

Identifying Brain Tumors

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Introduction to the problem



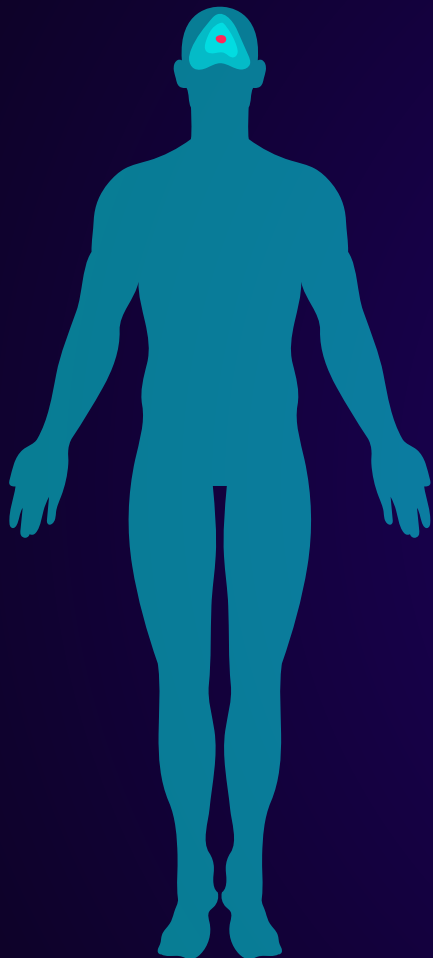
Introduction

- According to the American Brain Tumor Association, approximately 700,000 people in the United States are living with a primary brain tumor, and nearly 80,000 new cases are diagnosed each year .
- Magnetic Resonance Imaging (MRI) is one of the most effective and non-invasive tools for brain tumor detection.
- However, the manual analysis of MRI images by radiologists can be time-consuming.
- Despite advancements in medical technology, the five-year survival rate for patients with malignant brain tumors remains low, at around 36%.
This underscores the importance of early detection, which can lead to more effective treatment options, such as surgery, radiation, and chemotherapy.

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Research Question





Research Question

Can we train a model to use MRI brain scans to reliably predict the presence of no brain tumor vs. presence of a glioma, meningioma or pituitary tumor?

Implications for ML Tool:

- Earlier tumor detection for patients undergoing MRI scans
- Validation support for physicians writing a diagnosis/prognosis
- Improved surgical planning
- Patient-tailored brain tumor management
- Opportunities for researchers to build on tool for more advanced models



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About the Data



Brain Tumor MRI Dataset:




- **Data Sources:**

- Combined image datasets from figshare.com & other similar kaggle projects

- **Data Shape:**

- **7023** human brain MRI images total (including no tumor, meningioma, glioma, pituitary)
 - **1621** images featuring a glioma
 - **1645** images featuring a meningioma
 - **1757** images featuring a pituitary tumor
 - **2000** images featuring no tumor

- **Main Features:**

- Greyscale color // color values
 - 224 x 224 pixels // pixel labels
 - Scan orientation
 - Ensure no individual patient's images are separated across training and test sets
- 

Three types of tumors

Glioma

A type of brain tumor that starts in the glial cells, which are the supportive cells that help protect and nourish nerve cells in the brain.

