Additionally, FT-VGG16 and FT-InceptionV3 contribute their expertise in multi-scale feature extraction and advanced representation power.

This hybrid ensemble strategy mitigates the limitations of individual models, improving classification accuracy, reducing overfitting, and ultimately providing a more robust system for accurately classifying brain tumors such as gliomas, meningiomas, pituitary tumors, and non-tumor cases. Figure 9 presents the architectures of the proposed hybrid models Ens-CNN-VIT, Ens-FT-VGG16-FT-InceptionV3 and Ensvit-FT-InceptionV3 based on the application of the ensemble learning.

- Ens-FT-VGG16-FT-InceptionV3 Model

The Ens-FT-VGG16-FT-InceptionV3 model is a hybrid ensemble designed to integrate the strengths of FT-VGG16 and FT-InceptionV3. FT-VGG16 excels at extracting fine-grained spatial features, while FT-InceptionV3 captures multiscale patterns through its inception modules. MRI images are processed separately through both models, and their extracted feature vectors are concatenated into a unified representation. This combined vector is passed through fully connected layers and classified into four tumor categories using a softmax layer.

- Ens-VIT-FT-InceptionV3 Model

The Ens-ViT-FT-InceptionV3 model is a hybrid ensemble architecture that combines the ViT and FT-InceptionV3 models to enhance brain tumor classification accuracy. Specifically, the ViT model captures DATA 2025 - 14th International Conference on Data Science, Technology and Applications



Figure 5: Architecture of the proposed Vit Model.



spatial attention features, while InceptionV3 excels at extracting high-level features. For each input image, the outputs of both models are obtained—ViT generates class probabilities, and InceptionV3 provides feature maps. These outputs are then flattened and concatenated into a single feature vector, which is passed to a dense layer for the final classification. The Adam optimizer is used for training the ensemble model, and the training process is carried out over 30 epochs. This combination of two different models allows the ensemble to take advantage of the complementary information provided by both networks, potentially improving classification accuracy.

- Ens-CNN-VIT Model

The Ens-CNN-ViT model combines CNNs and ViTs to enhance brain tumor classification. CNNs extract local spatial features like edges and textures, and spatial details through their convolutional and pooling layers while ViTs excel at capturing global relationships within the data using their self-attention mechanism and process MRI images as patches, capturing global context and long-range dependencies. MRI images are processed through both models, generating feature maps that are then concatenated into a unified feature vector. This fused representation is passed through fully connected layers, with a softmax layer classifying images into four tumor categories. By combining these two architectures, the model benefits from both local and global feature extraction capabilities, improving classification accuracy and reducing the risk of overfitting.

4 EXPERIMENTAL RESULTS

In this section, we present and analyze the results obtained from the hybrid models developed for brain tumor classification using MRI images.

Performance evaluation was carried out using several standard metrics, including accuracy, recall, F1-score, and confusion matrices. These results demonstrate the effectiveness and robustness of the proposed models.

Table 2 presents the evaluation results for brain tumor classification using the proposed models, including the Proposed-CNN, ViT model, Fine-Tuned VGGNet16, Fine-Tuned VGGNet19, Fine-Tuned InceptionV3, and the new three ensemble learning hy-



Figure 8: Architecture of the Proposed Fine-Tuned Vgg19 Model.

brid models, which are Ens-CNN-ViT, Ens-VGG16-InceptionV3, and Ens-ViT-InceptionV3 models. This table displays the performance metrics—accuracy, precision, recall, and F1-score—for each model.

As indicated in the table, the Ens-VGG16-InceptionV3 hybrid model achieved the highest overall performance with an accuracy of 99%, followed by FT-InceptionV3, CNN, Ens-ViT-InceptionV3, Ens-CNN-ViT, Vit model, FT-VGG16, and FT-VGG19 with 93%, 92%, 90%, 88%, 83%, 81% and 75% respectively. The Ens-ViT-InceptionV3 model also demonstrated a strong performance with an accuracy of 90%, while the Ens-CNN-ViT model attained an accuracy of 88%. These results highlight the superior performance of the Ens-VGG16-InceptionV3 model, with a notable improvement of 11 % over the Ens-CNN-ViT model and a 9% improvement over the Ens-ViT-InceptionV3 Model. In comparison, the CNN model achieved an accuracy of 92%, while the ViT model reached 83%. Among the fine-tuned pre-trained models, FT-VGG16 and FT-VGG19 performed the least, with accuracies of 81% and 75%, respectively.

Figure 10 illustrates the confusion matrices for the three hybrid ensemble learning models: (A) Ens-ViT-FT-InceptionV3, (B) Ens-FT-VGG16-FT-InceptionV3, and (C) Ens-CNN-ViT. As an example, Figure 10 (B) presents the matrix includes four classes: suspected glioma, suspected meningioma, not suspected tumor, and pituitary. The diagonal elements represent the correct model predictions: 826 correct predictions for suspected glioma, 815 correct predictions for suspected tumor, and 826 correct predictions for suspected pituitary. The off-diagonal elments indicate that there are some misclassifications, but these misclassifications are rare.

