

BRAIN TUMOR DETECTION SYSTEM

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Abstract - Detecting brain tumors plays a vital role in medical image analysis crucial for timely diagnosis and treatment planning. An inventive project is introduced here, merging Convolutional Neural Networks (CNNs) with machine learning to automate brain tumor detection. By utilizing Python libraries like TensorFlow, a comprehensive framework is built, integrating CNN-based feature extraction with conventional machine learning algorithms for binary tumor classification. The methodology includes pre-processing MRI scans, extracting significant features using a CNN structure, and inputting these features into machine learning classifiers to identify the presence or absence of a brain tumor. Extensive experimentation is conducted on a diverse dataset containing MRI images of brains with and without tumors to assess the performance of various CNN architectures and machine learning models. The results exhibit promising accuracy and efficiency in tumor detection tasks, with the developed framework achieving remarkable sensitivity and specificity rates. This system holds substantial potential in aiding healthcare professionals in precise diagnosis and treatment planning, ultimately enhancing patient outcomes in neuro-oncology.

Key Words: Tumor, CNN, MRI, accuracy, detection

1. INTRODUCTION

Among the various organs present in the human body, the brain particularly stands out as being the most crucial. An issue commonly seen leading to dysfunction in the brain is the development of a brain tumor, characterized by the uncontrolled growth of excess cells. These abnormal cells tend to consume essential nutrients that healthy brain cells and tissues require, ultimately resulting in brain failure. At present, doctors rely on manual examination of MRI scans to detect and evaluate brain tumors, a method known for its susceptibility to inaccuracies and inefficiencies. Brain cancer, being a severe and life-threatening condition, claims numerous lives due to delayed or incorrect diagnoses. The primary goal behind detecting brain tumors lies in enabling early diagnosis and treatment. This project aims to develop an automated system capable of identifying the presence of brain tumors in MRI scans. By utilizing CNNs, this system processes MRI images to determine the existence of a tumor, offering a reliable diagnostic tool for healthcare professionals. Such an automated system presents substantial advantages

within clinical settings, assisting doctors in providing quick and precise diagnoses. By concentrating on the binary classification task—determining whether a tumor is present or not—this initiative strives to improve early detection, hence enhancing treatment outcomes and patient survival rates in neuro-oncology.

1.1 Motivation

Current techniques for identifying brain tumors heavily depend on the manual analysis of MRI scans by radiologists and healthcare experts. While this traditional method is somewhat effective, it comes with its own set of. These include the risk of human mistakes, subjective judgment, and the considerable time needed for a thorough evaluation. The manual assessment process may result in inconsistent findings, especially considering the intricate nature of brain tumors and their subtle display in imaging. Furthermore, the growing number of MRI scans in medical setups can overwhelm healthcare providers, causing delays in diagnosing and treating patients. Several current systems integrate basic image processing methods and computer-aided diagnosis (CAD) tools to support physicians. However, these systems often lack the advanced features necessary to precisely and effectively identify tumors.

One common method to detect brain tumors is Magnetic Resonance Imaging (MRI). However, the manual interpretation of MRI scans by radiologists can be time-consuming and subjective. This can lead to errors or delays in diagnosis. This project aims to develop an automated system using computer-based methods to identify brain tumors in MRI images. By utilizing Convolutional Neural Network (CNN) algorithms, the system can efficiently analyze MRI scans to determine the presence of a tumor. The process involves several stages: image preprocessing, feature extraction, and classification. During image preprocessing, MRI scans are enhanced to make it easier to identify relevant features. Feature extraction focuses on identifying significant characteristics within the images that indicate the presence of a tumor. In the classification stage, neural network techniques are utilized to determine if a tumor is present. This innovative approach aims to streamline the detection process and improve diagnostic accuracy in identifying brain tumors using MRI technology.

1.2 Objectives

The proposed system for detecting brain tumors aims to achieve several important goals to enhance the diagnostic process:

- **Early Detection:** Focus on finding brain tumors early to improve patient outcomes.
- **Enhanced Accuracy:** Improve diagnostic accuracy using advanced machine learning techniques.
- **Binary Classification:** Develop a reliable system for distinguishing between tumor and non-tumor cases.
- **Seamless Integration:** Ensure smooth integration into existing clinical workflows.
- **Ongoing Improvement:** Keep performance current with the latest advancements in medical imaging and machine learning.
- **Collaborative Decision Support:** Provide comprehensive diagnostic insights and recommendations through collaborative decision-making.
- **Multi-Modal Imaging Data:** Combine multiple types of imaging data for a more thorough analysis.
- **Scalability and Adaptability:** Make sure the system can handle large amounts of data and adapt to technological advancements.

