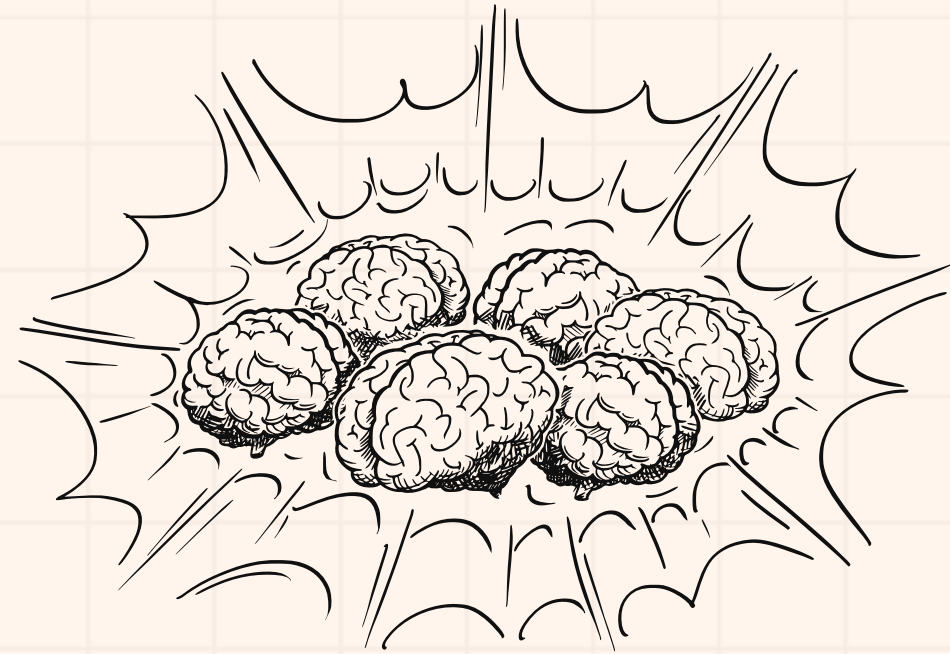


BRAIN TUMOR



Classification & Object Detection
System

TEAM MEMBERS

Sarah Alaa

CNN Model From Scratch

Menna Mahmoud

Applied RESNET Versions

Abdullah Yasser

Applied Effiecent Versions

Nour Yahia

Applied Desnet Versions

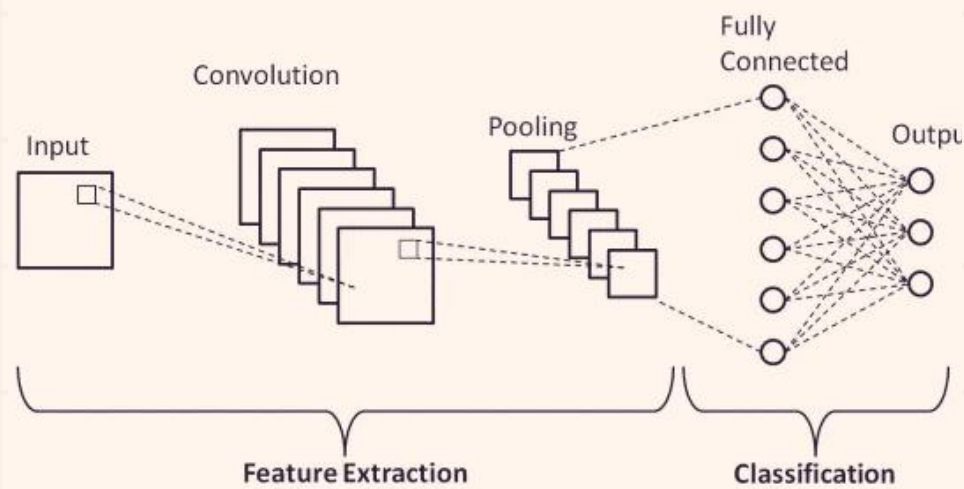
Ahmed Abo ELnaga

Object Detection Model

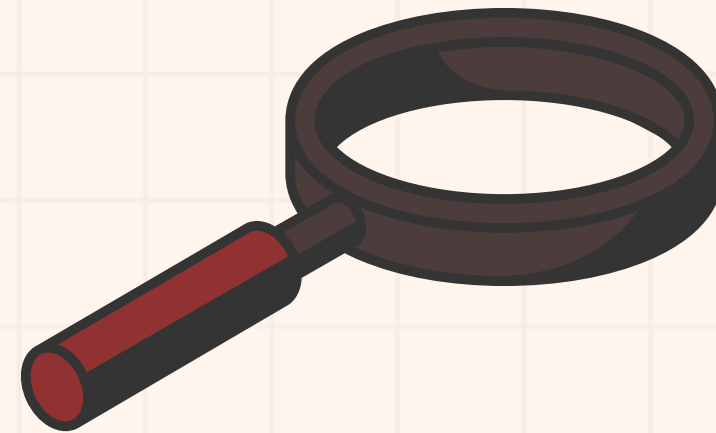
Amr Ibrahim

Model Deployment

PROJECT COMPONENTS



**IMAGE
CLASSIFICATION**



**OBJECT
DETECTION**



**MODEL
DEPLOYMENT**

DATA SELECTION

Brain Tumor MRI Images 44 Classes

collection of T1, contrast-enhanced T1, and T2 magnetic

Images without any type of marking or patient identification, interpreted by radiologists and provided for study purposes.

The images are separated by astrocytoma, carcinoma, ependymoma, ganglioglioma, germinoma, glioblastoma, granuloma, medulloblastoma, meningioma, neurocytoma, oligodendroglioma, papilloma, schwannoma and tuberculoma.

[Artificial intelligence in neuro-oncology: advances and challenges in brain tumor diagnosis, prognosis, and precision treatment](#)

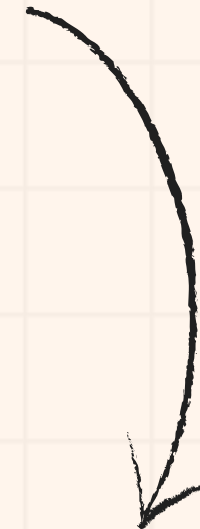


CNN Model From Scratch

ResNet Pretrained Model

CLASSIFICATION

DenseNet Pretrained Model



CNN Model FROM SCRATCH

- Built a deep CNN model from scratch with 4 convolutional blocks, dropout, and batch normalization to classify brain tumor MRI images.
- Used Adamax optimizer, categorical_crossentropy loss, and callbacks (EarlyStopping, ReduceLROnPlateau) for stable and efficient training.

**We tried using pretrained Models
for better performane**

Classification Report:				
	precision	recall	f1-score	support
Astrocitoma	0.90	0.95	0.92	58
Carcinoma	0.96	0.96	0.96	25
Ependimoma	1.00	0.93	0.97	15
Ganglioglioma	1.00	0.83	0.91	6
Germinoma	1.00	1.00	1.00	10
Glioblastoma	1.00	1.00	1.00	21
Granuloma	1.00	0.88	0.93	8
Meduloblastoma	1.00	1.00	1.00	13
Meningioma	0.99	0.92	0.95	88
Neurocitoma	0.96	0.98	0.97	45
Oligodendroglioma	1.00	0.95	0.98	22
Papiloma	0.88	1.00	0.94	23
Schwannoma	0.96	1.00	0.98	47
Tuberculoma	1.00	0.93	0.97	15
_NORMAL	0.96	1.00	0.98	52
accuracy			0.96	448
macro avg	0.97	0.96	0.96	448
weighted avg	0.96	0.96	0.96	448

