



Brain Tumor Detection

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Road Map

1. Data Discovery & Formulation
2. Data Preparation & Processing
3. Design a Model
4. Model Building
5. Results
6. Measuring of Effeteness



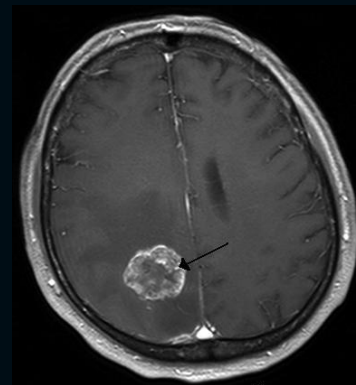
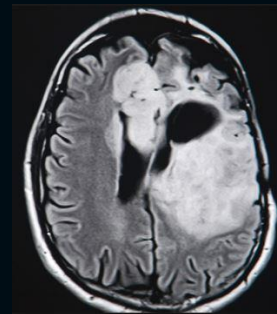
1. Data Discovery & Formulation

- **Business Problem-** Identifying the Brain Tumor by predict whether the patient has a brain tumor or not from automate the process of brain cropping from MRI scans without human intervention.
- **Solution (Critical Objective)-** This project is a combination of CNN model classification problem & Computer Vision problem that would classify and predict if subject has a tumor or not base on MRI scan's existing and new brain images.

1. Data Discovery & Formulation- cont.

- **About Brain Tumor (Background)-**

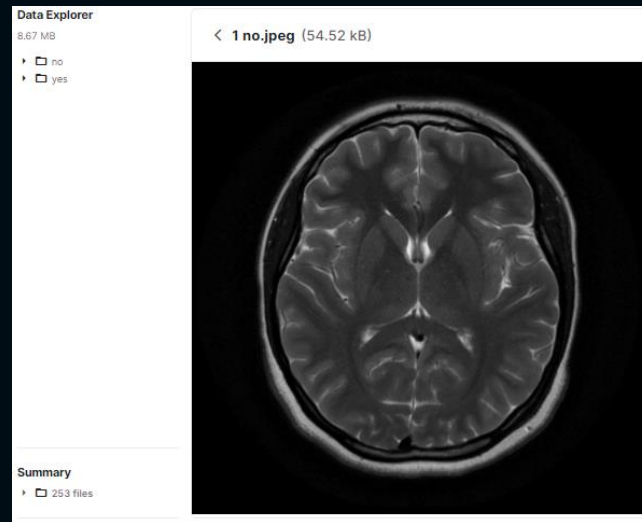
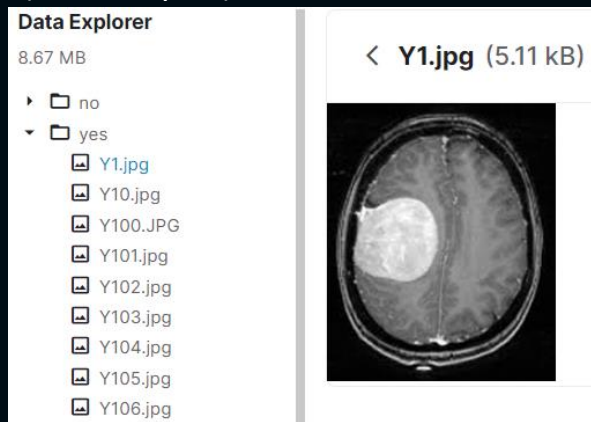
- A brain tumor occurs when abnormal cells form within the brain.
- There are 2 main types of tumors:
 - I. Cancerous (malignant) tumors
 - II. Benign tumors.
- Cancerous tumors can be divided into primary tumors, which start within the brain, and secondary tumors, which have spread from elsewhere, known as brain metastasis tumors.
- All types of brain tumors may produce symptoms that vary depending on the part of the brain involved.
- These symptoms may include headaches, seizures, problems with vision, vomiting and mental changes.