2. RELATED WORKS

In [1], image processing is crucial in identifying and diagnosing brain tumors through MRI and CT scans. The process starts with preprocessing steps like reducing noise, normalizing intensity, registering images, and enhancing them to make them suitable for analysis. Once the preprocessing is done, segmentation comes into play to outline tumor regions using various algorithms such as thresholding, region growing, level sets, active contours, and graph cuts. Features are then extracted to describe the tumors using statistical, shape, and spatial characteristics necessary for classification. Different machine learning algorithms like support vector machines (SVMs), random forests, and artificial neural networks (ANNs) are utilized for tumor classification. The performance of these algorithms is assessed using metrics like sensitivity, specificity, accuracy, and area under the curve (AUC).

In [2], the article centers on the early classification of brain tumors using MRI with the help of deep learning (DL) and transfer learning (TL) methods. By utilizing an automated framework, there is a reduction in the necessity for extensive human involvement. This framework follows a structured process involving data preprocessing, data enhancement, feature extraction, and classification. Various models like Xception, NasNet Large, DenseNet121, and InceptionResNetV2 are utilized, with their

performance being assessed using metrics such as accuracy, sensitivity, precision, specificity, and F1-score. The study accentuates the significance of timely detection and exhibits the remarkable efficiency of the Xception model when coupled with the ADAM optimizer. The research makes a valuable contribution to brain MRI image analysis by refining preprocessing techniques, optimizing data augmentation strategies, and employing cutting-edge DL models for the precise identification of brain tumors.

In [3], the proposed system offers a thorough examination of studies on identifying and classifying brain tumors from MRI scans using deep learning models. It underscores the significant impact of deep learning in medical imaging, especially with convolutional neural networks capturing key tumor features. The study delves into an extensive review of deep learning techniques, discussing their advantages and disadvantages. It explains how deep learning models leverage MRI data to improve tumor detection accuracy. This paper is a valuable guide to current trends and future possibilities in brain tumor diagnosis with deep learning models.

In [4], the proposed system introduces a dual-module approach for detecting brain tumors. In the first module, MRI images are enhanced using adaptive Wiener filtering, neural networks, and independent component analysis to improve clarity and contrast. The second module utilizes Support Vector Machines (SVM) for segmenting and classifying tumors, targeting various types of brain tumors. This method strives to generate clearer MRI images and enhance classification accuracy, contributing to improved clinical diagnoses' reliability. The study validates these techniques on the CE-MRI image database, showcasing significant potential in enhancing the quality of MRI images and the precision of tumor detection endeavors.

In [5], the study discusses the complex challenge of categorizing brain tumors using computer-aided diagnostics, particularly focusing on magnetic resonance imaging (MRI). A method is proposed that emphasizes multi-level feature extraction and combination for early detection due to the diverse nature of tumor cells. By utilizing two well-known deep learning models, Inception-v3 and DenseNet201, two different scenarios were assessed for tumor detection and classification. Firstly, features from various Inception modules are extracted, combined, and classified using a SoftMax classifier. The second scenario entails extracting features from different DenseNet blocks, combining them, and employing a SoftMax classifier for classification as well. By applying these techniques to a publicly accessible three-class brain tumor dataset resulted in impressive testing accuracies of 99.34 percent and 99.51 percent with Inception-v3 and DenseNet201, correspondingly. These results highlight the superior performance of the proposed feature combination approach compared to existing deep learning and machine

learning methods in precisely diagnosing brain tumors through medical imaging.

3. METHODOLOGY

3.1 Overview

The system proposed comprises five main modules. Firstly, the Dataset module, then Pre-processing, followed by Splitting the data, Building the CNN model, training the Deep Neural network for epochs, and finally classification.