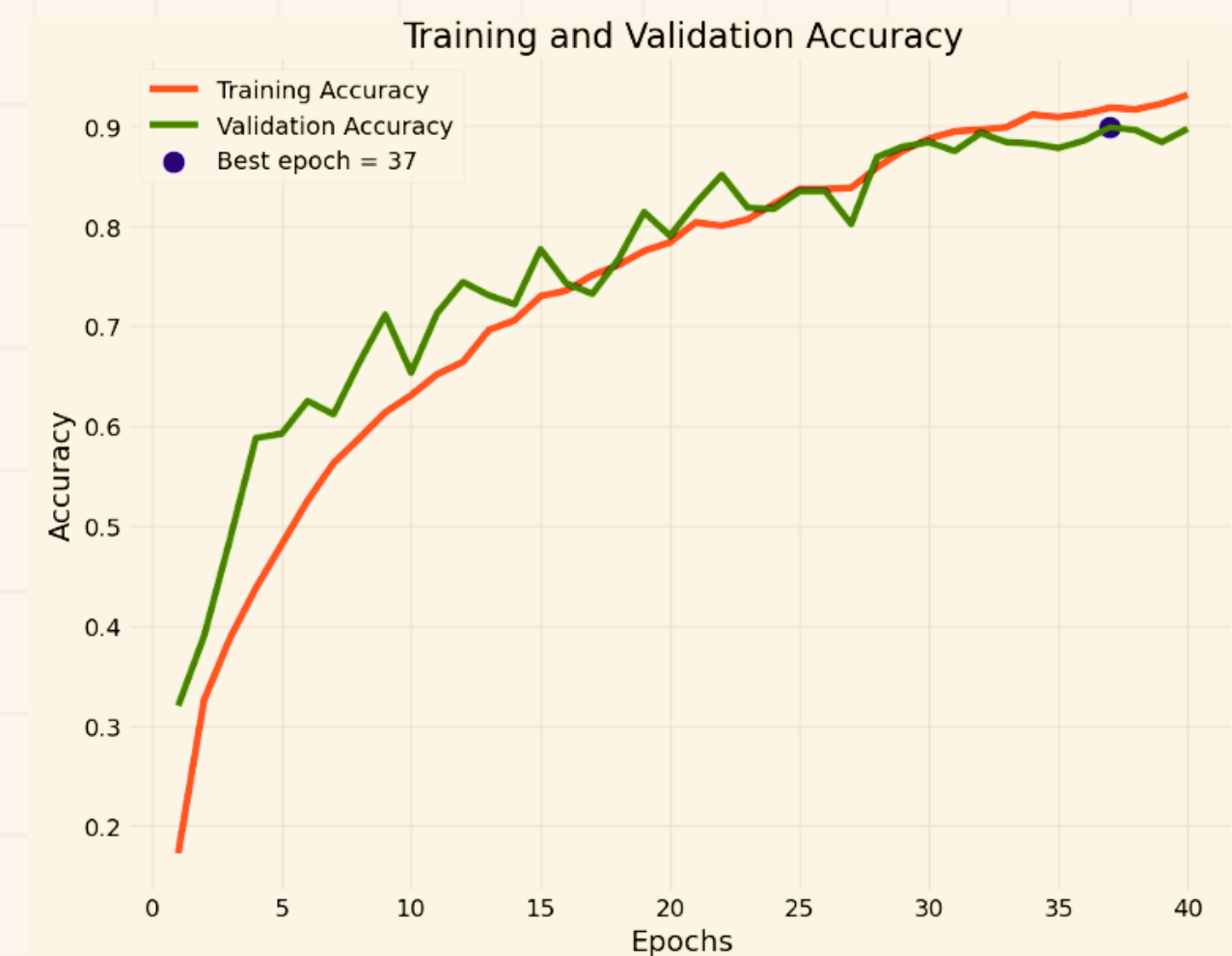
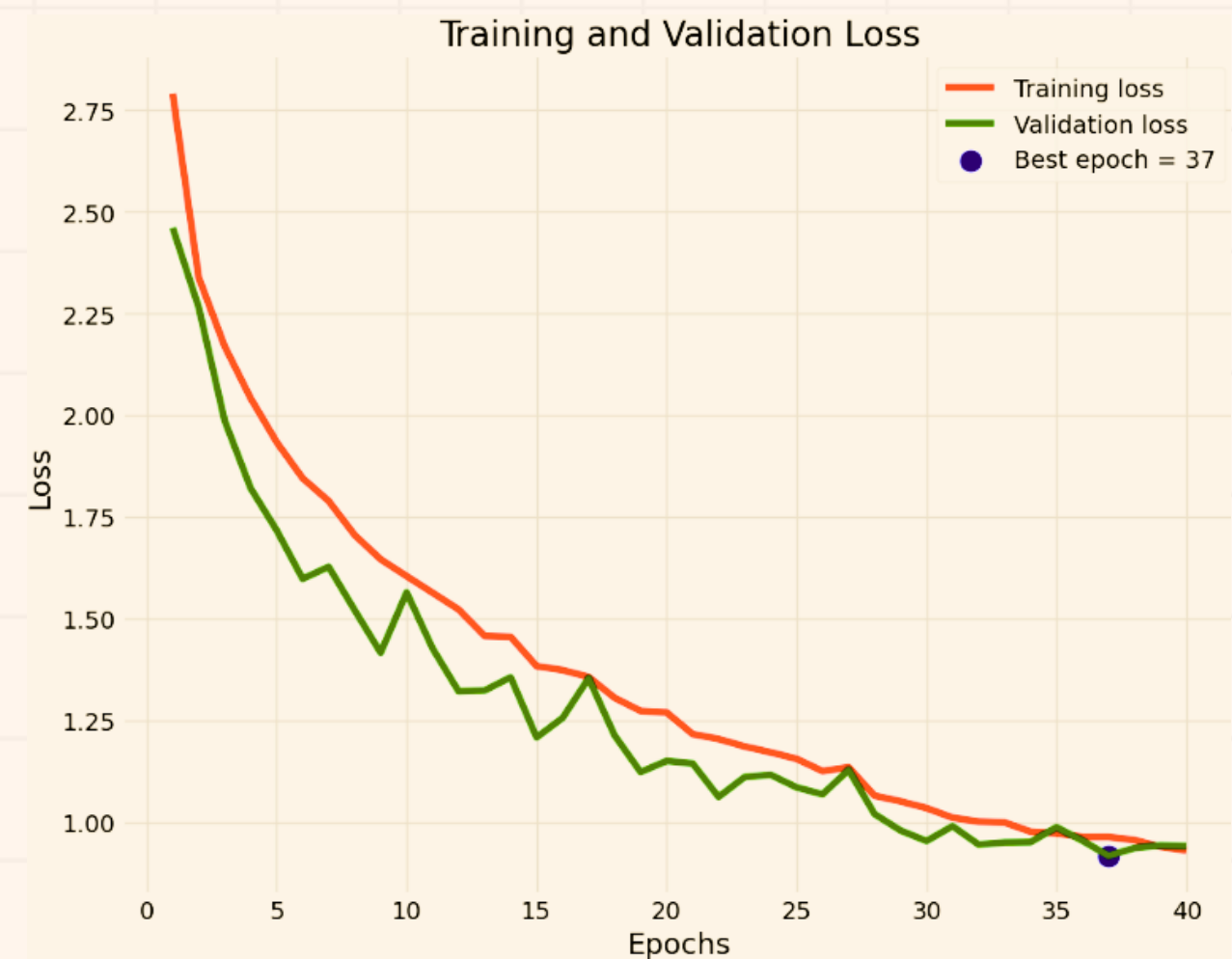
ResNet PRETRAINED MODEL

Explored various versions of the ResNet architecture to identify the most effective one for our classification task.

After testing and fine-tuning models like , ResNet50, and ResNet101... etc.

ResNet50V2 stood out by delivering the best accuracy and overall best performance of resnet versions

**Still not the best Performance Metrics
So we tried Another pretrained Model**



DenseNet

PRETRAINED MODEL

Next, we employed the DenseNet121 & DenseNet169 architecture, leveraging its pre-trained weights on ImageNet. We fine tuned the model to adapt it to our specific classification task.

DenseNet121's design, characterized by dense connections between layers, facilitated efficient feature propagation and reuse, leading to improved performance.

Throughout training, we monitored the model's performance using accuracy and loss metrics, ensuring optimal convergence and generalization

Still not the best Performance Metrics
So we tried Another pretrained Model

Evaluation: DenseNet-121				
Accuracy: 69.07%				
	precision	recall	f1-score	support
Astrocitoma T1	0.87	0.75	0.81	36
Astrocitoma T1C+	0.72	0.66	0.69	47
Astrocitoma T2	0.64	0.40	0.49	35
Carcinoma T1	1.00	0.86	0.92	14
Carcinoma T1C+	0.96	0.96	0.96	23
Carcinoma T2	1.00	0.67	0.80	15
Ependimoma T1	1.00	0.11	0.20	9
Ependimoma T1C+	0.80	0.40	0.53	10
Ependimoma T2	0.00	0.00	0.00	12
Ganglioglioma T1	1.00	0.50	0.67	4
Ganglioglioma T1C+	1.00	0.50	0.67	4
Ganglioglioma T2	1.00	0.40	0.57	5
Germinoma T1	0.50	0.17	0.25	6
Germinoma T1C+	0.75	0.38	0.50	8
Germinoma T2	0.00	0.00	0.00	7
Glioblastoma T1	1.00	0.55	0.71	11
Glioblastoma T1C+	0.94	0.79	0.86	19
Glioblastoma T2	0.67	0.36	0.47	11
Granuloma T1	0.00	0.00	0.00	6
Granuloma T1C+	0.44	0.57	0.50	7
Granuloma T2	0.00	0.00	0.00	4
accuracy			0.69	915
macro avg	0.71	0.54	0.58	915
weighted avg	0.72	0.69	0.67	915