1. Data Discovery & Formulation- cont.

- **About Data -**

- Data Source- Brain MRI Images for Brain Tumor Detection Dataset from Kaggle Open Source.
- It consists of MRI scans of two classes:
 - NO - no tumor, encoded as 0 (98 samples)
 - YES - tumor, encoded as 1 (155 samples)



1. Data Discovery & Formulation- cont.

- **Tools & Technologies-**

- Python Language
- Python libraries- Numpy, Pandas, Matplotlib, OpenCV, scikit-learn, plotly & tqdm
- Python Utility Modules- os, shutil, itertools & imutils
- Python-based frameworks- keras & tensorflow
- Python IDE- Jupyter Notebook with Anaconda
- CNN- Neural Network Model for Images
- VGG-16 – CNN Architecture

- **Target People-** Hospital Director Board, Clinicians and Physicians- Directly and Patients- indirectly

2. Data Preparation & Processing

- Already, our required Data is in digital form from MRI Scan (we have information in digital).
- We are using Kaggle data. So, Already collection, processing and cleaning parts are done. So, we are using Data Acquisition method for data collection.
- Finally, we have our data in 2 folders for this Project which are tumor and no tumor.

3. Design a Model

- In this phase, we will discuss each step by step up to Model Building phase.
- In here, we will explore how, we prepare our data for building CNN model.

3. Design a Model- cont.

- **Creating Folder Directories**

- TRAIN, TEST and VAL with YES and NO Folders (9 Directories).
- TRAIN → 193 Images, TEST → 10 Images & VAL → 50 Images

```
TRAIN_DIR = 'TRAIN/'
TEST_DIR = 'TEST/'
VAL_DIR = 'VAL/'
IMG_SIZE = (224,224)

# use predefined function to load the image data into workspace
X_train, y_train, labels = load_data(TRAIN_DIR, IMG_SIZE)
X_test, y_test, _ = load_data(TEST_DIR, IMG_SIZE)
X_val, y_val, _ = load_data(VAL_DIR, IMG_SIZE)
```

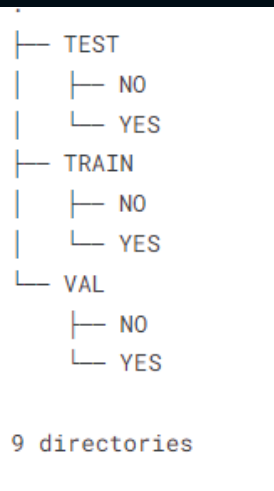
100% | 2/2 [00:02<00:00, 1.12s/it]
D:\Anaconda\lib\site-packages\ipykernel_launcher.py:18: VisibleDeprecationWarning:

Creating an ndarray from ragged nested sequences (which is a list-or-tuple of lists-or-tuples-or ndarrays with different lengths or shapes) is deprecated. If you meant to do this, you must specify 'dtype=object' when creating the ndarray.

193 images loaded from TRAIN/ directory.

100% | 2/2 [00:00<00:00, 34.41it/s]
10 images loaded from TEST/ directory.

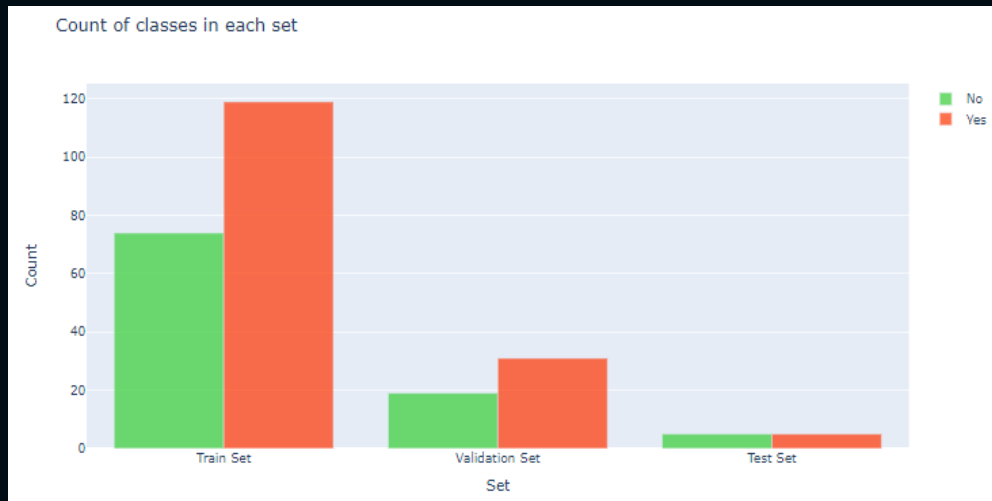
100% | 2/2 [00:00<00:00, 4.73it/s]
50 images loaded from VAL/ directory.



