```
# Importing Libraries 💵 💻
import tensorflow as tf
from tensorflow.keras.preprocessing import image
from tensorflow.keras.preprocessing.image import ImageDataGenerator
from tensorflow.keras.layers import Flatten, Dense, Dropout, Activation, Input
from tensorflow.keras.layers import Conv2D, MaxPooling2D, BatchNormalization
from tensorflow.keras.models import Model, Sequential
from tensorflow.keras.optimizers import Adam
from tensorflow.keras.utils import plot model
from tensorflow.keras.applications import VGG16
import numpy as np
import os
import matplotlib.pyplot as plt
from PIL import Image
import zipfile
import warnings
warnings.filterwarnings("ignore")
%matplotlib inline
from google.colab import files
uploaded = files.upload() # Select the downloaded zip file
# Extract the dataset
import zipfile
with zipfile.ZipFile(list(uploaded.keys())[0], 'r') as zip_ref:
    zip ref.extractall('dataset')
    Choose Files archive (9).zip

    archive (9).zip(application/x-zip-compressed) - 155791278 bytes, last modified: 4/13/2025 - 100% done

     Saving archive (9).zip to archive (9).zip
from google.colab import drive
drive.mount('/content/drive')
# Copy dataset to Colab
!cp "/content/dataset.zip" ./
# Extract
!unzip -q dataset.zip -d dataset
→ Mounted at /content/drive
     cp: cannot stat '/content/dataset.zip': No such file or directory
     unzip: cannot find or open dataset.zip, dataset.zip.zip or dataset.zip.ZIP.
```

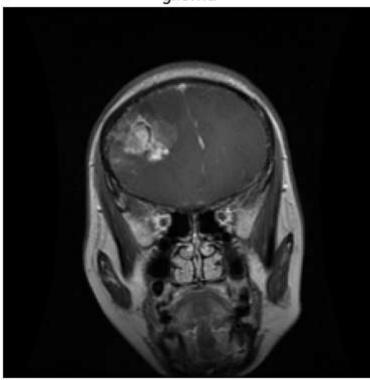
```
# Set dataset paths
train_datasets = "/content/dataset/Training"
validation_datasets = "/content/dataset/Testing"

# Display a sample image
img_path = '/content/dataset/Training/glioma/Tr-glTr_0000.jpg'
img = Image.open(img_path)
img = img.resize((224, 224)) # Resize to consistent size

# Display image
plt.imshow(img)
plt.title("glioma")
plt.axis('off') # Hide axis
plt.show()
```



glioma

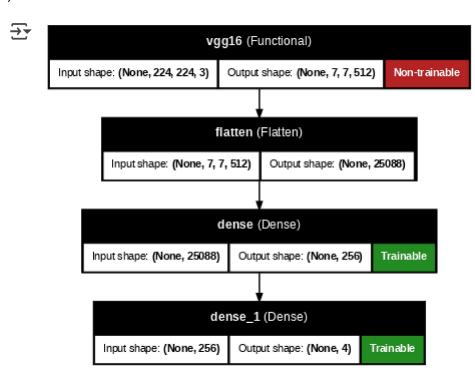


```
4/13/25, 4:00 PM
                                                  BrainTumor ipynb - Colab
        validation_datasets_generator_data = validation_datasets_generator.flow_from_directory(
            batch size=batch size,
            directory=validation_datasets,
            shuffle=True,
            target_size=(image_size, image_size),
            class_mode="categorical" # One-Hot-Encoded
        )
        return train datasets generator data, validation datasets generator data
    batch size = 64
    image size = 224
    train data , validation data = prepare the datasets(train datasets, validation datasets, bat
    \rightarrow Found 5712 images belonging to 4 classes.
         Found 1311 images belonging to 4 classes.
    # Load pre-trained VGG16 model without the top layers (include top=False) 🛂
    conv_base = VGG16(input_shape=(224, 224, 3), include_top=False, weights='imagenet')
    # Freeze the pre-trained layers to prevent training (transfer learning without fine-tuning)
    conv base.trainable = False
    # Build a new model on top of the pre-trained base 🧱
   model = Sequential()
    # Add the pre-trained VGG16 base model
   model.add(conv base)
    # Flatten the output for fully connected layers
   model.add(Flatten())
    # Add a dense layer with ReLU activation
   model.add(Dense(256, activation='relu'))
    # Add output layer with softmax activation for 4 classes
    model.add(Dense(4, activation='softmax'))
    # Compile the model with Adam optimizer and categorical crossentropy loss 🋠
    model.compile(optimizer='adam',
                  loss='categorical_crossentropy',
                  metrics=['accuracy'])
```