Evaluation: DenseNet-169				
Accuracy: 73.77%				
	precision	recall	f1-score	support
Astrocitoma T1	0.74	0.72	0.73	36
Astrocitoma T1C+	0.78	0.62	0.69	47
Astrocitoma T2	0.56	0.57	0.56	35
Carcinoma T1	0.80	0.86	0.83	14
Carcinoma T1C+	0.96	0.96	0.96	23
Carcinoma T2	1.00	0.67	0.80	15
Ependimoma T1	0.56	0.56	0.56	9
Ependimoma T1C+	0.67	0.40	0.50	10
Ependimoma T2	0.36	0.42	0.38	12
Ganglioglioma T1	0.40	0.50	0.44	4
Ganglioglioma T1C+	1.00	0.75	0.86	4
Ganglioglioma T2	0.00	0.00	0.00	5
Germinoma T1	0.50	0.67	0.57	6
Germinoma T1C+	1.00	0.62	0.77	8
Germinoma T2	0.75	0.43	0.55	7
Glioblastoma T1	0.88	0.64	0.74	11
Glioblastoma T1C+	0.84	0.84	0.84	19
Glioblastoma T2	0.33	0.36	0.35	11
Granuloma T1	0.86	1.00	0.92	6
Granuloma T1C+	1.00	0.57	0.75	7
Granuloma T2	0.00	0.00	0.00	4
accuracy			0.74	915
macro avg	0.69	0.64	0.65	915
weighted avg	0.74	0.74	0.73	915

EffiecentNet

PRETRAINED MODEL

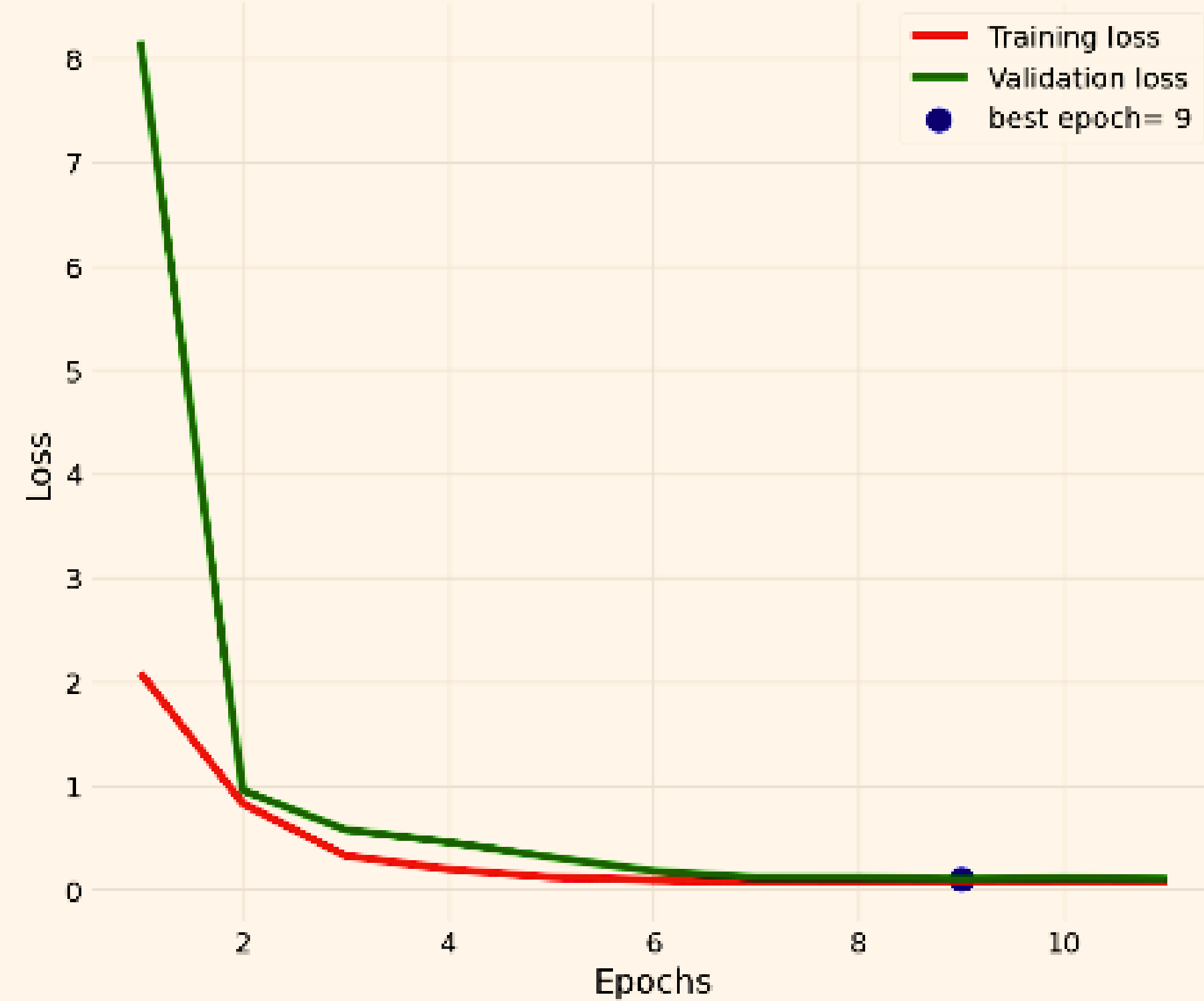
Applied the EfficientNet versions to leverage its optimized scaling capabilities for image classification.

After evaluating different variants, EfficientNetB5 was selected as the final model due to its excellent balance of accuracy and computational efficiency, achieving the best results among the tested versions.