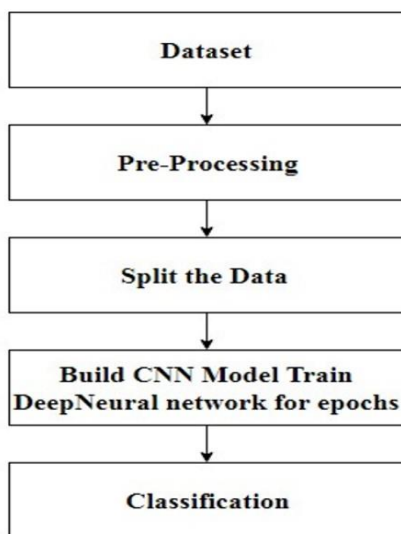


Fig-1: Proposed System

In the dataset module, multiple MRI images can be taken with one as the input. During pre-processing, the image is encoded with labels and resized accordingly. When splitting the data, 80 percent is assigned to Training Data and 20 percent to Testing Data. Further steps involve building the CNN model followed by training the deep neural network for epochs. The images are then classified as either Tumor detected or no tumor. In cases where a tumor is detected, it returns as such while in those without tumors, it returns as no tumor detected.

3.2 Working of CNN Model

A Convolutional Neural Network (CNN) works with images and multiple layers to extract and understand features. Filters in Convolution 2D layers help create maps to display spatial patterns. MAX Pooling2D layers simplify things by shrinking these maps, grabbing the highest values from various areas. Dropout layers help with generalization by turning off random neurons during training.

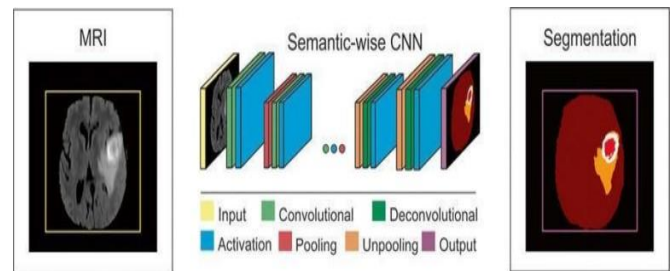


Fig-2: Working of CNN model

Flatten layers get the data ready for Dense layers, linking neurons through all levels for more complex feature learning. Activation layers bring in non-linearity to assist in spotting intricate patterns. The model gets fine-tuned with binary cross-entropy loss and the Adam optimizer for flexible learning rates and effective convergence. The SmallerVGGNet model is utilized in this implementation, which serves as a more compact version influenced by the VGGNet architecture. The primary goal is to decrease computational complexity while retaining essential structural elements. The model is comprised of several crucial components:

- 1) Input Layer: This layer establishes the shape of the input.
- 2) Convolutional Layers: Three sets of convolutional blocks are included:
 - a) Each block consists of Conv2D layers, followed by Activation (ReLU), Batch Normalization, Max-Pooling2D, and Dropout.
- 3) Fully Connected Layer: A Dense layer with 1024 units is present, followed by Activation (ReLU), Batch Normalization, and Dropout.
- 4) Output Layer: This layer features a Dense layer with a sigmoid activation function tailored for binary classification.

This design allows the SmallerVGGNet model to efficiently learn and extract image features, making it suitable for tasks such as binary classification of brain tumor images. The data undergoes preprocessing with various augmentations to enhance model robustness, and the learning rate adjusts dynamically using the ReduceLROnPlateau callback to improve model performance.

3.3 Working of SmallerVGGNet

The specialized convolutional neural network, SmallerVGGNet, has been designed to handle 48x48 RGB images effectively. Unlike VGG16, it opts for fewer convolutional layers and filters to reduce computational requirements while still maintaining spatial intricacies. This is made possible through consistent configurations of

stride and padding. Beginning with 3x3 convolutional filters, the network aims to uncover detailed spatial features such as edges and textures. Following each convolutional layer, max-pooling steps are implemented to decrease the size of feature maps using 2x2 windows with a stride of 2. This approach not only enhances computational efficiency but also helps in preventing overfitting.