Pituitary

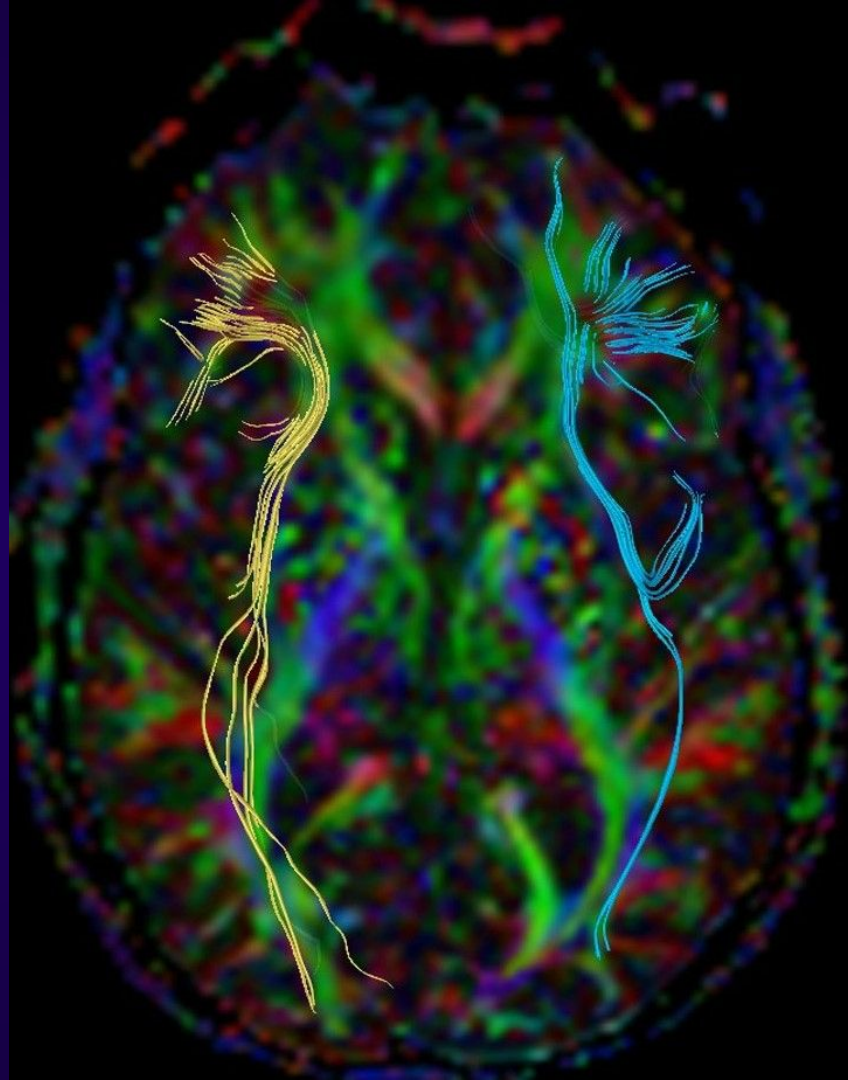
A growth in the pituitary gland, which can affect hormone levels and overall body functions.

Meningioma

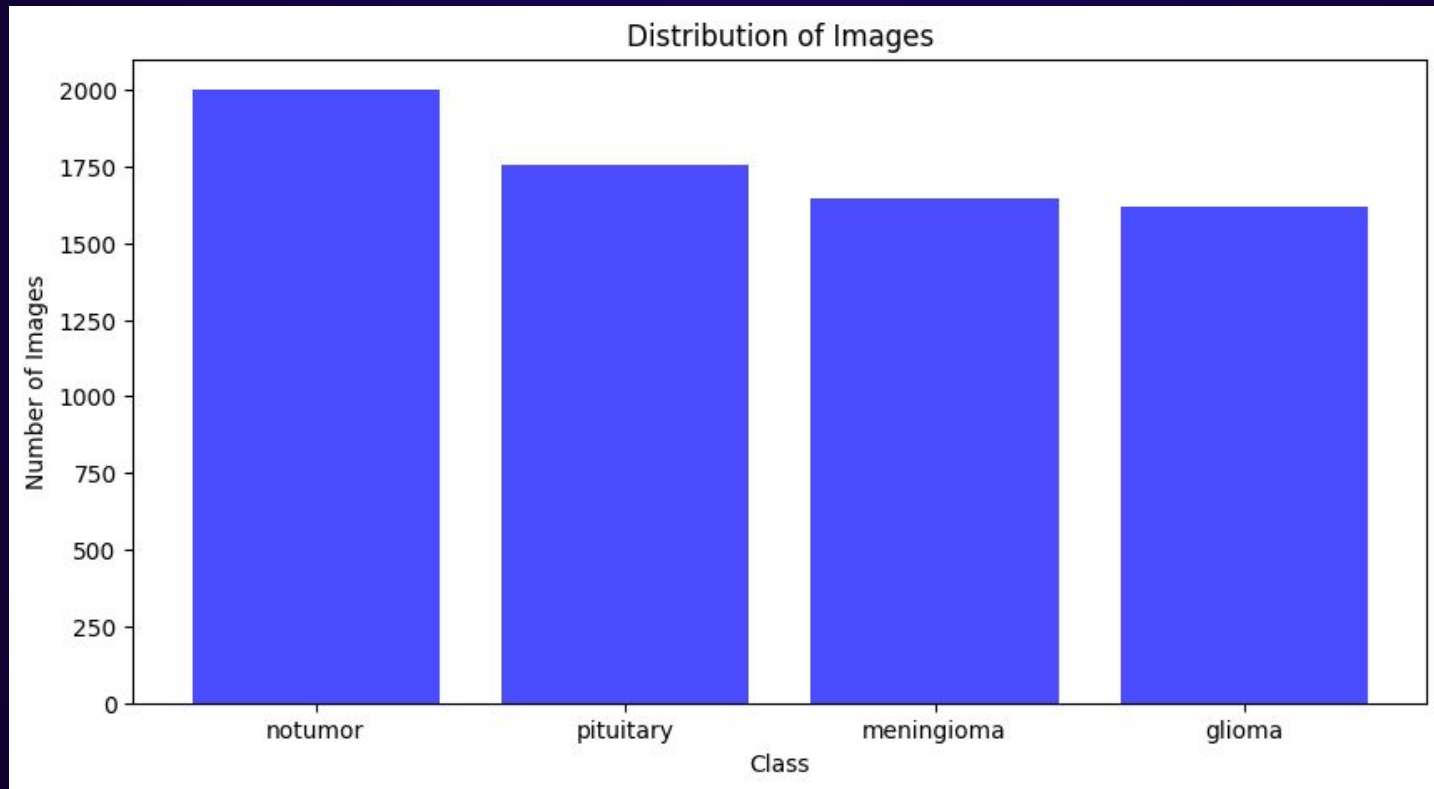
A tumor that forms on the membranes covering the brain and spinal cord, usually benign and slow-growing.