3. Design a Model- cont.

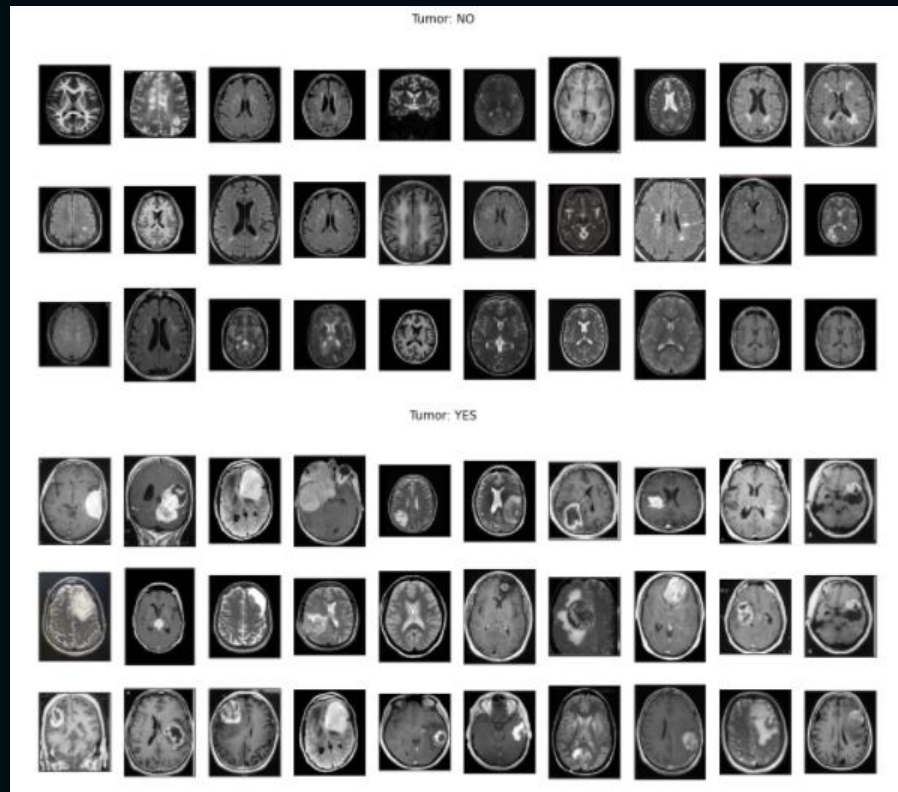
- **Plot the of Count Directories-**

- TRAIN, TEST and VAL with YES and NO Folders (9 Directories).
- TRAIN→ Yes-119 Images & NO- 74 Images
- VAL→ Yes-31 Images & NO- 19 Images
- TEST→ Yes-5 Images & NO- 5 Images