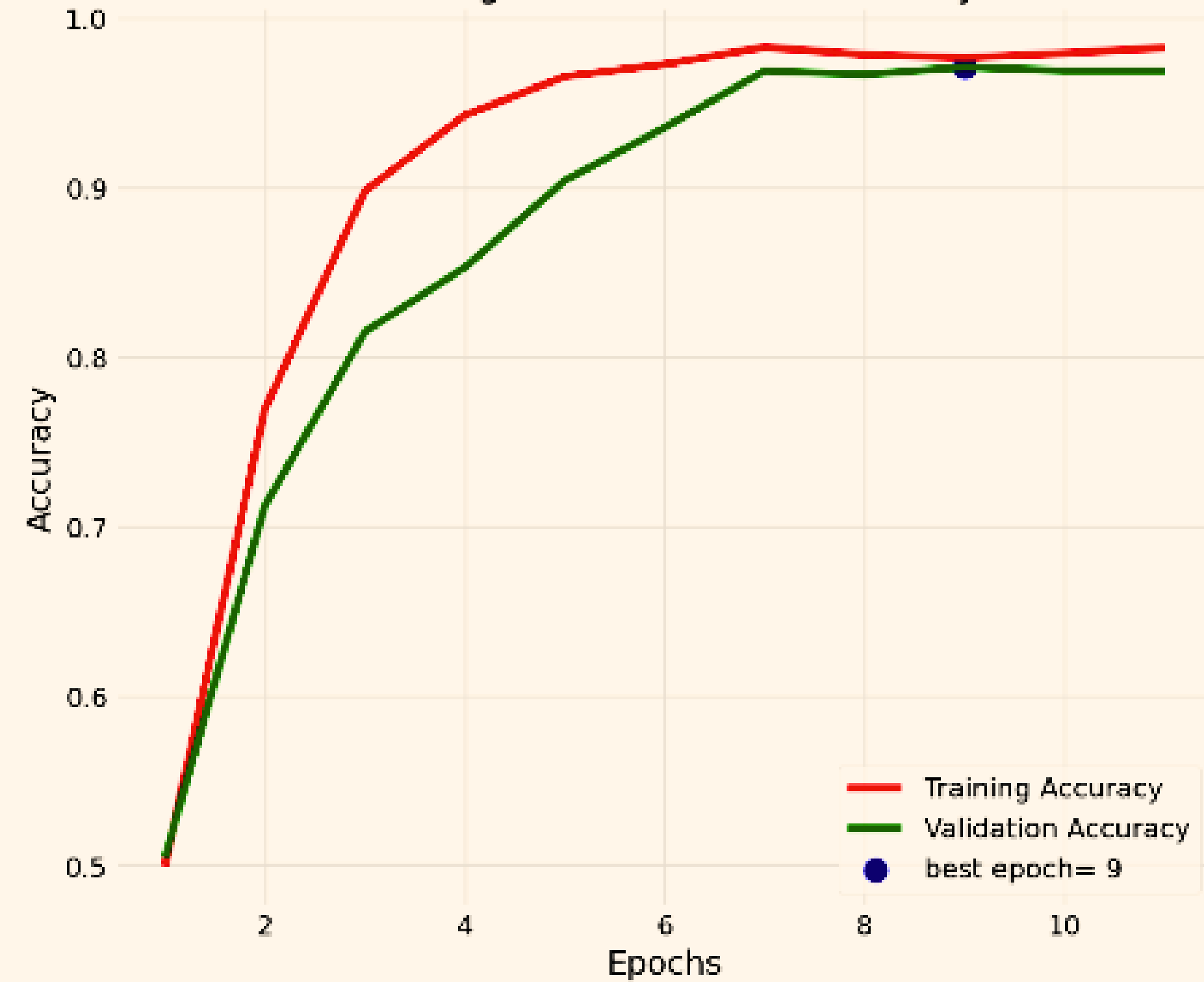
Best Performance Metrics
Approved

Evaluating model: EfficientNetB5				
Classification Report:				
	precision	recall	f1-score	support
Astrocitoma	0.93	0.97	0.95	58
Carcinoma	1.00	1.00	1.00	25
Ependimoma	1.00	0.93	0.97	15
Ganglioglioma	1.00	0.83	0.91	6
Germinoma	1.00	1.00	1.00	10
Glioblastoma	1.00	1.00	1.00	20
Granuloma	1.00	1.00	1.00	8
Meduloblastoma	1.00	1.00	1.00	13
Meningioma	0.98	0.99	0.98	87
Neurocitoma	1.00	0.96	0.98	46
Oligodendroglioma	0.96	0.96	0.96	23
Papiloma	1.00	0.96	0.98	24
Schwannoma	0.98	1.00	0.99	46
Tuberculoma	0.93	0.93	0.93	14
_NORMAL	0.98	1.00	0.99	53
accuracy			0.98	448
macro avg	0.98	0.97	0.98	448
weighted avg	0.98	0.98	0.98	448

Training and Validation Loss



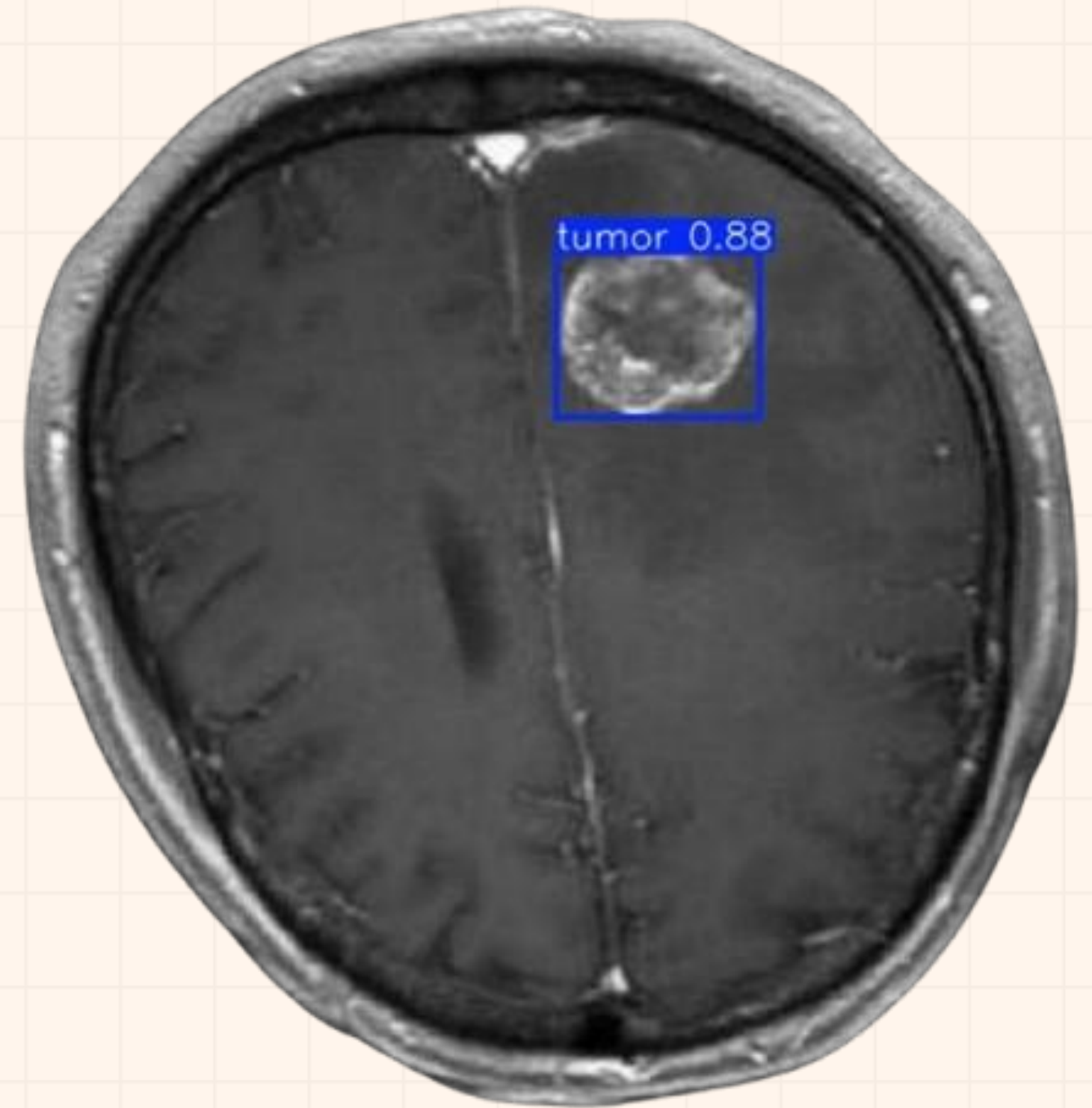
Training and Validation Accuracy



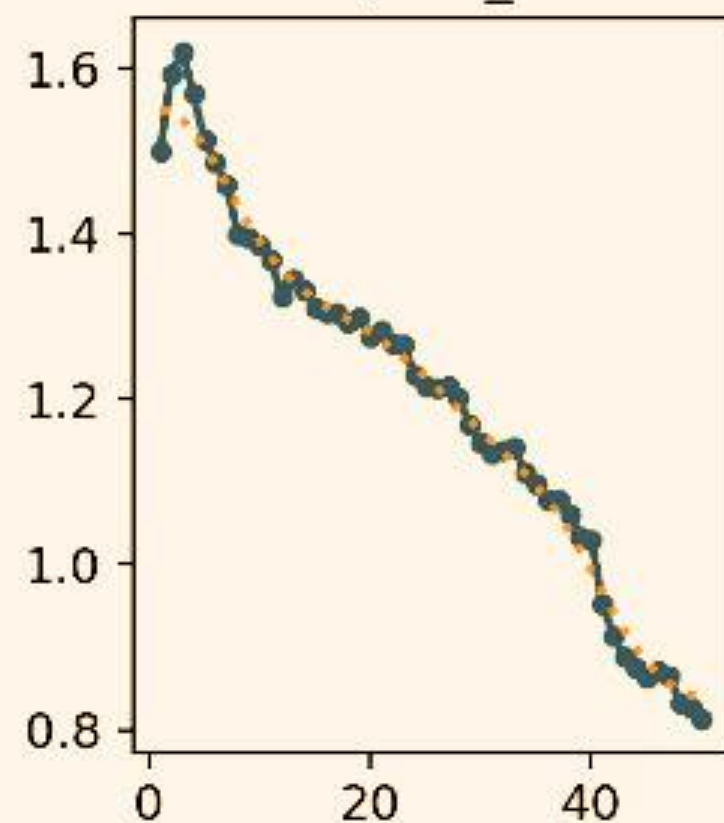
OBJECT DETECTION

Developed a deep learning-based object detection model to automatically detect and classify brain tumors from medical images using the YOLOv8 architecture.