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Exploratory Data Analysis



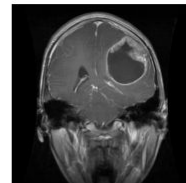
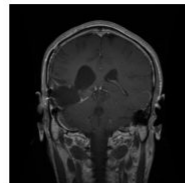
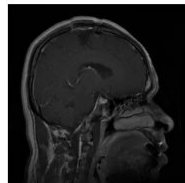
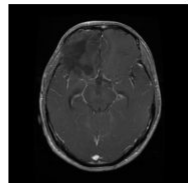
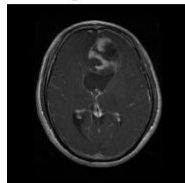
Data Distribution



First Five Images from Each Class (Training Set)

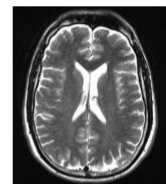
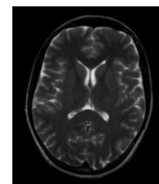
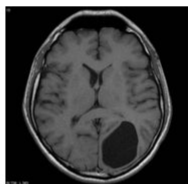
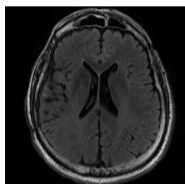
glioma

glioma



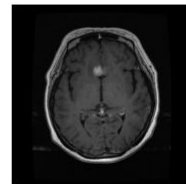
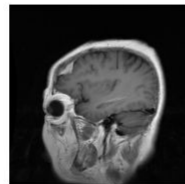
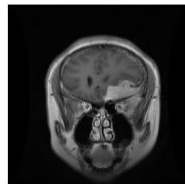
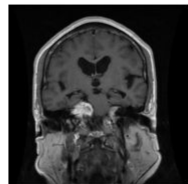
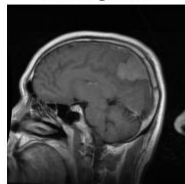
no tumor

notumor



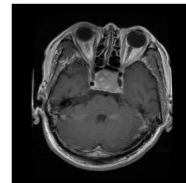
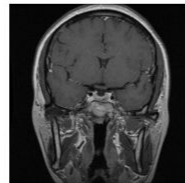
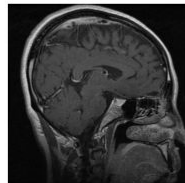
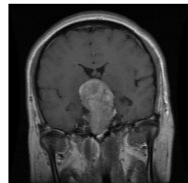
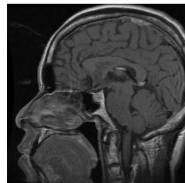
meningioma