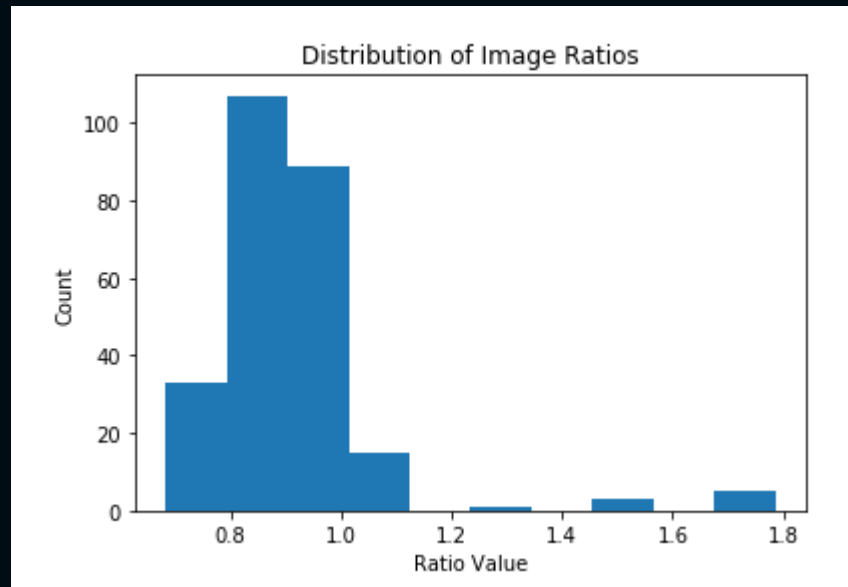
3. Design a Model- cont.

- **Plot for 30 images from the specified set (YES & NO)**
 - As we can see, images have different width and height and different size of "black corners".
 - Since the image size for VGG-16 input layer is (224,224)
 - some wide images may look weird after resizing.



3. Design a Model- cont.

- **Plot the Distribution of Image Ratios**
 - Histogram of ratio distributions (ratio = width/height)
 - So, to solve this problem as a first step of "normalization" would be to crop the brain out of the images.



3. Design a Model- cont.

- **Crop the Images**

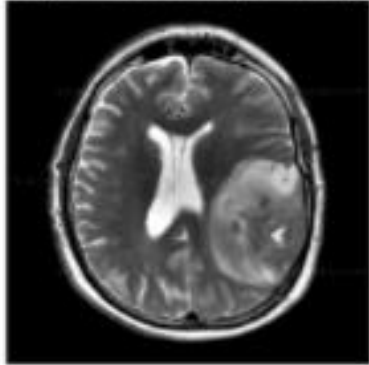
- Finds the extreme points on the image and crops the rectangular out of them.
- We do some steps to crop the images. They are;
 - Change RGB (Color) Image to Grey Image
 - Then, We Apply Gaussian Blur Filter (threshold the image)
 - Then, we Apply Erosion and Dilation filters (to remove any small regions of noise)
 - Next, find contours in threshold image, then grab the largest one
 - Finally, find the extreme points.
- Apply this crop function with one specific image

3. Design a Model- cont.

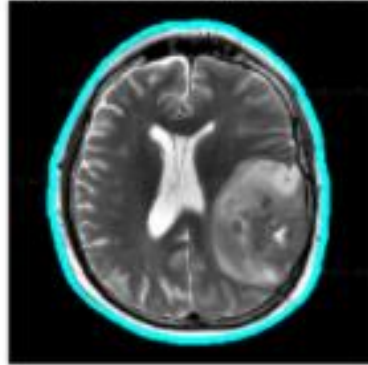
- **Crop the Images- cont.**

- Plot that image with 4 stages. They are the original image, Find the biggest contour, Find the extreme points and Crop the image.