Downloading data from https://storage.googleapis.com/tensorflow/keras-applications/vgg16
58889256/58889256

Os Ous/step

```
plot_model(
    model,
    to_file='model.png',
    show_shapes=True,
    show_dtype=False,
    show_layer_names=True,
    dpi=55,
    show_trainable=True
)
```



```
\rightarrow \overline{\phantom{a}} Epoch 1/10
    89/89 -
                             74s 602ms/step - accuracy: 0.6126 - loss: 2.1328 - val_accura
    Epoch 2/10
                             - 7s 74ms/step - accuracy: 0.8750 - loss: 0.3358 - val_accuracy
    89/89 -
    Epoch 3/10
    89/89 -
                             - 36s 400ms/step - accuracy: 0.9162 - loss: 0.2359 - val_accura
    Epoch 4/10
                             7s 74ms/step - accuracy: 0.8594 - loss: 0.4297 - val_accuracy
    89/89 -
    Epoch 5/10
    89/89
                             • 34s 377ms/step - accuracy: 0.9395 - loss: 0.1696 - val_accura
    Epoch 6/10
                             7s 73ms/step - accuracy: 0.9688 - loss: 0.1105 - val_accuracy
    89/89 -
    Epoch 7/10
    89/89 —
                             Epoch 8/10
```

```
      89/89
      7s 74ms/step - accuracy: 0.9844 - loss: 0.0891 - val_accuracy

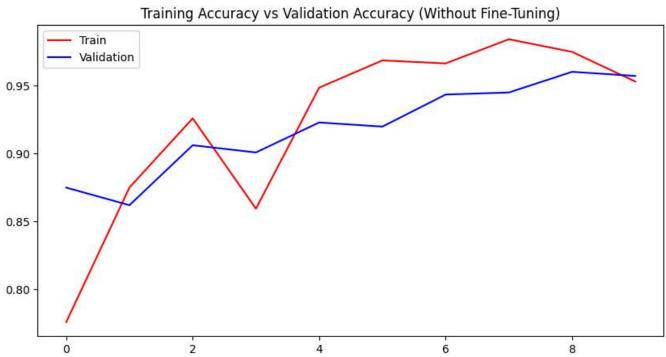
      Epoch 9/10
      76s 394ms/step - accuracy: 0.9723 - loss: 0.0867 - val_accuracy

      Epoch 10/10
      7s 78ms/step - accuracy: 0.9531 - loss: 0.0922 - val_accuracy
```

```
# Train the model
print("\nTraining model without fine-tuning...")
# Plotting the graph of Accuracy and Validation Accuracy
plt.figure(figsize=(10, 5))
plt.title('Training Accuracy vs Validation Accuracy (Without Fine-Tuning)')
plt.plot(history.history['accuracy'], color='red', label='Train')
plt.plot(history.history['val_accuracy'], color='blue', label='Validation')
plt.legend()
plt.show()
```

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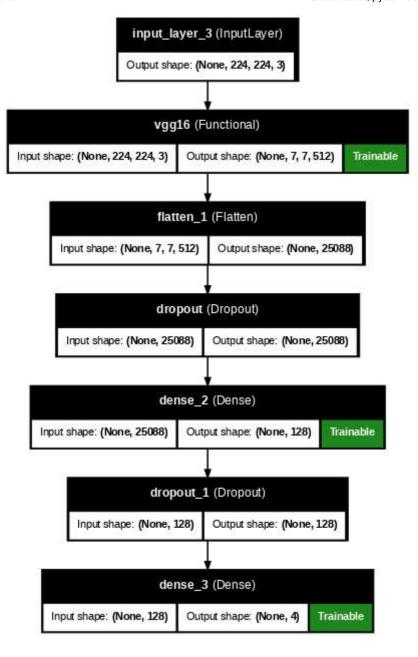
raining model without fine-tuning...



```
print("\nTransfer Learning With Fine-Tuning ( ) ( ) ( )
# Load VGG16 model
base_model = VGG16(input_shape=(224, 224, 3), include_top=False, weights='imagenet')
# Freeze layers of base model
for layer in base_model.layers:
```

