

Project 3 MSU AI Boot Camp

April 2024

Brain Tumor Classification Using Deep Neural Networks

Brain tumor classification is a critical task in medical imaging, aiding in the diagnosis and treatment planning for patients. Deep Learning, particularly Convolutional Neural Networks (CNNs), has shown remarkable success in this area. Here, we explore two effective design strategies for constructing deep neural networks capable of classifying brain tumors into four distinct types.

Custom CNN Architecture

Design Overview

The Custom CNN Architecture is tailored specifically for the dataset at hand, allowing for a flexible design that can be iteratively refined to meet the unique requirements of brain tumor classification.

Input Layer

- **Description:** The entry point for input images, resized to a uniform dimension (e.g., 224×224 pixels) for consistency.

Convolutional Layers

- **Purpose:** Extract features from images using small kernel sizes (e.g., 3×3 or 5×5) with ReLU activation for introducing non-linearity.

Pooling Layers

- **Function:** Employ max pooling to reduce the spatial dimensions and computational load, improving efficiency.

Normalization Layers

- **Role:** Stabilize learning by applying batch normalization following convolutional layers.

Fully Connected Layers (Dense Layers)

- **Action:** Flatten the output from convolutional and pooling layers, using dense layers for classification. Dropout layers are included to mitigate overfitting.

Output Layer

- **Goal:** Classify images into one of the four brain tumor categories using a softmax activation function.

Key Considerations

- **Iterative Design:** Start simple and increase complexity based on performance and computational limits.
 - **Prevention of Overfitting:** Incorporate dropout layers strategically within the network.
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Transfer Learning with Pretrained Models

Design Overview

Leveraging pretrained models offers a shortcut to achieving high accuracy, especially with a limited dataset, by utilizing the feature extraction capabilities of models trained on extensive datasets like ImageNet.

Pretrained Base Model

- **Core:** Utilize models such as VGG16, ResNet50, or InceptionV3, removing the top layer to adapt them for feature extraction specific to brain tumor classification.

Global Average Pooling (GAP)

- **Strategy:** Reduce spatial dimensions to a single vector per channel with a GAP layer, following the last convolutional layer of the base model.

Dense Layers

- **Mechanism:** Add dense layers with ReLU activation atop the GAP to classify the features. Implement dropout or regularization to avoid overfitting.

Output Layer

- **Objective:** A softmax activation layer for classifying features into four brain tumor types.

Key Considerations

- **Data Augmentation:** Essential for enhancing training data diversity, improving model generalization across varied and complex medical images.
- **Optimization and Loss Function:** Adam or SGD with momentum are recommended optimizers, with categorical crossentropy as the loss function, suitable for multi-class classification.

Selecting the Right Approach

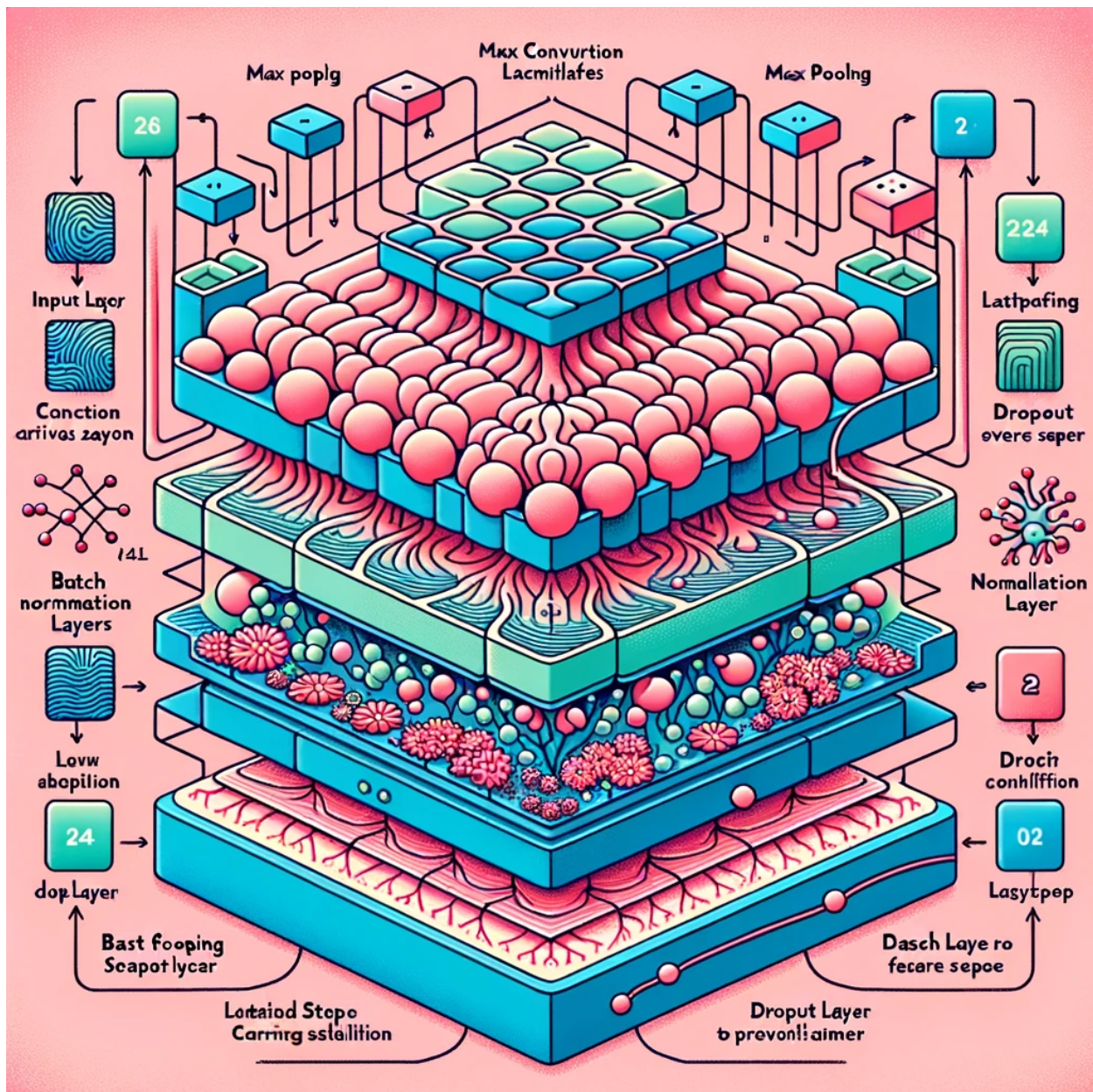
- **Custom CNN Architecture:** Best if you possess the computational resources and domain expertise necessary for developing and refining a specialized model.
- **Transfer Learning with Pretrained Models:** Ideal for scenarios with limited data or when aiming to leverage pre-established patterns from extensively trained models, offering faster training and enhanced accuracy.