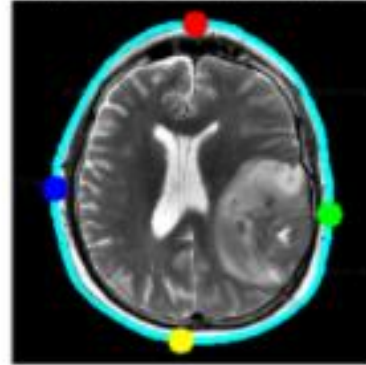
Step 1. Get the original image



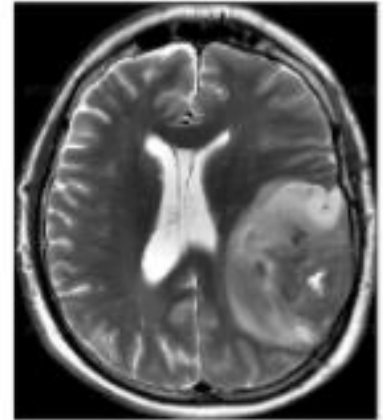
Step 2. Find the biggest contour



Step 3. Find the extreme points

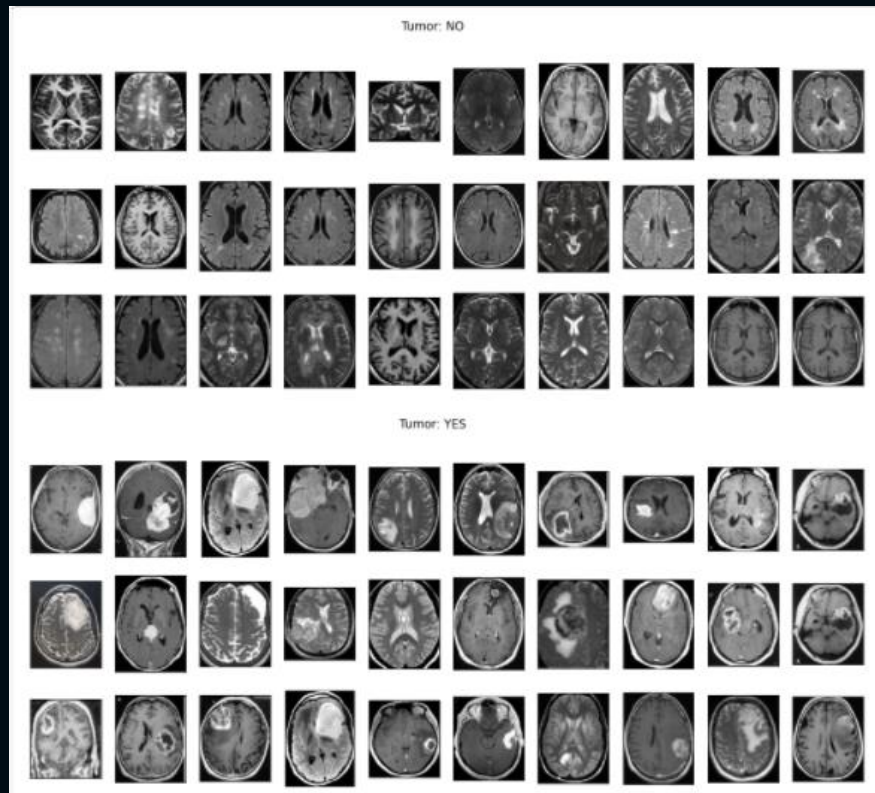


Step 4. Crop the image



3. Design a Model- cont.

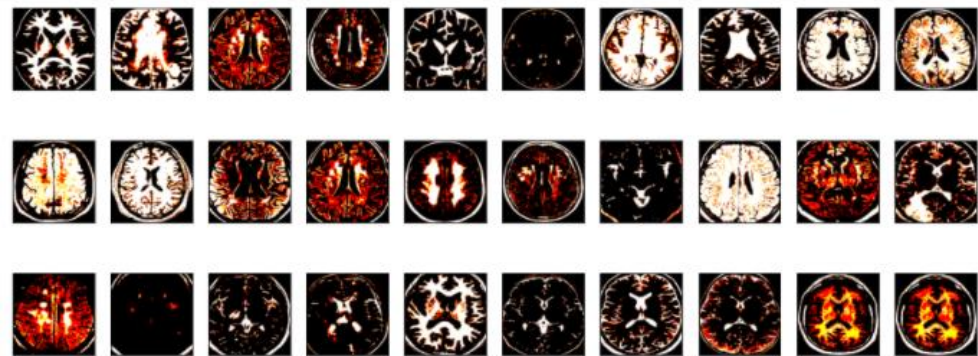
- **Crop the Images- cont.**
 - Apply this crop function to all the images in TRAIN, TEST and VAL directories.
 - Creating new directories called TRAIN_CROP, TEST_CROP, VAL_CROP with YES and NO directories and saving the new cropped images in them.