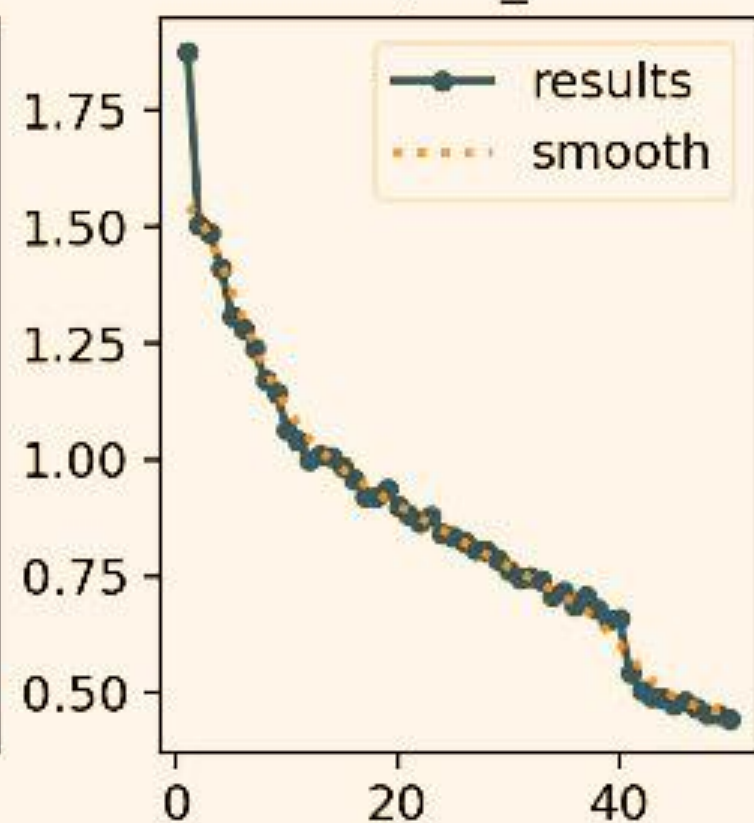
The goal is to assist radiologists and medical practitioners by providing fast and accurate detection to support early diagnosis and treatment planning



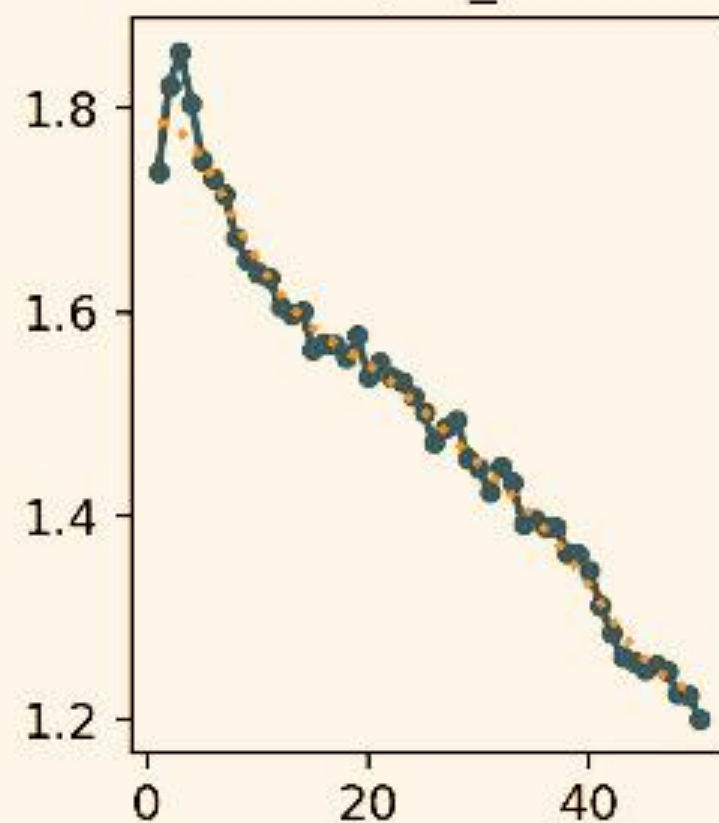
train/box_loss



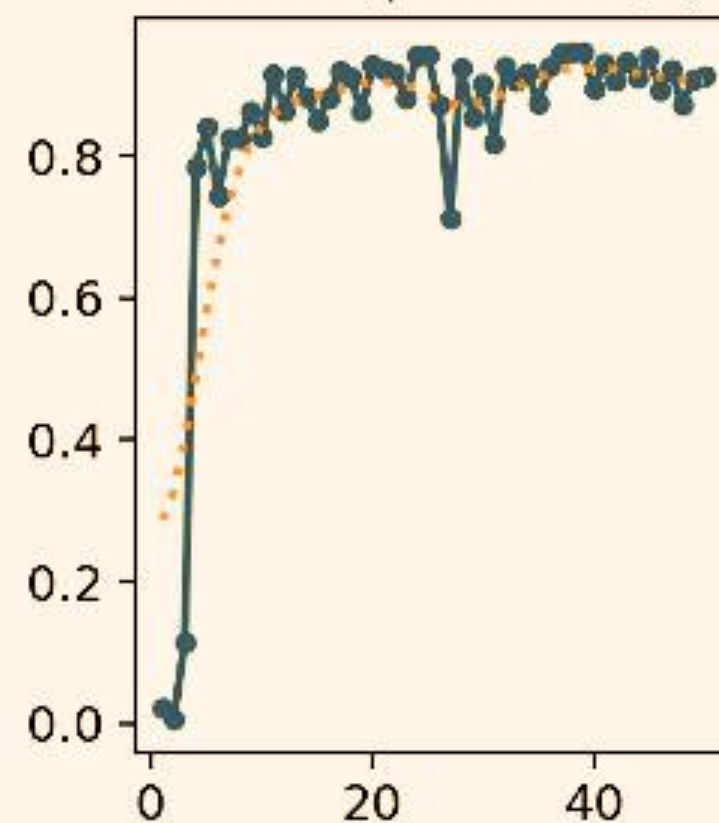
train/cls_loss



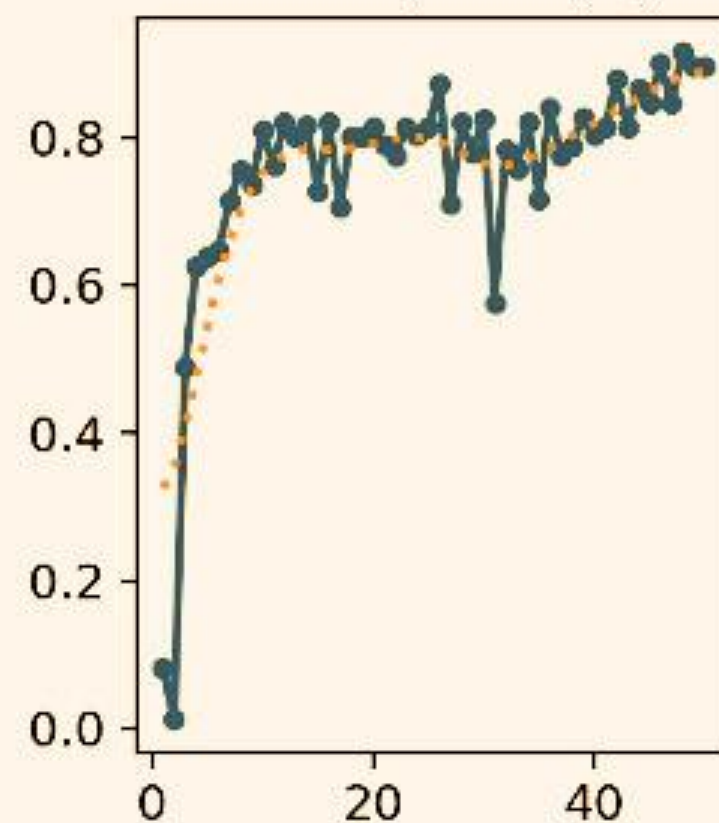
train/dfl_loss



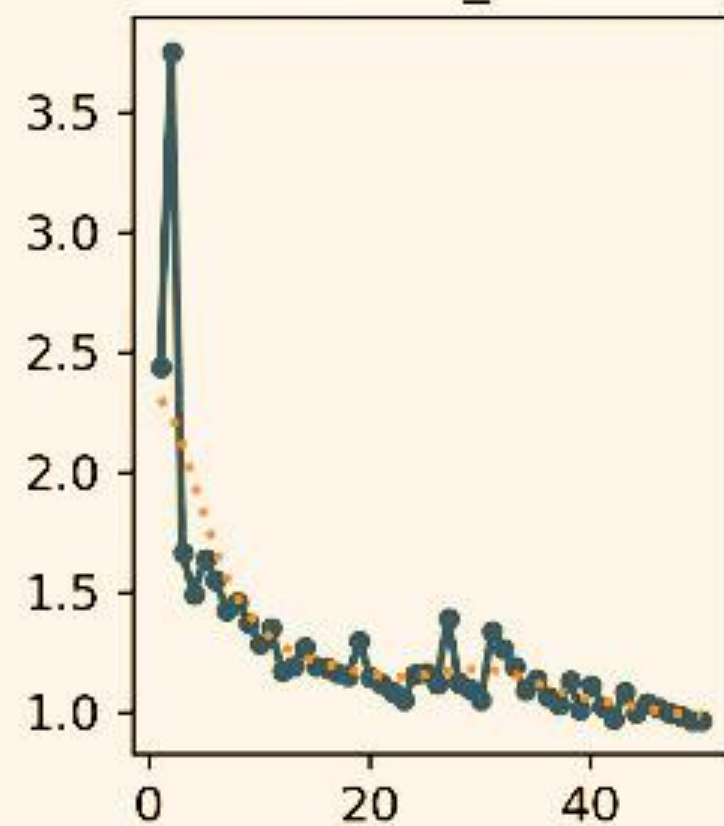
metrics/precision(B)