meningioma

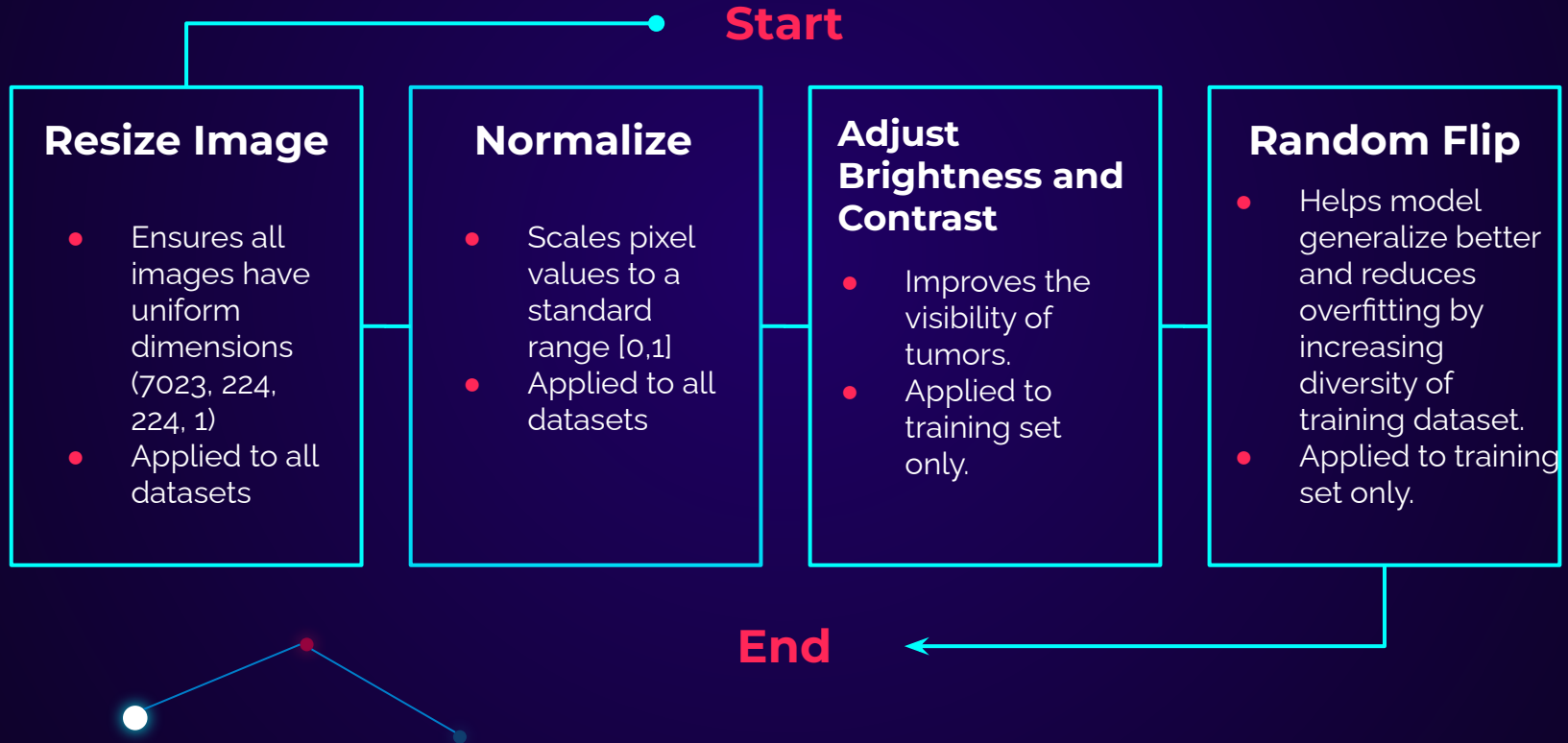


pituitary

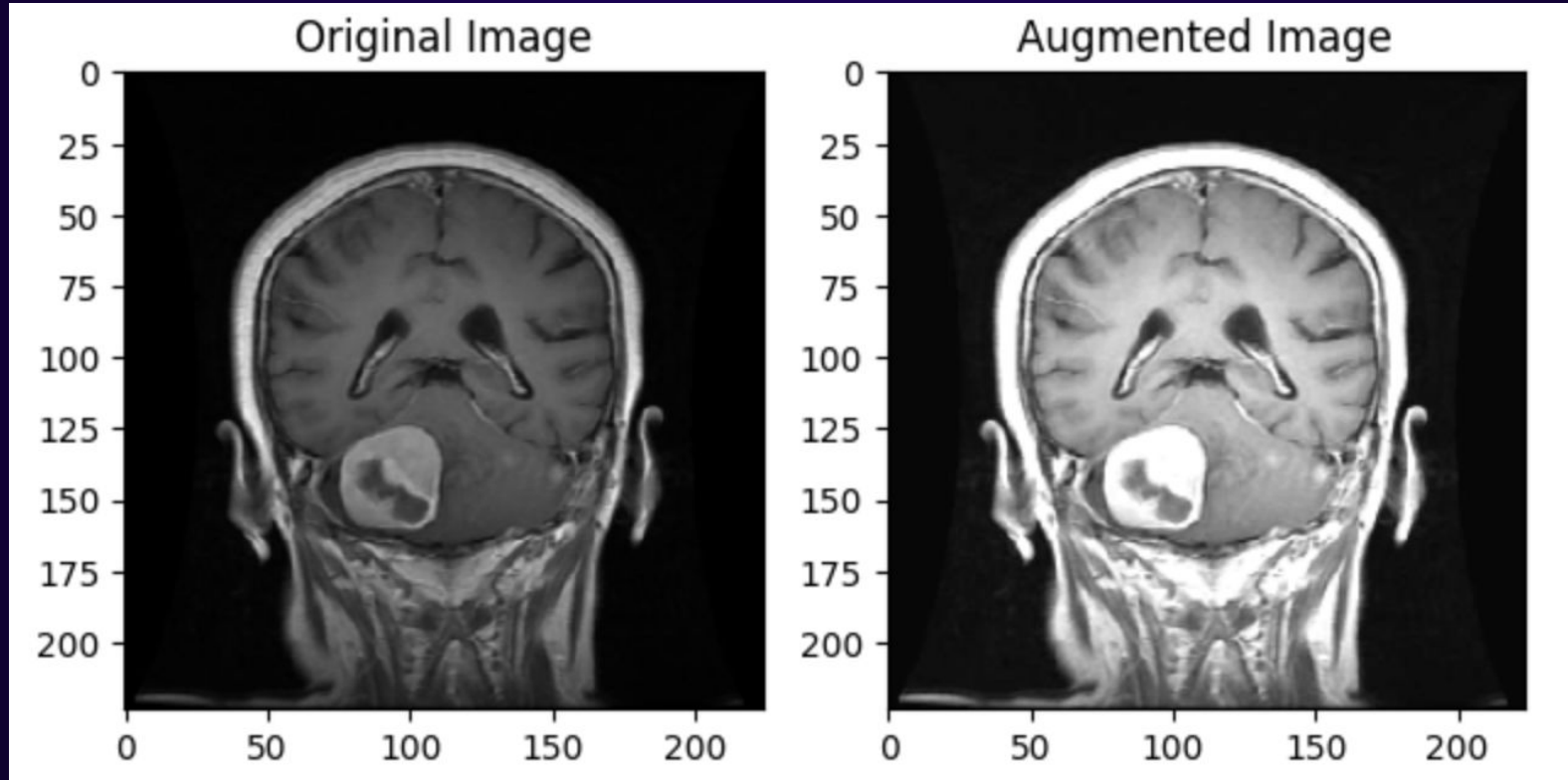
pituitary



Data Preprocessing



Data Preprocessing (Example)





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Prediction Algorithm



Prediction Algorithm:

CNN Model

Feature Generation
- Convolution, Activation
and Pooling layers

Pros
- Well established
techniques
- Interpretable



Transformer Model

Feature Generation
- Self-attention,
Feed-forward,
multi-head and positional
layers

Pros
- Scalability and flexibility
- Recent success with
ViTs



Suggested Hybrid Model

Start with a CNN feature Extractor that would include Convolution Layers and Pooling Layers and then layer Transformer Encoding which would include Positional Encoding and Self-Attention Mechanisms. Lastly, we will include a fully connected layer an an output layer.

Pros: Takes strengths of each model - using localized feature extraction from CNNs with contextual understanding of Transformers.

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Evaluate Results



Baseline Model

- Evaluate performance of our image classification algorithm with a good baseline model appropriate for comparison of performance.
- Use the majority class classifier, which would predict 'no tumor' in each case.
- Defaulting our baseline prediction to 'no tumor' is beneficial since brain tumor diagnosis is rare.