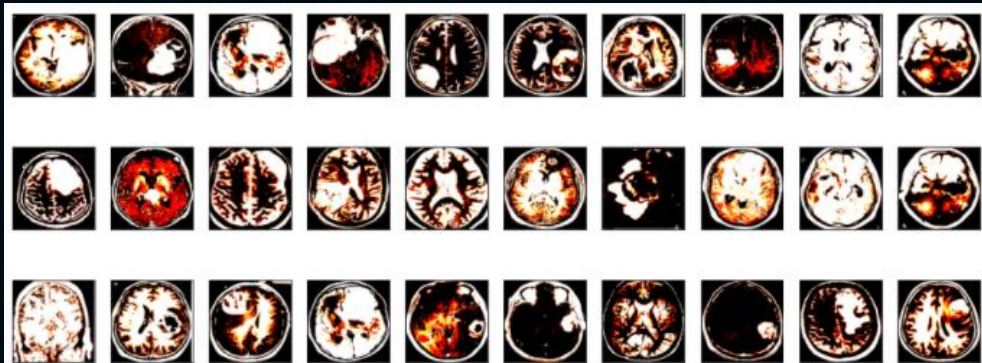
3. Design a Model- cont.

- **Resize the images for VGG-16**
 - We have to resize the images for $224 * 224$ size.
 - Then, Plot the images.

NO



YES



3. Design a Model- cont.

- **Data Augmentation**

- we used many parameter for this. They are:

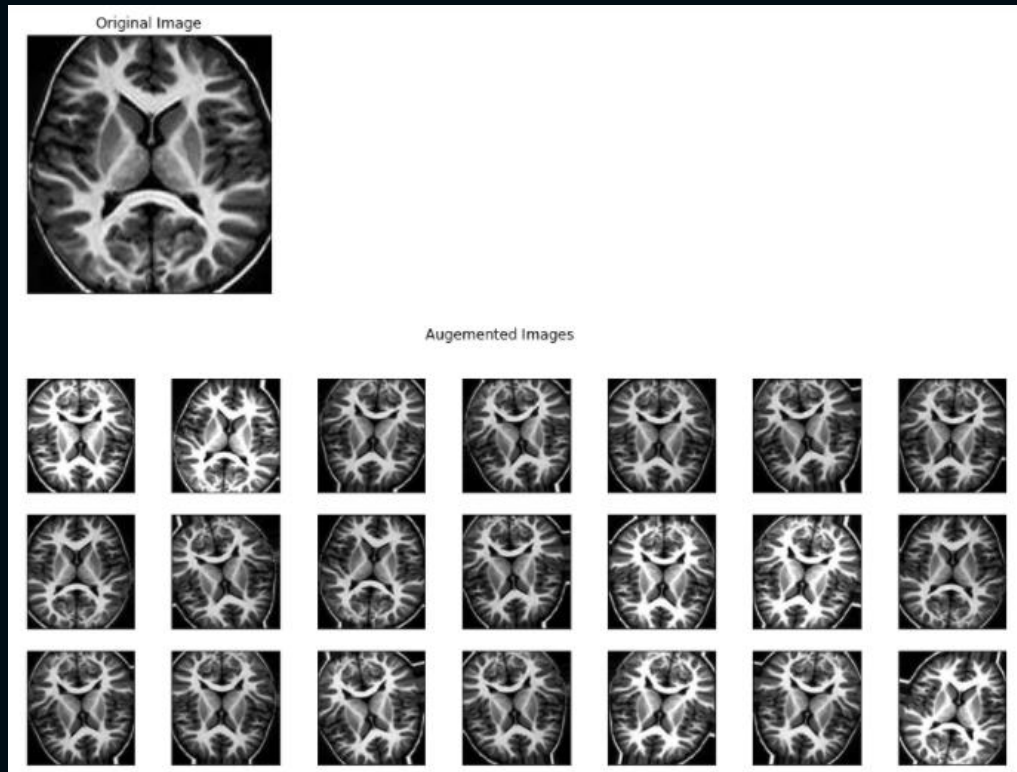
- Rotation range
 - Width shift range
 - Height shift range
 - Rescale
 - Shear range
 - Brightness range
 - Horizontal flip
 - Vertical flip

```
In [32]: # set the paramters we want to change randomly
demo_datagen = ImageDataGenerator(
    rotation_range=15,
    width_shift_range=0.05,
    height_shift_range=0.05,
    rescale=1./255,
    shear_range=0.05,
    brightness_range=[0.1, 1.5],
    horizontal_flip=True,
    vertical_flip=True
)|
```

3. Design a Model- cont.

- **Data Augmentation- cont.**

- Applying this function to one image and plotting after transforming images.
- Then, saving them in 'preview' directory (creating).
- Then, applying same function for whole images in TRAIN_CROP & VAL_CROP.