Both strategies necessitate meticulous hyperparameter tuning (learning rate, batch size, number of epochs) and ongoing performance evaluation using a validation set to ensure the model's effectiveness and adjust as needed.

Deep Neural Network Diagrams

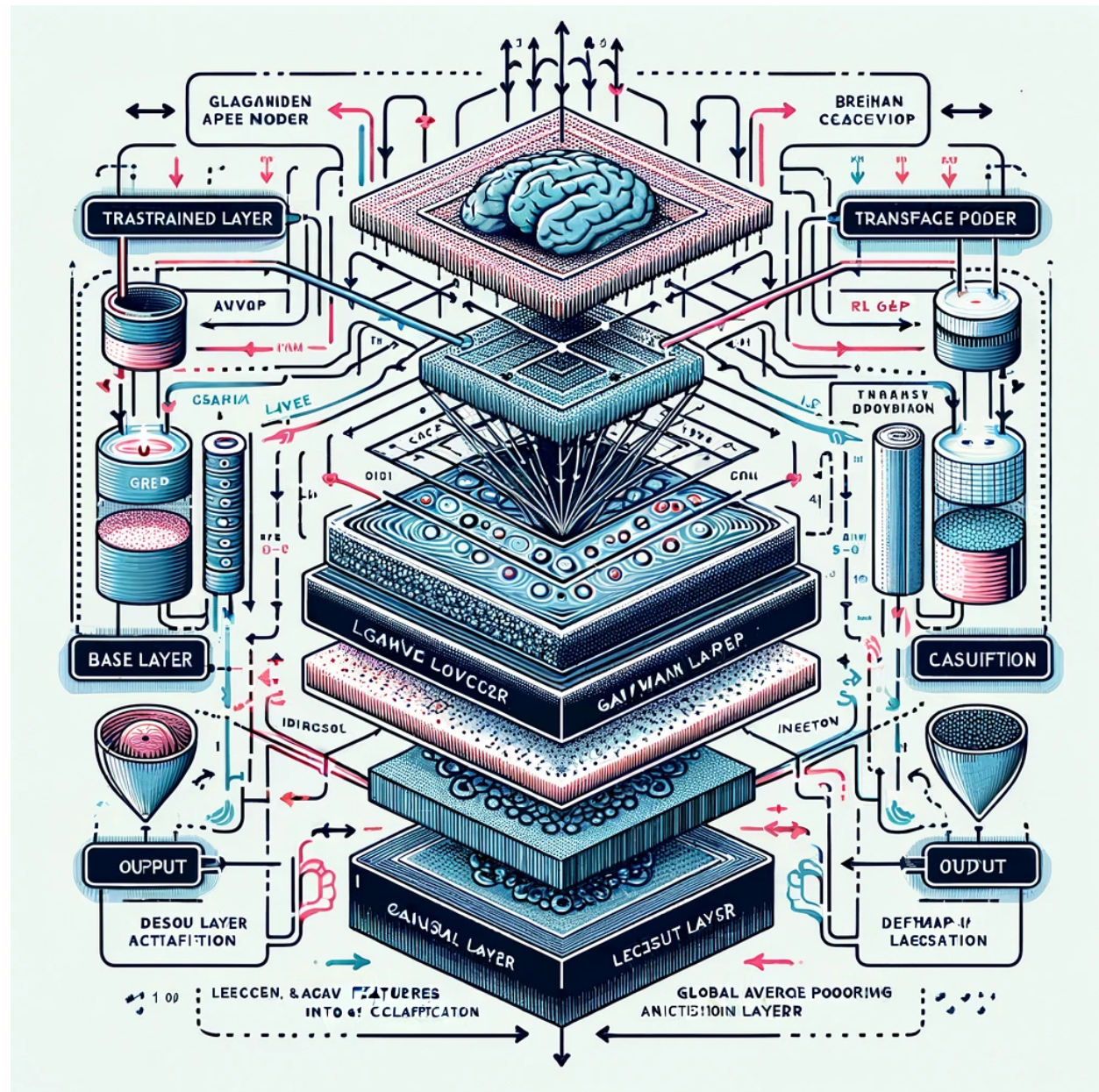
Custom CNN Architecture: This diagram illustrates a custom Convolutional Neural Network (CNN) specifically designed for the task of classifying brain tumors. It highlights the sequential layers including convolutional, max pooling, batch