Among them, 7 suspected meningiomas were mis-

Table 2: Evaluation of brain tumor classification using the proposed CNN, ViT model, fine-tuned pre-trained models, and three ensemble learning hybrid models.

| Classifier | Accuracy | Precision | Recall | F1- |
|-------------|----------|------------|--------|-------|
| | | | | score |
| CNN | 92% | 92% | 92% | 92% |
| ViT | 83% | 85% | 83% | 84% |
| FT- | 93% | 93% | 93% | 93% |
| InceptionV3 | 7 | | | |
| FT-VGG16 | 81% | 81% | 81% | 81% |
| FT-VGG19 | 75% | 77% | 75% | 76% |
| Ens-CNN- | 88% | 89% | 88% | 88% |
| ViT | | | | |
| Ens-ViT- | 90% | 91% | 90% | 90% |
| InceptionV3 | PUBL | .ICA' | וסוד | VS |
| Ens-VGG16- | 99% | 99% | 99% | 99% |
| InceptionV3 | | | | |

classified as suspected gliomas, and 1 suspected pituitary tumor was misclassified as suspected glioma. Overall, the model showed excellent accuracy and correctly classified most cases with few misclassifications. This strong performance shows that the integrated model (FT-VGG16-FT-InceptionV3) can distinguish these four categories very effectively.

Figure 11, shows the training and validation accuracy and loss curves for the hybrid ensemble model, highlighting the superior performance of Ens-FT-VGG16-FT-InceptionV3, which achieves the highest validation accuracy and lowest loss, indicating its robustness.

5 COMPARATIVE EVALUATION

The primary objective of this study is to propose hybrid deep learning models to enhance the classification of brain tumors from MRI images. These tu-



Figure 9: Architectures of the proposed hybrid models based on Ensemble learning.



Figure 10: Confusion Matrices obtained by (A) Ens-ViT-FT-InceptionV3 (B) Ens-FT-VGG16-FT-InceptionV3 (C) Ens-CNN-ViT.



Figure 11: Training accuracy and loss curves for the Ens-FT-VGG16-FT-InceptionV3 ensemble models.

mors—including glioma, meningioma, pituitary tumor, and healthy brains—pose significant diagnostic challenges due to their often similar characteristics. Our results, summarized in Table 3, demonstrate that the proposed ensemble models, particularly Ens-VGG16-InceptionV3, achieved remarkable performance, attaining 99% accuracy and a 99% F1score. These findings underscore the effectiveness of leveraging multiple architectures to improve classification performance and robustness.

Table 3 provides a comparative analysis of the

performance of our proposed models against previous works using the same Brain Tumor Classification (MRI) dataset. Earlier studies, such as Gómez-Guzmán et al., which employed a combination of generic CNN and InceptionV3 model, achieved an accuracy of 97.12% and an F1-score of 96.59%. Similarly, Zafer Khaliki et al., and Dhakshnamurthy et al., reported comparable results, with both achieving 98% accuracy using VGG16 and VGG16-ResNet50 combination models, respectively. Mahmud et al., reported a lower accuracy of 93.30% and an F1-score of 91% using a standalone CNN. In contrast, our ensemble models significantly outperform these earlier methods, demonstrating the superiority of hybrid approaches in leveraging complementary features for more precise brain tumor classification.

The superior performance of ensemble learning hybrid lies in their ability to combine the feature extraction strengths of different architectures. For example, CNNs excel at detecting local features but may struggle with global relationships, which are effectively captured by Vision Transformers. By integrating these architectures, models like Ens-FT-VGG16-FT-InceptionV3 and Ens-ViT-FT-InceptionV3 capi-

| Authors | Models | Best Model | Accuracy (%) | F1- score |
|--|--|---------------------------------|--------------|--------------|
| Proposed approaches | Ens-FT-VGG16-FT- InceptionV3 Ens-ViT-FT-InceptionV3 Ens-CNN-ViT | Ens-FT-VGG16-FT- InceptionV3 | 99% | 99% |
| Gómez-Guzmán et al.,(Gómez-Guzmán et al., 2023) | Generic CNN and six TL mod- els | InceptionV3 | 97.12% | 96.59% |
| Khaliki and Basarslan(Khaliki and Başarslan, 2024) | EfficientNetB4 InceptionV3 VGG19 VGG16 CNN | VGG16 | 98% | 97% |
| Dhakshnamurthy <i>et</i> <i>al.</i> ,(Dhakshnamurthy et al., 2024) | AlexNet VGG16 ResNet50 VGG16–ResNet50 | VGG16– ResNet50 | 98% | 98% |
| Mahmud et al., 2023) | VGG16 ResNet-50 CNN nceptionV3 | CNN | 93.30% | 91% |

Table 3: Comparative analysis of proposed work with previous works.

talize on both local and global feature extraction, achieving balanced and superior performance.

6 CONCLUSION AND FUTURE WORK

This study proposed a hybrid deep learning approach for brain tumor classification from MRI images, integrating CNNs, ViTs, and fine-tuned pre-trained models (FT-InceptionV3, FT-VGG16, and FT-VGG19). By leveraging ensemble learning, we combined the strengths of different architectures, leading to significant improvements in classification accuracy, robustness, and generalization. The proposed hybrid models-Ens-FT-VGG16-FT-InceptionV3, Ens-ViT-FT-InceptionV3, and Ens-CNN-ViT-demonstrated superior performance compared to standalone models and previous studies. Notably, Ens-FT-VGG16-FT-InceptionV3 achieved the highest accuracy and F1-score of 99%, underscoring the effectiveness of hybrid architectures in medical imaging applications. These findings highlight the potential of AI-driven solutions to enhance early detection and classification of brain tumors, minimizing diagnostic delays and improving patient outcomes. Future research will focus on enhancing model interpretability through Explainable AI (XAI), expanding datasets to improve generalization across diverse populations, and optimizing computational efficiency for real-time clinical deployment. This study reinforces the transformative impact of deep learning in medical diagnostics, paving the

way for more precise, scalable, and reliable brain tumor classification systems.

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