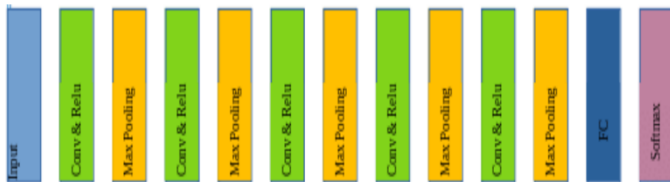


Fig-3: SmallerVGGNet Architecture

Following the convolutional and pooling layers, SmallerVGGNet combines simplified fully connected (FC) layers with ReLU activation functions. This integration aims to bring in non-linearity and support high-level feature learning. Each FC layer includes Batch Normalization to ensure stability and Dropout to prevent overfitting. The network ends with a dense output layer employing sigmoid activation, designed for precise binary classification tasks like identifying brain tumors. By excluding Local Response Normalization, SmallerVGGNet enhances memory usage and computational speed while maintaining classification accuracy. This characteristic makes it ideal for scenarios where computational resources are limited yet accurate image analysis is crucial.

4. SYSTEM DESIGN

4.1 System Overview

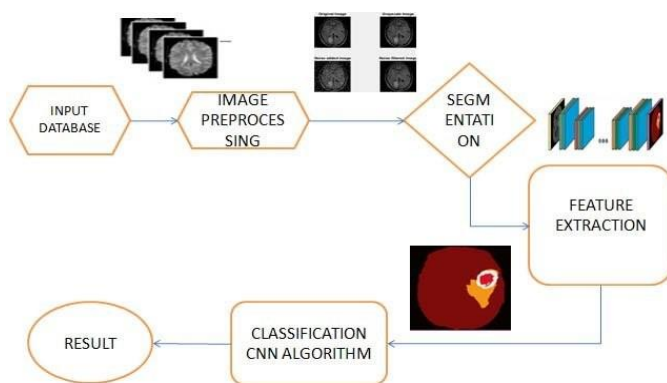


Fig-4: System Architecture

The diagram of the system’s architecture shows how operations flow through the Brain Tumor Detection system. It maps out the steps from getting data (datasets

from Kaggle.com) to preprocessing images, segmenting, extracting features, using CNNs for classification, and showing results on the interface. This visual guide helps us understand how each part works together to detect brain tumors effectively and accurately.

The Brain Tumor Detection interface is a big step forward in medical imaging tech to boost finding and treating brain tumors. With top-notch technology and easy-to-use design, it helps medical pros to spot and tackle brain tumors quickly. This technique can raise diagnostic accuracy, better patient results, and push forward medical research, ultimately improving treatment plans and patient results.

- 1) Input: The Brain Tumor Detection system relies on datasets from Kaggle.com, a well-known platform for accessing and working on datasets. These datasets are carefully selected to contain various medical imaging data essential for training and validating the brain tumor detection model. Through utilizing these datasets, the system guarantees that the model is strong and able to accurately assess a broad spectrum of brain tumor cases. This improves its dependability and efficiency in clinical settings.
- 2) Image Processing: The preparation of images before analysis processing is important to enhance quality. Techniques like noise reduction, contrast enhancement, resizing, color correction, segmentation, and feature extraction are used in sequence. Noise reduction helps clarity while resizing color correction maintains consistency. Segmentation feature extraction isolates tumor features for accurate analysis by the classification model.
- 3) Segmentation: Segmentation represents a crucial stage in digital image processing. It involves dividing an image into various distinct regions or segments, depending on pixel traits. In the realm of brain tumor identification, segmentation plays a key role in pinpointing and separating tumor areas from normal brain tissue. This methodology allows for the detailed outlining and examination of tumor attributes like dimensions, appearance, and position. Through precise segmentation of tumor zones, the system boosts diagnostic precision and streamlines the planning of specialized treatments for individuals dealing with brain tumors.
- 4) Feature Extraction: The process of feature extraction entails capturing and transforming raw image data into numerical features that preserve vital information for analysis. This

conversion turns pixel-based data into significant features that machine learning algorithms, specifically Convolutional Neural Networks (CNNs), can process. Techniques for feature extraction encompass edge detection, texture analysis, and statistical measurements. These methods extract discriminative features that signal the presence or absence of tumors. These features play a crucial role in training the classification model, allowing it to differentiate between images with tumors and those without tumors accurately.

