

```
import tensorflow as tf
from tensorflow import keras
from tensorflow.keras import layers
from tensorflow.keras.preprocessing.image import ImageDataGenerator
```

```
from google.colab import drive
drive.mount('/content/drive')
```

Drive already mounted at /content/drive; to attempt to forcibly remount, call drive.mount("/content/drive", force\_remount=True).

```
IMG_SIZE=224
BATCH_SIZE=32
```

```
train_datagen =
ImageDataGenerator(rescale=1./255,validation_split=0.2)
```

```
train_generator=train_datagen.flow_from_directory('/content/drive/
MyDrive/
brain_tumor_dataset',target_size=(IMG_SIZE,IMG_SIZE),batch_size=BATCH_
SIZE,class_mode='binary',subset='training')
```

Found 915 images belonging to 2 classes.

```
val_generator=train_datagen.flow_from_directory('/content/drive/
MyDrive/
brain_tumor_dataset',target_size=(IMG_SIZE,IMG_SIZE),batch_size=BATCH_
SIZE,class_mode='binary',subset='validation')
```

Found 227 images belonging to 2 classes.

```
model=keras.Sequential([layers.Conv2D(32,
(3,3),activation='relu',input_shape=(IMG_SIZE,IMG_SIZE,3)),
layers.MaxPooling2D((2,2)),layers.Conv2D(64,
(3,3),activation='relu'),layers.MaxPooling2D((2,2)),layers.Conv2D(128,
(3,3),activation='relu'),layers.MaxPooling2D((2,2)),
layers.Flatten(),layers.Dense(128,activation='relu'),layers.Dense(1,ac
tivation='sigmoid')
])
```

```
/usr/local/lib/python3.11/dist-packages/keras/src/layers/
convolutional/base_conv.py:107: UserWarning: Do not pass an
`input_shape`/`input_dim` argument to a layer. When using Sequential
models, prefer using an `Input(shape)` object as the first layer in
the model instead.
```

```
    super().__init__(activity_regularizer=activity_regularizer,
**kwargs)
```

```
model.summary()
```

```
Model: "sequential_2"
```

Layer (type) Param #	Output Shape
conv2d_8 (Conv2D) 896	(None, 222, 222, 32)
max_pooling2d_8 (MaxPooling2D) 0	(None, 111, 111, 32)
conv2d_9 (Conv2D) 18,496	(None, 109, 109, 64)
max_pooling2d_9 (MaxPooling2D) 0	(None, 54, 54, 64)
conv2d_10 (Conv2D) 73,856	(None, 52, 52, 128)
max_pooling2d_10 (MaxPooling2D) 0	(None, 26, 26, 128)
flatten_2 (Flatten) 0	(None, 86528)
dense_4 (Dense) 11,075,712	(None, 128)
dense_5 (Dense) 129	(None, 1)

Total params: 11,169,089 (42.61 MB)

Trainable params: 11,169,089 (42.61 MB)

Non-trainable params: 0 (0.00 B)

```
model.compile(optimizer='adam',loss='binary_crossentropy',metrics=['accuracy'])
```

```
model.fit(train_generator,epochs=5,validation_data=val_generator,batch_size=BATCH_SIZE)
```

```
/usr/local/lib/python3.11/dist-packages/keras/src/trainers/data_adapters/py_dataset_adapter.py:121: UserWarning: Your `PyDataset` class should call `super().__init__(**kwargs)` in its constructor. `**kwargs` can include `workers`, `use_multiprocessing`, `max_queue_size`. Do not pass these arguments to `fit()`, as they will be ignored.
```

```
self._warn_if_super_not_called()
```

Epoch 1/5

29/29 ————— 0s 5s/step - accuracy: 0.6000 - loss: 1.1251

```
/usr/local/lib/python3.11/dist-packages/keras/src/trainers/data_adapters/py_dataset_adapter.py:121: UserWarning: Your `PyDataset` class should call `super().__init__(**kwargs)` in its constructor. `**kwargs` can include `workers`, `use_multiprocessing`, `max_queue_size`. Do not pass these arguments to `fit()`, as they will be ignored.
```

```
self._warn_if_super_not_called()
```

29/29 ————— 206s 7s/step - accuracy: 0.6017 - loss: 1.1133 - val\_accuracy: 0.7004 - val\_loss: 0.5595

Epoch 2/5

29/29 ————— 125s 4s/step - accuracy: 0.7902 - loss: 0.4291 - val\_accuracy: 0.7489 - val\_loss: 0.4913

Epoch 3/5

29/29 ————— 134s 5s/step - accuracy: 0.8504 - loss: 0.3417 - val\_accuracy: 0.7709 - val\_loss: 0.4451

Epoch 4/5

29/29 ————— 126s 4s/step - accuracy: 0.9036 - loss: 0.2602 - val\_accuracy: 0.8106 - val\_loss: 0.3933

Epoch 5/5

29/29 ————— 128s 4s/step - accuracy: 0.9149 - loss: 0.2039 - val\_accuracy: 0.8634 - val\_loss: 0.2856

<keras.src.callbacks.history.History at 0x7f1d24503e90>

```
model.save('/content/drive/MyDrive/brain_tumor_model/brain_tumor_ml.h5')
```

WARNING:absl:You are saving your model as an HDF5 file via `model.save()` or `keras.saving.save\_model(model)`. This file format is considered legacy. We recommend using instead the native Keras format, e.g. `model.save('my\_model.keras')` or `keras.saving.save\_model(model, 'my\_model.keras')`.

```

from tensorflow.keras.models import load_model
from tensorflow.keras.preprocessing import image
import matplotlib.pyplot as plt
import numpy as np
model
=load_model('/content/drive/MyDrive/brain_tumor_model/brain_tumor_ml.h5')
print("Model Loaded")