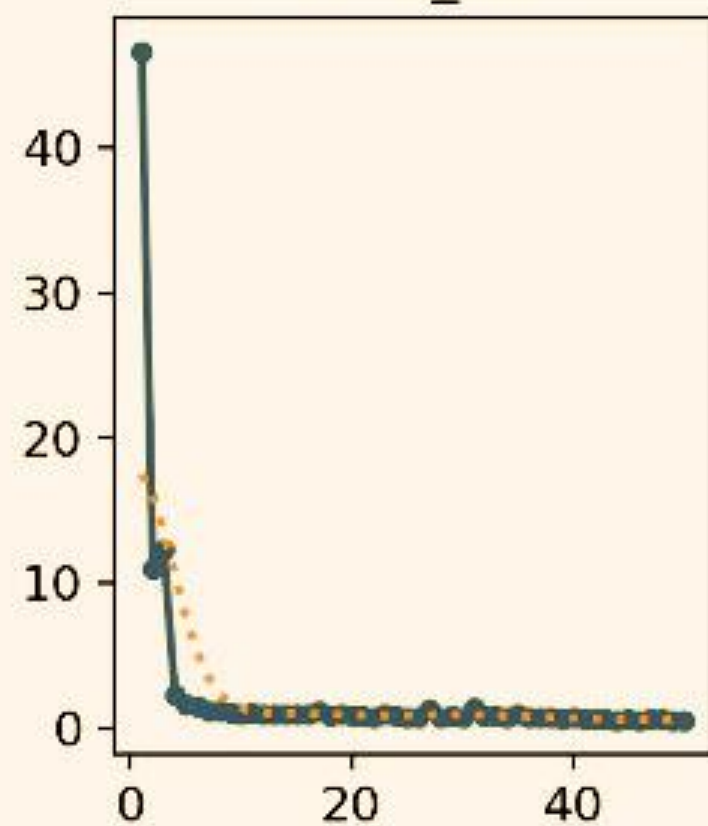
metrics/recall(B)



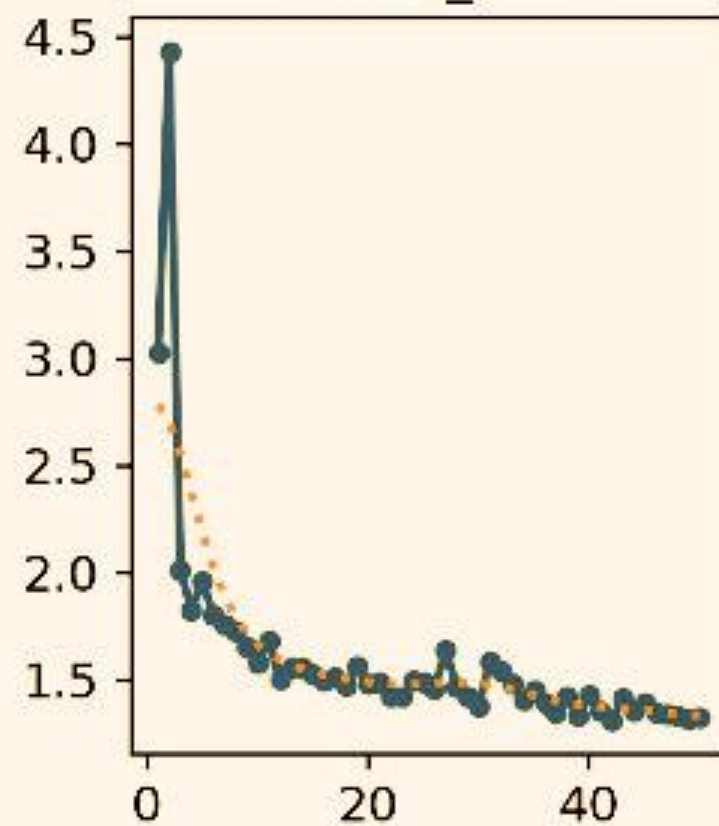
val/box_loss



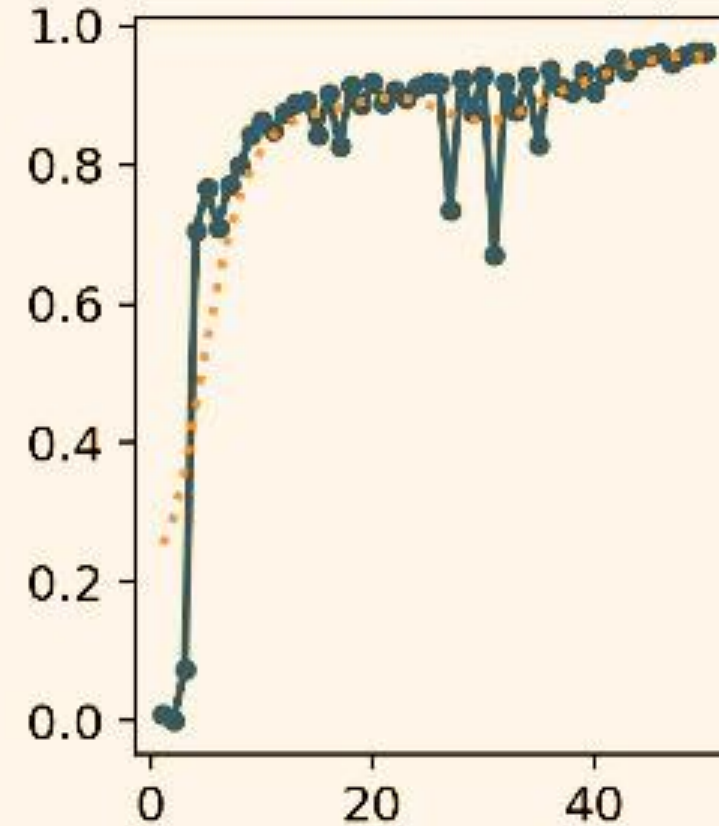
val/cls_loss



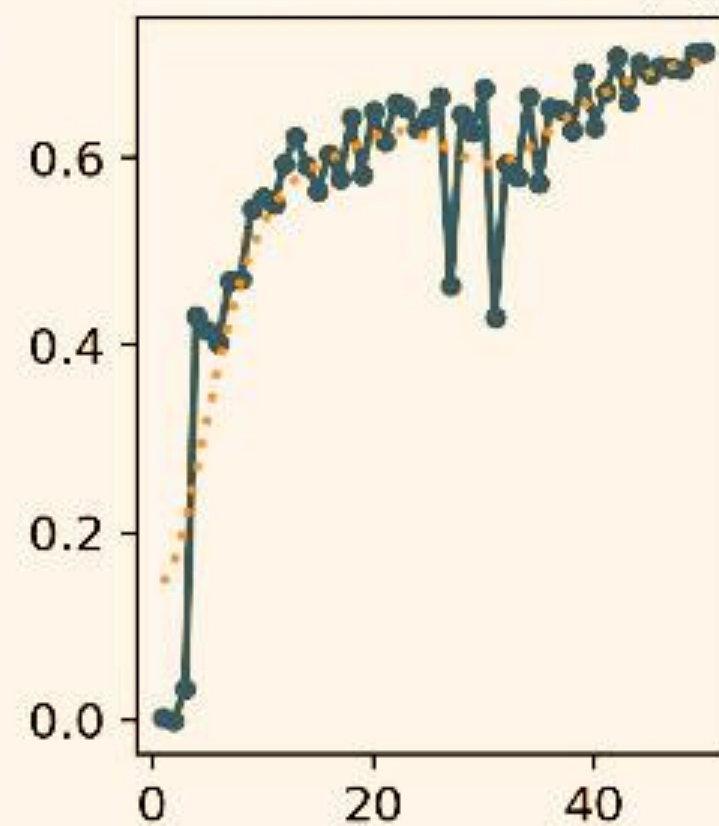
val/dfl_loss



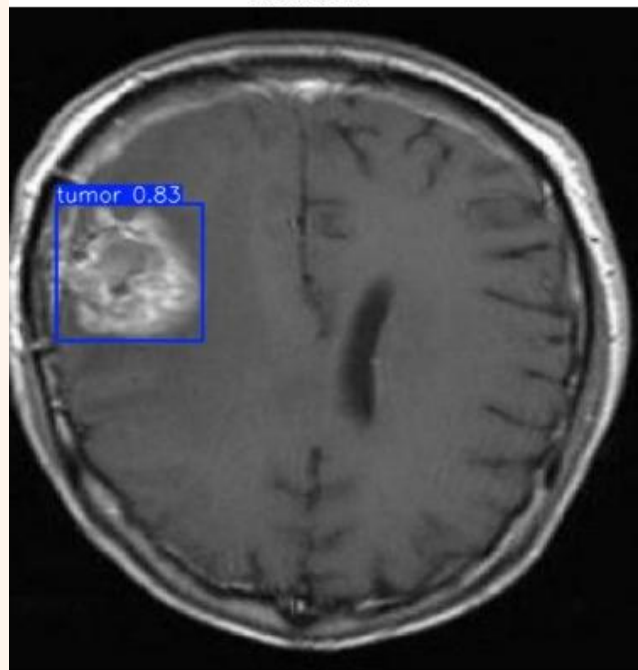
metrics/mAP50(B)



