

```

import tensorflow as tf

import tensorflow as tf
from tensorflow import keras
from tensorflow.keras import layers
from tensorflow.keras.preprocessing.image import ImageDataGenerator
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/HI/Brain_Tumor_Dataset',
    target_size=(IMG_SIZE,IMG_SIZE),
    batch_size=BATCH_SIZE,
    class_mode='binary',
    subset='training'
)

```

Found 4221 images belonging to 2 classes.

```

val_generator=train_datagen.flow_from_directory(
    '/content/drive/MyDrive/HI/Brain_Tumor_Dataset',
    target_size=(IMG_SIZE,IMG_SIZE),
    batch_size=BATCH_SIZE,
    class_mode='binary',
    subset='validation'
)

```

Found 1055 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,activation='sigmoid')
])
model.summary()

```

/usr/local/lib/python3.10/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: "sequential"

Layer (type) Param #	Output Shape
conv2d (Conv2D) 896	(None, 222, 222, 32)
max_pooling2d (MaxPooling2D) 0	(None, 111, 111, 32)
conv2d_1 (Conv2D) 18,496	(None, 109, 109, 64)
max_pooling2d_1 (MaxPooling2D) 0	(None, 54, 54, 64)
conv2d_2 (Conv2D) 73,856	(None, 52, 52, 128)
max_pooling2d_2 (MaxPooling2D) 0	(None, 26, 26, 128)
flatten (Flatten) 0	(None, 86528)
dense (Dense) 11,075,712	(None, 128)
dense_1 (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=1,validation_data=val_generator,batch_size=BATCH_SIZE)
```

```
/usr/local/lib/python3.10/dist-packages/keras/src/trainers/data_adapters/py_dataset_adapter.py:122: 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()
```

```
132/132 _____ 654s 5s/step - accuracy: 0.8413 - loss: 0.3999 - val_accuracy: 0.9204 - val_loss: 0.1896
```

```
<keras.src.callbacks.history.History at 0x7c3940561690>
```

```
model.save('/content/drive/MyDrive/HI/Brain_Tumor_Dataset/braintumor.h5')
```

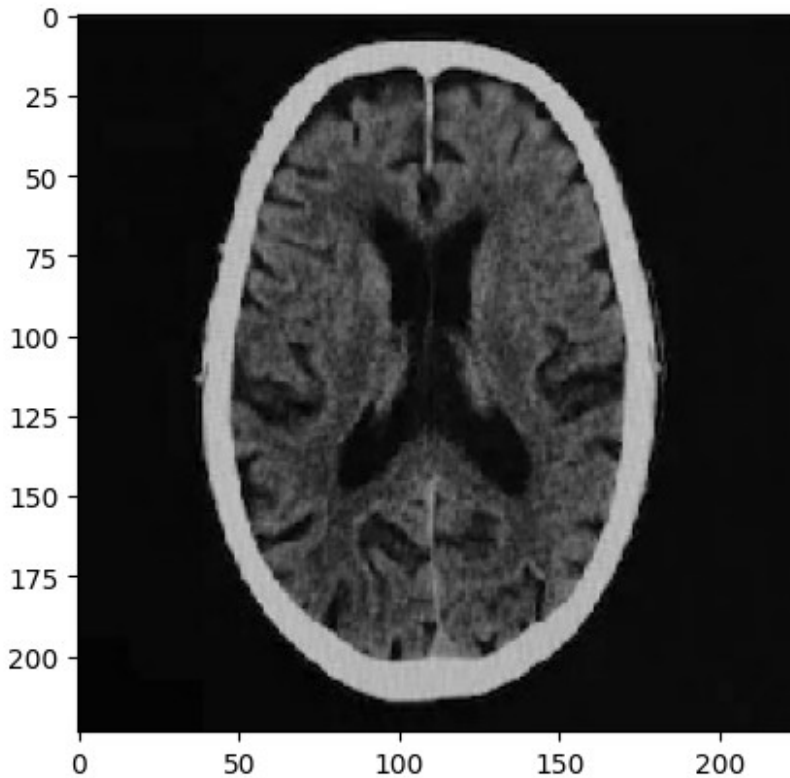
```
WARNING:absl:You are saving your model as an HDF5 file via `model.save()` or `keras.saving.save_model(model)`. This file format is considered legacy. We recommend using instead the native Keras format, e.g. `model.save('my_model.keras')` or `keras.saving.save_model(model, 'my_model.keras')`.
```

```
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/HI/Brain_Tumor_Dataset/braintumor.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/HI/Brain_Tumor_Dataset/Negative/Te-noTr_0005.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)
print(prediction)
```

```
1/1 _____ 0s 139ms/step
[[0.18178749]]
```

```
if(prediction>0.5):
    print("you have tumor")
else:
    print("you don't have tumor")
```

```
you don't have tumor
```

```
model.save('/content/drive/MyDrive/HI/Brain_Tumor_Dataset/
braintumor.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')`.
```

New section