normalization, and dense layers, culminating in an output layer with softmax activation for classification into four categories.



Transfer Learning Architecture: This diagram showcases a network utilizing transfer learning from a pre-trained base model (like VGG16, ResNet50, or InceptionV3), followed by a global average pooling layer, dense layers with

dropout for classification, and a softmax output layer. This approach leverages learned features from large datasets to improve classification accuracy.



Setting Up a Transfer Learning Architecture with the Brain Tumor MRI Dataset

This guide outlines a comprehensive approach to developing a transfer learning model using the Brain Tumor MRI Dataset available on Kaggle. By leveraging a pretrained model, we can efficiently learn from image data to classify brain tumors with high accuracy.

1. Prepare Your Environment

Before diving into the data and model, ensure your environment is correctly set up:

- **Required Software & Libraries:**
 - **Python:** The core programming language we'll be using.
 - **Libraries:** Make sure to have `tensorflow`, `keras`, `numpy`, `pandas`, and `matplotlib` installed.
 - **Dataset Acquisition:**
 - **Kaggle Dataset:** The Brain Tumor MRI Dataset is available on Kaggle. You'll need a Kaggle account and the Kaggle API installed on your machine. Use the Kaggle CLI for direct dataset download.
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2. Load and Preprocess the Data

Loading the Data

- **Image Libraries:** Utilize `PIL` or `opencv` to load the images.
- **Dataset Segmentation:** If the dataset isn't already divided, split it into training, validation, and test sets.

Preprocessing Steps

- **Resize Images:** Adjust the images to the expected size of the pretrained model (e.g., 224×224 for VGG16).
- **Normalization:** Scale pixel values to a 0-1 range for better model performance.

- **Data Augmentation:** Enhance your model's robustness and reduce overfitting with `ImageDataGenerator` from `keras.preprocessing.image`.
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3. Configure the Pretrained Model

- **Model Selection:** Choose from the pretrained models available in `tensorflow.keras.applications`, such as VGG16, ResNet50, or InceptionV3.
 - **Model Configuration:** Load the chosen model without its top layer (`include_top=False`) and set `input_shape` to match your image size.
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+ 4. Add Classification Layers

- **Global Average Pooling (GAP):** Integrate a `GlobalAveragePooling2D` layer to condense feature maps into a single vector per channel.
 - **Dense Layers:** Insert one or two dense layers to learn complex feature combinations, applying dropout or other regularization techniques to avoid overfitting.
 - **Output Layer:** The final layer should be a dense layer with a softmax activation function, having a neuron for each brain tumor class.
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5. Compile the Model

Configure your model for training:

- **Optimizer:** Use an optimizer like Adam for effective learning.
 - **Loss Function:** Apply `categorical_crossentropy` for multi-class classification.
 - **Metrics:** Evaluate your model with metrics such as accuracy.
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6. Train the Model

Train your model with the `.fit()` method, providing:

- **Training and Validation Data:** Ensure both datasets are included.

- **Epochs & Batch Size:** Set these based on your dataset size and computational resources.
 - **Callbacks:** Use `ModelCheckpoint` and `EarlyStopping` to monitor performance and save the best iteration.
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7. Evaluate the Model

- **Performance Analysis:** After training, use `.evaluate()` to test your model on the unseen test dataset.
 - **Prediction:** Employ `.predict()` to classify new images, further examining the model's predictive capabilities.
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8. Fine-tuning (Optional)

For potential improvements:

- **Layer Unfreezing:** Unfreeze and retrain some of the top layers of the pretrained model with a very low learning rate to refine feature learning on your specific dataset.

This step-by-step guide aims to streamline the development of a transfer learning model for classifying brain tumors using MRI images. By following these instructions, you can harness the power of deep learning to contribute valuable insights into brain tumor diagnosis and classification.