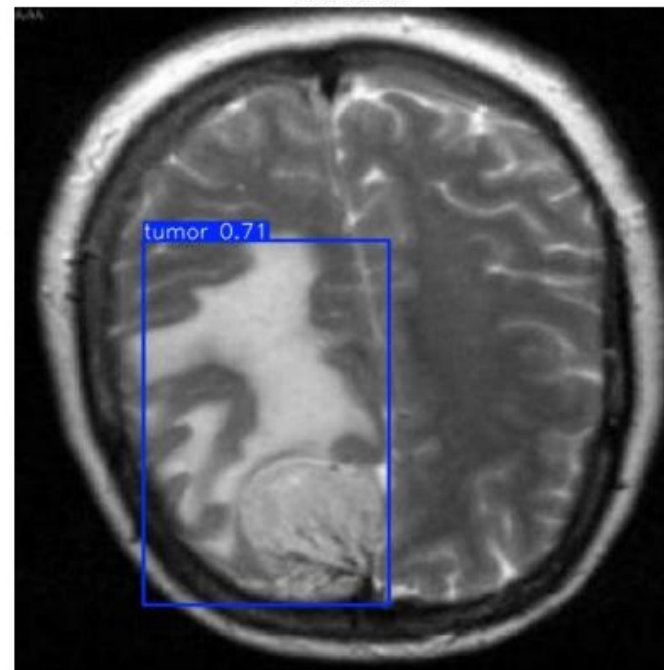
metrics/mAP50-95(B)