```

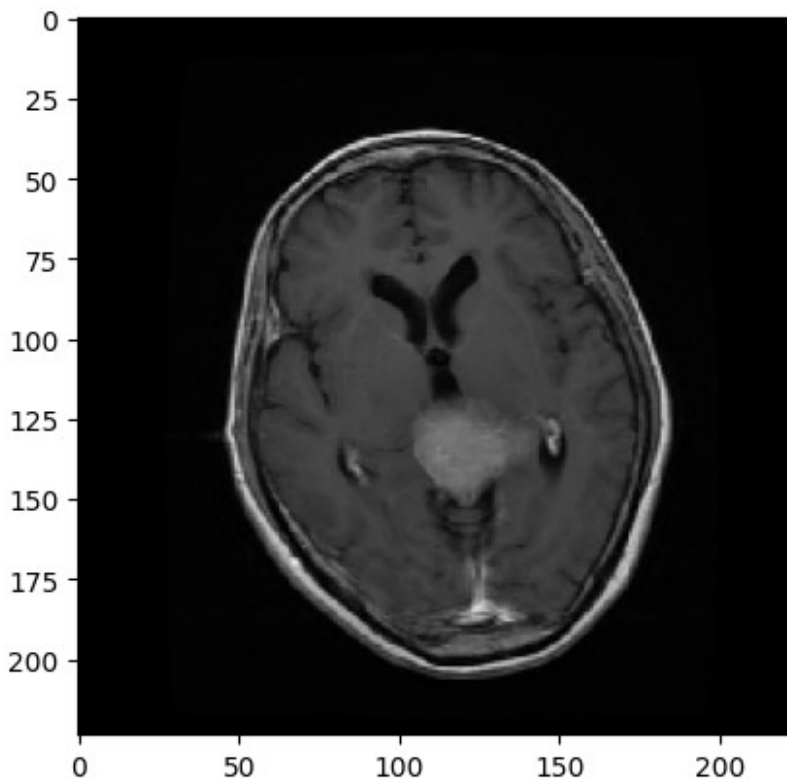
WARNING:absl:Compiled the loaded model, but the compiled metrics have yet to be built. `model.compile\_metrics` will be empty until you train or evaluate the model.

Model Loaded

```

test_image_path="/content/drive/MyDrive/brain_tumor_dataset/yes/Y1717.jpg"
img=image.load_img(test_image_path,target_size=(224,224))
plt.imshow(img)
plt.axis()
plt.show()

```



```

img_array=image.img_to_array(img)
img_array=np.expand_dims(img_array,axis=0)
img_array /= 255.

```

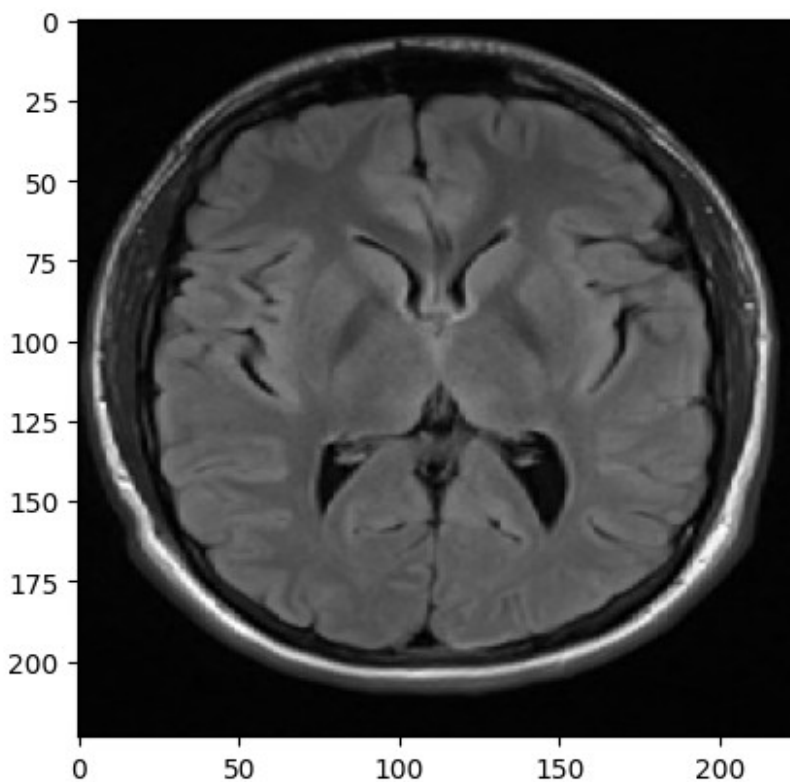
```

prediction = model.predict(img_array)
if prediction[0][0] > 0.5:
    print("Tumor Detected")
else:
    print("No Tumor Detected")

1/1 _____ 0s 189ms/step
Tumor Detected

test_image_path="/content/drive/MyDrive/brain_tumor_dataset/no/
N1435.jpg"
img=image.load_img(test_image_path,target_size=(224,224))
plt.imshow(img)
plt.axis()
plt.show()

```



```

img_array=image.img_to_array(img)
img_array=np.expand_dims(img_array,axis=0)
img_array /= 255.
prediction = model.predict(img_array)
if prediction[0][0] < 0.5:
    print("Tumor Detected")
else:
    print("No Tumor Detected")

```

1/1 ————— 0s 105ms/step  
No Tumor Detected