Class	Percentage (%)	Rate of Prevalence in Reality
No Tumor	28%	
Pituitary	25%	
Glioma	23%	
Meningioma	23%	

Evaluating Results

Minimize False Negatives & Maximize Recall

- Recall = positive cases that were correctly identified by the model
- Critical to early & effective treatment. We do not want the model to miss positive cases of cancer.

Multiclass ROC Analysis - Assess Model Discriminatory Ability

- **Approach:** Use One-vs-Rest (OvR) strategy for each tumor class:
 - Class A (no tumor) vs. (Class B + Class C + Class D)
 - Class B (meningioma) vs. (Class A + Class C + Class D)
 - Class C (glioma) vs. (Class A + Class B + Class D)
 - Class D (pituitary) vs. (Class A + Class B + Class C)
- **Metric:** Compute Area Under the Curve (AUC) for each class-specific ROC curve.
- **Micro-averaging:** Average AUC across all classes, weighted by their support.
- **Macro-averaging:** Average AUC independently for each class, then average these values.
- **Benefits:**
 - Each class-specific ROC curve plots the TPR (recall) against FPR at different thresholds
 - Comprehensive evaluation of model's ability to distinguish between tumor types.
 - Visual representation (ROC curves) aids in threshold selection and comparison.
 - Handles class imbalances naturally, crucial for varied tumor prevalence.