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Project: Brain Tumor Prediction

Importing Required Library

```
In [1]:
         import tensorflow as tf
         import numpy as np
         from tensorflow.keras import models, Sequential, layers
         import matplotlib.pyplot as plt
         from tensorflow.keras.preprocessing.image import ImageDataGenerator
```

Read Train dataset

```
In [4]:
         train_ds = tf.keras.preprocessing.image_dataset_from_directory(
             directory = "Training",
             shuffle = True,
             label_mode = 'int',
             batch_size = 32,
             image_size = (256, 256)
         )
```

Found 5712 files belonging to 4 classes.

Read Test dataset

```
In [11]:
          test_data = tf.keras.preprocessing.image_dataset_from_directory(
              directory = "Testing",
              shuffle = True,
              label_mode = 'int',
              batch_size = 32,
              image_size = (256, 256)
          )
```

Found 1311 files belonging to 4 classes.

Out[16]: ['glioma', 'meningioma', 'notumor', 'pituitary']

```
In [12]:
          train_size = int(0.9*len(train_ds))
          valid_size = int(0.1*len(train_ds))
          train_size, valid_size
Out[12]: (161, 17)
In [14]:
          train_data = train_ds.take(train_size)
          valid_data = train_ds.skip(train_size)
          len(train_data) , len(valid_data), len(test_data)
Out[14]: (161, 18, 41)
In [16]:
          class_name = train_ds.class_names
          class_name
```

```
In [21]:
    train_datagen = ImageDataGenerator(
        rescale = 1./255,
        horizontal_flip = True,
        vertical_flip = True,
        rotation_range = 10
)

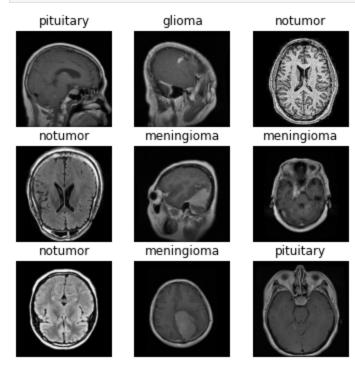
train_generator = train_datagen.flow_from_directory(
        directory="Training",
        target_size=(224, 224),
        batch_size=32,
        class_mode="sparse",
        shuffle=True
)
```

Found 5712 images belonging to 4 classes.

Visualize the Data

```
In [17]:
    plt.figure(figsize = (6,6))

    for image_batch, label_batch in train_ds.take(1):
        for i in range(9):
            ax = plt.subplot(3,3,i+1)
            plt.imshow(image_batch[i].numpy().astype('uint8'))
            plt.title(class_name[label_batch[i]])
            plt.axis('off')
```



Preprocessing

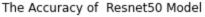
```
train_data = train_data.cache().shuffle(1000).prefetch(buffer_size = tf.data.AUTOTUNE)
test_data = test_data.cache().shuffle(1000).prefetch(buffer_size = tf.data.AUTOTUNE)
valid_data = valid_data.cache().shuffle(1000).prefetch(buffer_size = tf.data.AUTOTUNE)
```