- 5) Classification: The task is to categorize data with Convolutional Neural Networks (CNNs), which is a custom deep learning design for image sorting duties. The CNN structure deciphers the gathered traits across many neuron layers, mastering hierarchy levels of image features. This education process helps the design to split input images into two sections: "Tumor Spotted" or "No Tumor Spotted." With the usage of CNNs, the solution attains strong and trustworthy sorting outcomes, aiding healthcare practitioners in forming wise judgments founded on precise brain tumor spotting.
- 6) Result Display: The Brain Tumor Detection interface features user-friendly components to showcase classification results instantly. By clicking the "Upload Image" button, users kickstart the analysis procedure. Afterward, when users click on the "Predict" button, the system processes the uploaded image using a trained CNN model. Following this step, the interface promptly reveals the classification outcome, determining if a brain tumor is identified or not. This smooth integration of image upload, processing, and result presentation enriches the user experience and streamlines decision-making for medical experts.
- 7) User Interaction: The interface, designed with usability in mind, focuses on intuitive navigation and interaction for users. It includes easily understandable features like image upload and prediction buttons that streamline the process for medical professionals and researchers. This design promotes efficiency in accessing and interpreting brain tumor detection results, aiding in timely clinical decisions and enhancing user satisfaction.
- 8) Impact and Benefits: Brain Tumor Detection interface offers significant benefits in clinical practice and medical research. By diagnostic accuracy, it allows for early detection and intervention in brain tumor patients, ultimately leading to better treatment outcomes and patient

care. Additionally, the interface supports progress in medical imaging technology and brain tumor research by providing data-driven insights and fostering collaborative research efforts.

Overall, the Brain Tumor Detection interface represents a transformative tool in neurology and oncology contributing to healthcare advancements through innovation and technological integration.

5. RESULTS

5.1 Performance Metrics

To validate our proposed brain tumor classification model, we utilize standard performance metrics: accuracy, precision, recall, and F1. These metrics offer a thorough evaluation of the model's capacity to differentiate between tumor and non-tumor cases in medical imaging data.

- **Accuracy:** Reflects the overall correctness of the model's predictions across all classes. It is the ratio of correctly predicted instances to total instances.

$$\text{Accuracy} = \frac{\text{TP} + \text{TN}}{\text{TP} + \text{FN} + \text{FP} + \text{TN}}$$

True Positives (TP), True Negatives (TN), False Positives (FP), and False Negatives (FN) are obtained from the confusion matrix. A 97 percent accuracy implies that the model accurately classified 97 out of every 100 instances.

- **Precision:** Determines the proportion of correctly predicted positive cases (tumors) out of all predicted positive cases.

$$\text{Precision} = \frac{\text{TP}}{\text{TP} + \text{FP}}$$

- **Recall:** Shows the proportion of correctly predicted positive cases (tumors) out of all actual positive cases.

$$\text{Recall} = \frac{\text{TP}}{\text{TP} + \text{FN}}$$

- **F1-score:** A harmonic mean of precision and recall, providing a balanced metric between the two.

$$F1 = 2 \times \frac{\text{Precision} \times \text{Recall}}{\text{Precision} + \text{Recall}}$$

5.2 Confusion Matrix

The confusion matrix details the model's predictions versus actual outcomes, aiding in comprehending the distribution of misclassifications:

Table-1: Confusion Matrix

	Predicted: No tumor	Predicted: Tumor
Actual: No tumor	326	5
Actual: Tumor	11	269

This matrix reveals that the model correctly classified 326 'no tumor' cases and 269 'tumor' cases but also misclassified 5 'no tumor' cases as 'tumor' and 11 'tumor' cases as 'no tumor'. The confusion matrix's overall structure aids in visualizing the model's performance in specific classes and identifying areas for enhancement.

5.3 Performance Metric Table

The summarized performance metrics for our model are presented in the table below, offering a holistic evaluation of its ability to accurately classify both "tumor" and "no tumor" instances.

Table-2: Performance Metric Table

	Precision	Recall	F1- Score	Support
No tumor	0.97	0.98	0.98	331
Tumor	0.98	0.96	0.97	280
Accuracy	0.97			611

The analysis of the performance metric table is summarized below:

- **Precision:** Precision for "no tumor" is 0.97 and for "tumor" is 0.98, indicating correct predictions 97% and 98% of the time respectively.
- **Recall:** Recall for "no tumor" is 0.98 and for "tumor" is 0.96, signifying accurate identification of 98% and 96% actual cases respectively.
- **F1-Score:** The F1-Score combines precision and recall into a single metric and delivers values of 0.98 for "no tumor" and 0.97 for "tumor", showcasing a balanced high-performing model.
- **Support:** Reflecting the number of actual occurrences for each class in the dataset with 331 instances of "no tumor" and 280 instances of "tumor".