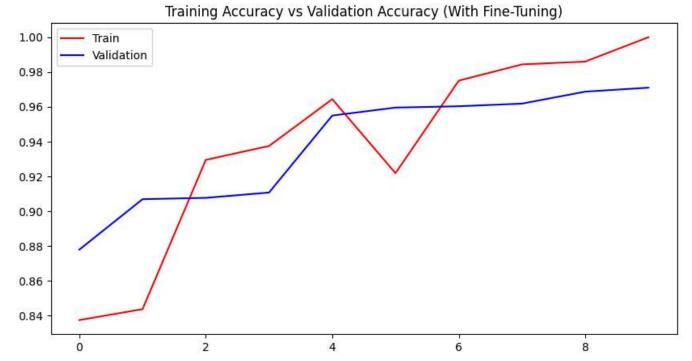
```
layer.trainable = False
# Optionally unfreeze layers of a specific block (fine-tuning)
for layer in base model.layers:
    if layer.name.startswith('block5'):
        layer.trainable = True
# Build the model
inputs = Input(shape=(224, 224, 3))
x = base model(inputs, training=False) # Base model in inference mode
x = Flatten()(x)
x = Dropout(0.3)(x)
x = Dense(128, activation='relu')(x)
x = Dropout(0.2)(x)
outputs = Dense(4, activation='softmax')(x)
model1 = Model(inputs, outputs)
# Compile model
model1.compile(optimizer=Adam(learning_rate=0.0001), # While Finetuning value of learning r
              loss='categorical crossentropy',
              metrics=['accuracy'])
\rightarrow
     Transfer Learning With Fine-Tuning 🖸 🧠
# Plot model architecture
plot_model(
    model1,
    to file='model1.png',
    show shapes=True,
    show_dtype=False,
    show layer names=True,
    dpi=55,
    show_layer_activations=False,
    show_trainable=True
)
```







Training model with fine-tuning... Epoch 1/10 - 58s 544ms/step - accuracy: 0.7400 - loss: 0.6818 - val_accura 89/89 -Epoch 2/10 **7s** 73ms/step - accuracy: 0.8438 - loss: 0.4931 - val_accuracy 89/89 -Epoch 3/10 89/89 -- **59s** 438ms/step - accuracy: 0.9264 - loss: 0.2038 - val_accura Epoch 4/10 89/89 -**7s** 78ms/step - accuracy: 0.9375 - loss: 0.1942 - val accuracy Epoch 5/10 89/89 -- **75s** 447ms/step - accuracy: 0.9557 - loss: 0.1253 - val accura Epoch 6/10 89/89 -• **7s** 76ms/step - accuracy: 0.9219 - loss: 0.1253 - val accuracy Epoch 7/10 89/89 -**- 38s** 427ms/step - accuracy: 0.9728 - loss: 0.0710 - val accura Epoch 8/10 89/89 -**- 7s** 73ms/step - accuracy: 0.9844 - loss: 0.0836 - val accuracy Epoch 9/10 89/89 -77s 443ms/step - accuracy: 0.9884 - loss: 0.0355 - val accura Epoch 10/10 - **7s** 76ms/step - accuracy: **1.**0000 - loss: **0.**0435 - val accuracy 89/89 -