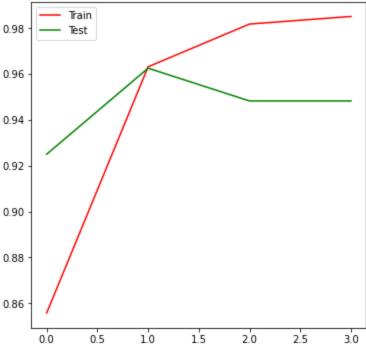
Convalution Neural Network Model

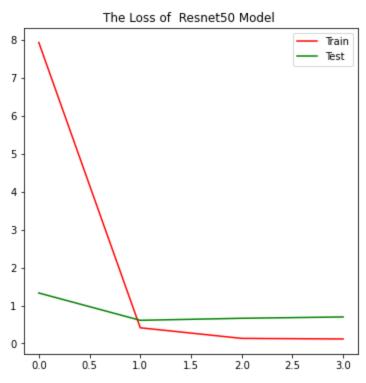
```
In [20]:
        Resnet50 = tf.keras.applications.ResNet50(
           include_top = False,
           weights="imagenet",
           input_shape = (256, 256, 3),
           classes = 4
         )
 In [21]:
        Resnet_Model = Sequential()
        Resnet50.trainable = False
        Resnet_Model.add(Resnet50)
        Resnet_Model.add(layers.Flatten())
         Resnet_Model.add(layers.Dense(512, activation = 'relu'))
        Resnet_Model.add(layers.Dense(4, activation = 'softmax'))
 In [22]:
        Resnet_Model.summary()
        Model: "sequential"
        Layer (type)
                             Output Shape
                                                 Param #
        ______
                             (None, 8, 8, 2048)
        resnet50 (Functional)
                                                 23587712
        flatten (Flatten)
                              (None, 131072)
        dense (Dense)
                              (None, 512)
                                                 67109376
        dense_1 (Dense)
                              (None, 4)
                                                 2052
        ______
        Total params: 90,699,140
        Trainable params: 67,111,428
        Non-trainable params: 23,587,712
 In [23]:
        Resnet_Model.compile(
           optimizer = 'adam',
           loss = tf.keras.losses.SparseCategoricalCrossentropy(from_logits = False),
           metrics = ['accuracy']
         )
 In [25]:
        Resnet50_history = Resnet_Model.fit(
           train_data,
           epochs = 4,
           validation_data = valid_data
        )
        Epoch 1/4
        - val_loss: 1.3296 - val_accuracy: 0.9250
        Epoch 2/4
        - val_loss: 0.6099 - val_accuracy: 0.9625
        Epoch 3/4
        - val_loss: 0.6636 - val_accuracy: 0.9482
        Epoch 4/4
        1 - val_loss: 0.6983 - val_accuracy: 0.9482
        Vertensions/Sefs is ory history keys()
Loading [MathJax]/extensions/Safe.js
```

```
Out[26]: dict_keys(['loss', 'accuracy', 'val_loss', 'val_accuracy'])

In [29]:
    plt.figure(figsize = (6,6))
    plt.plot(range(4), Resnet50_history.history['accuracy'], color = 'red', label = 'Train')
    plt.plot(range(4), Resnet50_history.history['val_accuracy'], color = 'green', label = 'Test
    plt.title('The Accuracy of Resnet50 Model ')
    plt.legend()
    plt.show()
    plt.figure(figsize = (6,6))
    plt.plot(range(4), Resnet50_history.history['loss'], color = 'red', label = 'Train')
    plt.plot(range(4), Resnet50_history.history['val_loss'], color = 'green', label = 'Test')
    plt.title('The Loss of Resnet50 Model ')
    plt.legend()
    plt.show()
```



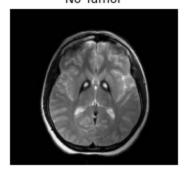




NoTumor Prediction

```
In [46]:
          import cv2
          import matplotlib.pyplot as plt
          import numpy as np
In [43]:
          # Read the image using OpenCV
          notumor = cv2.imread("no_tumor.jpg")
In [44]:
          # Convert the image from BGR (OpenCV format) to RGB (Matplotlib format)
          notumor_rgb = cv2.cvtColor(notumor, cv2.COLOR_BGR2RGB)
In [45]:
          plt.figure(figsize=(3, 3))
          plt.imshow(notumor_rgb)
          plt.title("No Tumor")
          plt.axis("off")
          plt.show()
```

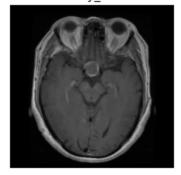
No Tumor



Pituitary Tumor Prediction

```
In [53]: pituitary_tumor = cv2.imread("pituitary.jpg")
# convert it RGB format
pituitary_tumor_rgb = cv2.cvtColor(pituitary_tumor, cv2.COLOR_BGR2RGB)
In [54]: plt.figure(figsize=(3, 3))
plt.imshow(pituitary_tumor_rgb)
plt.title("Pituitary_Tumor")
plt.axis("off")
plt.show()
```

Pituitary_Tumor



```
In [55]: # Resizze and reshape the image
   pituitary_tumor= cv2.resize(pituitary_tumor, (256, 256))
   pituitary_tumor = pituitary_tumor.reshape(1,256, 256, 3)
In [59]: Pred_tumor = Resnet_Model.predict(pituitary_tumor)
```

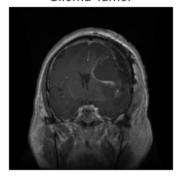
print("Predicted Tumor is : ", class_name[np.argmax(Pred_tumor[0])])

Predicted Tumor is : pituitary

Glioma Tumor Prediction

```
glioma_tumor = cv2.imread("glioma.jpg")
# convert it RGB format
glioma_tumor_rgb = cv2.cvtColor(pituitary_tumor, cv2.COLOR_BGR2RGB)
plt.figure(figsize=(3, 3))
plt.imshow(glioma_tumor_rgb)
plt.title("Glioma Tumor")
plt.axis("off")
plt.show()
```

Glioma Tumor



```
In [62]: # Resizze and reshape the image

Loading [MathJax]/extensions/Safe.js = cv2.resize(glioma_tumor, (256, 256))
```

```
glioma_tumor = glioma_tumor.reshape(1,256, 256, 3)
glioma_tumor = Resnet_Model.predict(glioma_tumor)
print("Predicted Tumor is : ", class_name[np.argmax(glioma_tumor[0])])
Predicted Tumor is : glioma
```

```
In [ ]:
```