5.4 Visual Results

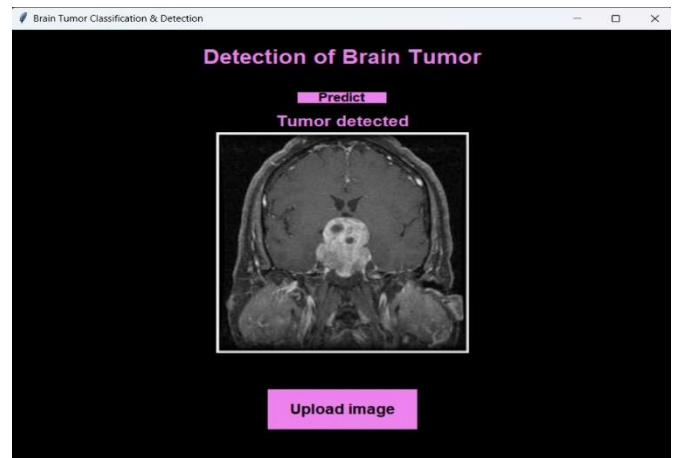


Fig-5: Result showing tumor detected

Visual results in Figure 5 exemplify the model's efficacy in brain tumor detection from medical images, displaying correct identifications or misses, illustrating its adeptness in analyzing complex medical imaging data and making precise predictions.

5.5 Evaluation

Our proposed model for classifying brain tumors achieves an impressive accuracy rate of 97 percent, showcasing its strong performance in distinguishing between tumor and non-tumor cases. The high precision scores (0.97 for 'no tumor' and 0.98 for 'tumor') along with recall rates (0.98 for 'no tumor' and 0.96 for 'tumor') emphasize the model's effectiveness in minimizing both false positives and false negatives.

The detailed confusion matrix offers valuable insights into specific misclassifications, highlighting areas where the model could benefit from further enhancements. It's worth noting that despite the complexity of the task, the model only makes a few minor errors by misclassifying a small number of cases.

Further bolstering these results is Figure 5's visual analysis, which vividly showcases the model's performance across various scenarios. The images provide clarity on how the model accurately identifies tumors based on distinct features while also highlighting potential challenges in detection.

6. DISCUSSIONS

6.1 Limitations

The present Brain Tumor Detection System encounters several challenges, despite significant. These challenges include relying on the quality and variety of

training data, the complex nature of interpreting results from deep learning models, and limitations in computational resources that hinder real-time processing capabilities. Another complicated factor is the variability in MRI acquisition protocols among medical institutions, affecting consistency and reliability.

6.2 Future Scope

Looking forward, there are significant opportunities for improving the Brain Tumor Detection System. Future efforts could concentrate on refining neural network architectures, incorporating explainable AI techniques to enhance transparency and trust among healthcare providers, capitalizing on advancements in hardware technology for quicker processing speeds, standardizing MRI acquisition protocols for data consistency, and integrating multimodal data sources for more personalized diagnostics. These improvements aim to enhance overall accuracy, reliability, and user-friendliness in clinical settings, ultimately advancing early detection and treatment outcomes for patients with brain tumors.

7. CONCLUSIONS

Brain tumors present a significant challenge in medical diagnosis due to their complexity and potential impact on patient health. Early and accurate detection is crucial for improving treatment outcomes and survival rates. Magnetic Resonance Imaging (MRI) offers detailed brain images. However, manual interpretation by radiologists is time-consuming and prone to errors. Automated systems utilizing machine learning and deep learning provide high accuracy, consistency, and efficiency, revolutionizing brain tumor detection and classification. The objective of this project is to develop an automated detection system that focuses on early detection, accuracy improvement, binary classification, and automated analysis. The system aims to offer reliable and rapid diagnostic support. It emphasizes seamless integration into clinical workflows and continuous improvement to stay current with advancements in medical imaging and machine learning. Furthermore, the project includes personalized risk assessments and collaborative decision support by integrating multimodal imaging data and patient-specific information. This ensures comprehensive diagnostic insights and recommendations for patients.

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