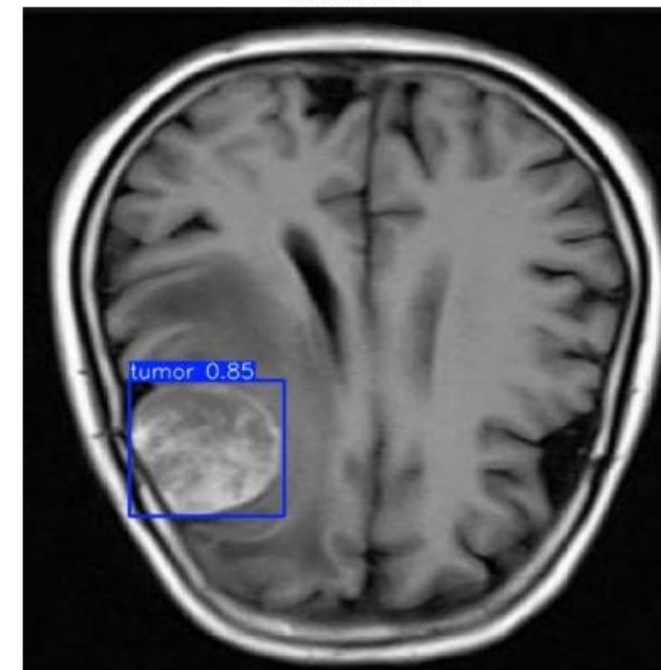
Prediction 1



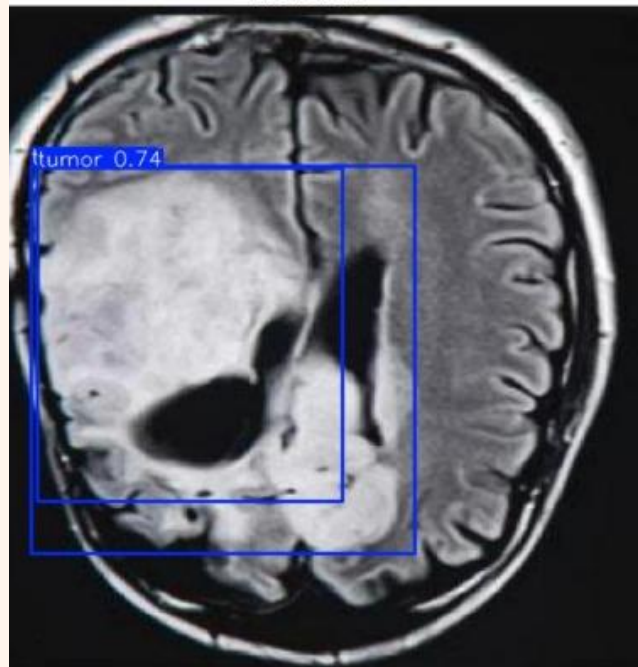
YoloV8 Predictions
Prediction 2



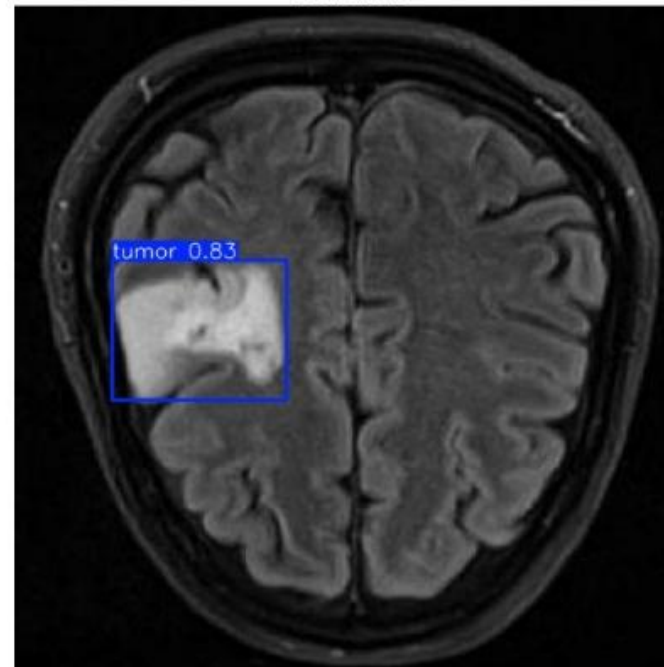
Prediction 3



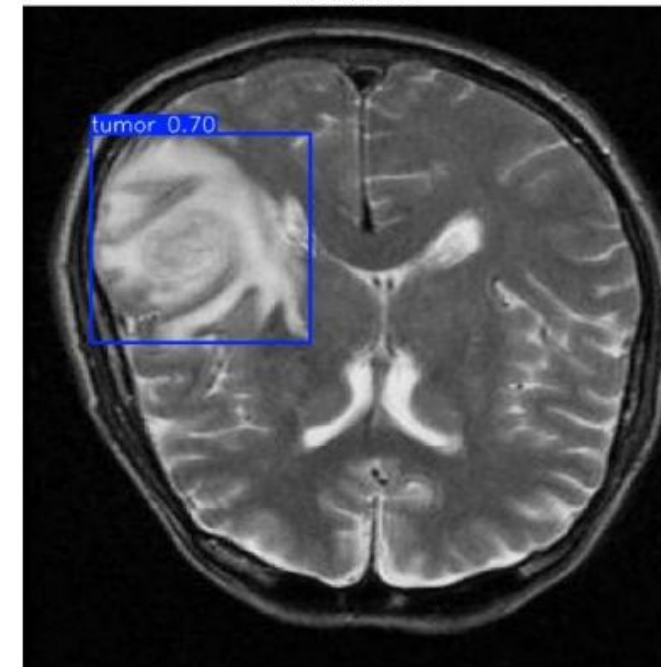
Prediction 4



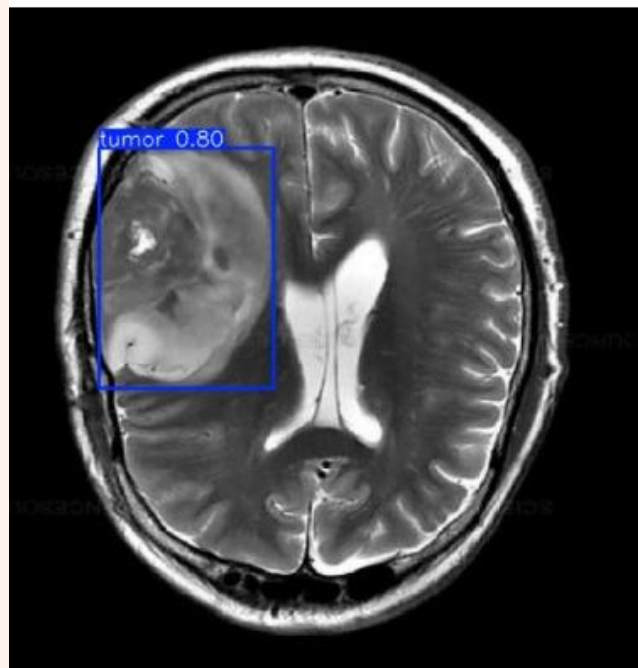
Prediction 5



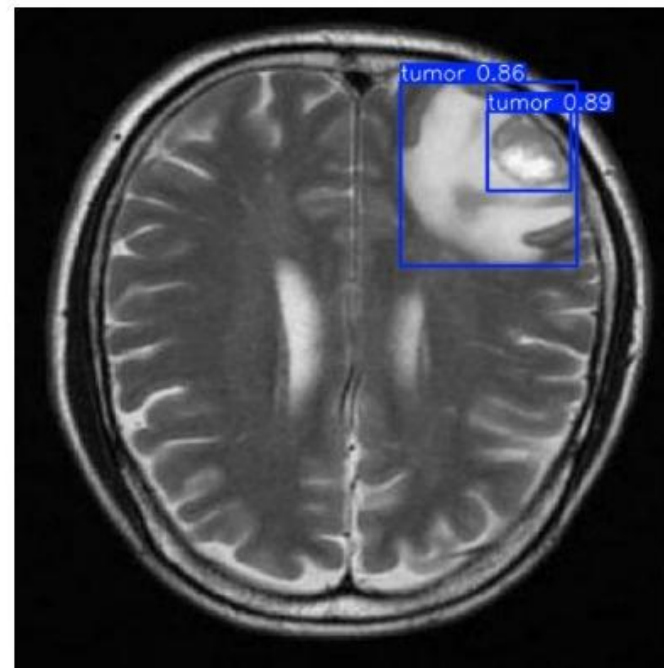
Prediction 6



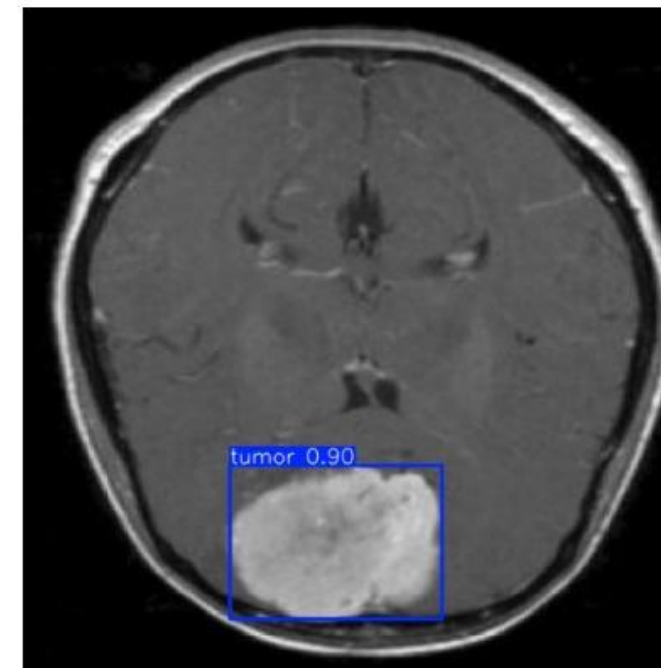
Prediction 7



Prediction 8



Prediction 9



DEPLOYMENT

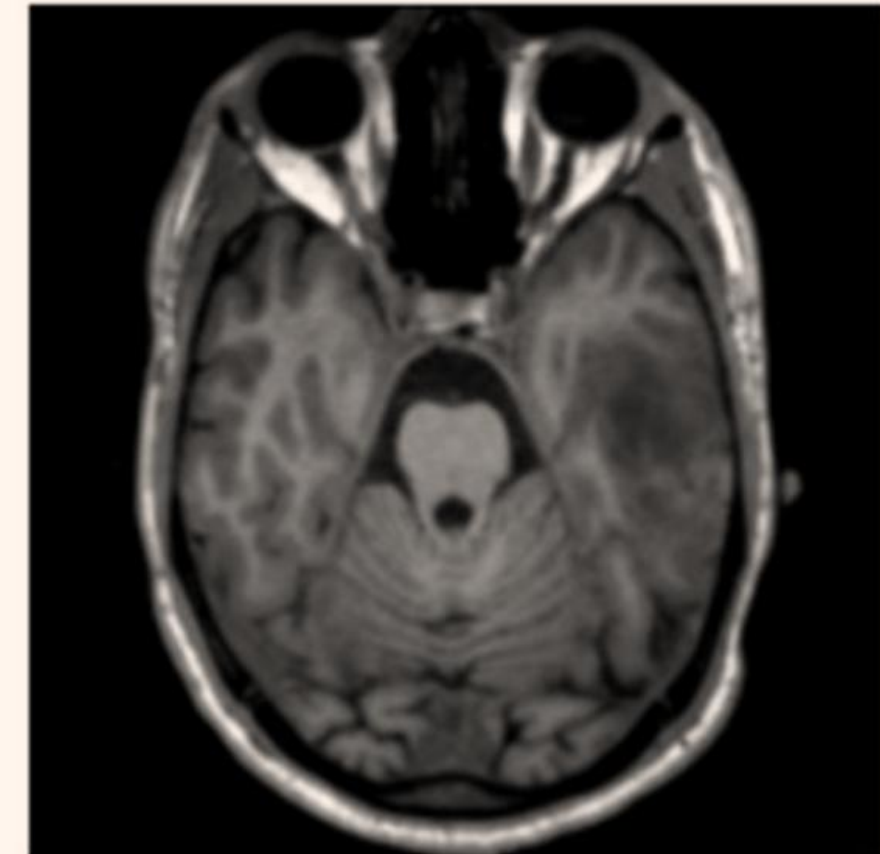
This Python script is a Flask web application that provides a user interface for classifying brain tumor images using a pre-trained deep learning model based on EfficientNetB5.

The app loads the model, processes uploaded images, performs predictions, and returns the tumor type along with the prediction probability.

Users can upload images via a web interface, where the backend resizes and preprocesses the images before running them through the model.

The prediction results are then displayed on a results page along with the uploaded image, which is encoded in base64 for embedding in the HTML.

Image Prediction Result



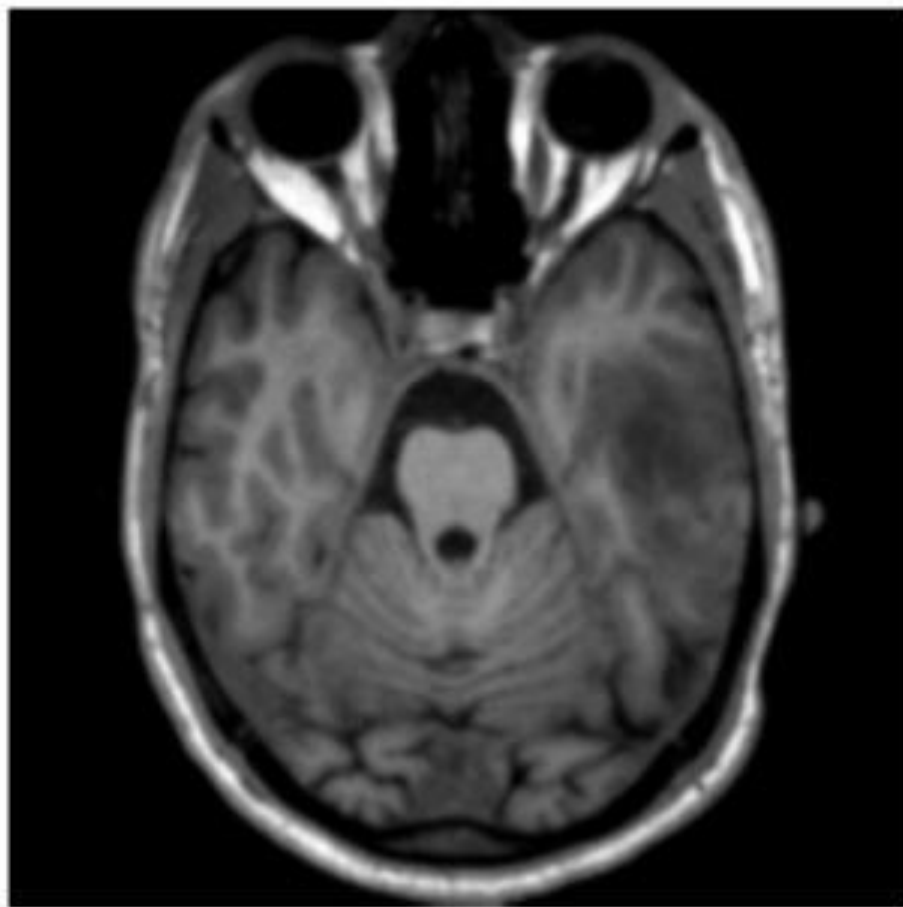
Prediction:

Astrocitoma

Probability: 0.99

Back

Image Prediction Result



Prediction:

Astrocitoma

Probability: 0.99

Back

THANK
YOU