Print final accuracies

```
BrainTumor ipynb - Colab
final_acc_no_finetune = history.history['val_accuracy'][-1] * 100
final acc with finetune = history1.history['val accuracy'][-1] * 100
print("\nFinal Results:")
print(f"VGG16 Without Fine-Tuning = Validation Accuracy: {final_acc_no_finetune:.2f}%")
print(f"VGG16 With Fine-Tuning = Validation Accuracy: {final acc with finetune:.2f}%")
→ ve Keras format, e.g. `model.save('my model.keras')` or `keras.saving.save model(model,
    ve Keras format, e.g. `model.save('my_model.keras')` or `keras.saving.save_model(model,
# First, let's check the actual class indices from our training data generator
train data.class indices
→ {'glioma': 0, 'meningioma': 1, 'notumor': 2, 'pituitary': 3}
```

```
# Load the saved models
from tensorflow.keras.models import load model
import numpy as np
from tensorflow.keras.preprocessing import image
import matplotlib.pyplot as plt
from google.colab import files
from PIL import Image
# Load both models (you can choose which one to use)
model no ft = load model('model.h5')
model_with_ft = load_model('model1.h5')
# CORRECTED CLASS NAMES ORDER
class_names = ['glioma', 'meningioma', 'no tumor', 'pituitary'] # Changed order
def predict_image(img_path, model=model_with_ft):
    # Load and preprocess image
    img = image.load_img(img_path, target_size=(224, 224))
    img_array = image.img_to_array(img)
    img_array = np.expand_dims(img_array, axis=0)
    img_array /= 255.0 # Normalize
    # Make prediction
    prediction = model.predict(img_array)
    predicted_class = class_names[np.argmax(prediction)] # Now uses correct order
    confidence = np.max(prediction) * 100
    # Display results
    plt.imshow(img)
    plt.title(f"Predicted: {predicted_class}\nConfidence: {confidence:.2f}%")
```

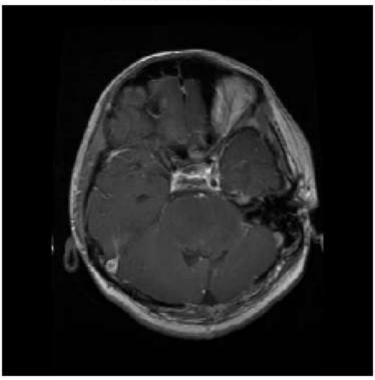
```
plt.axis('off')
    plt.show()
    return predicted_class, confidence
→ WARNING:absl:Compiled the loaded model, but the compiled metrics have yet to be built. `
     WARNING:absl:Compiled the loaded model, but the compiled metrics have yet to be built. `
# Test with sample images from the dataset
test images = [
    '/content/dataset/Testing/glioma/Te-gl_0010.jpg',
    '/content/dataset/Testing/meningioma/Te-me_0010.jpg',
    '/content/dataset/Testing/pituitary/Te-pi_0010.jpg',
    '/content/dataset/Testing/notumor/Te-no_0010.jpg'
]
print("\nTesting with sample images from dataset:")
for img_path in test_images:
    print(f"\nAnalyzing: {img_path}")
    predict_image(img_path)
```



Testing with sample images from dataset:

Analyzing: /content/dataset/Testing/glioma/Te-gl_0010.jpg
1/1 ______ 1s 581ms/step

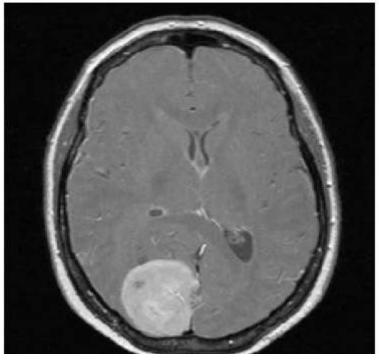
Predicted: glioma Confidence: 99.88%



Analyzing: /content/dataset/Testing/meningioma/Te-me_0010.jpg

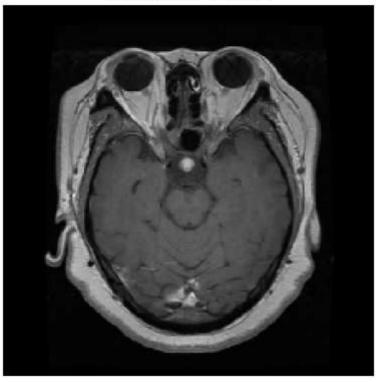
1/1 ______ 0s 44ms/step

Predicted: meningioma Confidence: 99.99%



Analyzing: /content/dataset/Testing/pituitary/Te-pi_0010.jpg
1/1 ______ 0s 43ms/step

Predicted: pituitary Confidence: 99.84%



Analyzing: /content/dataset/Testing/notumor/Te-no_0010.jpg

1/1 ______ 0s 48ms/step

Predicted: no tumor Confidence: 99.91%



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