3. Design a Model- cont.

- **Data Augmentation- cont.**

- Finally, original images were transformed in TRAIN_CROP & VAL_CROP.
- So, TRAIN_CROP has 193 images with 2 classes and VAL_CROP has 50 images with 2 classes.

```
In [35]: TRAIN_DIR = 'TRAIN_CROP/'
         VAL_DIR = 'VAL_CROP/'

         train_datagen = ImageDataGenerator(
             rotation_range=15,
             width_shift_range=0.1,
             height_shift_range=0.1,
             shear_range=0.1,
             brightness_range=[0.5, 1.5],
             horizontal_flip=True,
             vertical_flip=True,
             preprocessing_function=preprocess_input
         )

         test_datagen = ImageDataGenerator(
             preprocessing_function=preprocess_input
         )

         train_generator = train_datagen.flow_from_directory(
             TRAIN_DIR,
             color_mode='rgb',
             target_size=IMG_SIZE,
             batch_size=32,
             class_mode='binary',
             seed=RANDOM_SEED
         )

         validation_generator = test_datagen.flow_from_directory(
             VAL_DIR,
             color_mode='rgb',
             target_size=IMG_SIZE,
             batch_size=16,
             class_mode='binary',
             seed=RANDOM_SEED
         )
```

Found 193 images belonging to 2 classes.
Found 50 images belonging to 2 classes.

3. Design a Model- cont.

- **Model Selection-**

- We can use many transfer learning techniques and CNN from the scratch.
- But, we used VGG16 because, VGG16 is used in many deep learning image classification problems and it is a great building block for learning purpose as it is easy to implement.
- So, In our Model we have used VGG-16 and CNN to solve classification problem.

4. Model Building

- In this phase,
 - First, we load VGG-16 weights and build this architecture
 - Then, we have created Feed Forward Network (Final layer) of CNN

4. Model Building- cont.

- First Step (VGG-16)-

```
In [36]: # Load base model
vgg16_weight_path = 'C:/Users/Dell/Desktop/Brain_Tumor_Detection/vgg16_weights_tf_dim_ordering_tf_kernels_notop.h5'
base_model = VGG16(
    weights=vgg16_weight_path,
    include_top=False,
    input_shape=IMG_SIZE + (3,)
)
```

4. Model Building- cont.

- **Second Step (CNN- FFN)-**
 - VGG-16 Base Model
 - Flatten layer
 - Dropout layer
 - Dense layer
 - Model Compiling
 - Model Summary

```
In [38]: NUM_NURON = 1

model = Sequential()
model.add(base_model)
model.add(layers.Flatten())
model.add(layers.Dropout(0.5))
model.add(layers.Dense(NUM_NURON, activation='sigmoid'))

model.layers[0].trainable = False

model.compile(
    loss='binary_crossentropy',
    optimizer=RMSprop(lr=1e-4),
    metrics=['accuracy']
)

model.summary()
```

Model: "sequential_1"

Layer (type)	Output Shape	Param #
=====		
vgg16 (Functional)	(None, 7, 7, 512)	14714688
flatten_1 (Flatten)	(None, 25088)	0
dropout_1 (Dropout)	(None, 25088)	0
dense_1 (Dense)	(None, 1)	25089

```
=====
Total params: 14,739,777
Trainable params: 25,089
Non-trainable params: 14,714,688
```

4. Model Building- cont.

- **Second Step (CNN- FFN)- cont.-**

- Model Fit

- Epochs- 30
- Steps per epochs- 6
- Validation steps- 3,
- Early Stopping

```
In [47]: EPOCHS = 30
es = EarlyStopping(
    monitor='val_accuracy',
    mode='max',
    patience=6
)

history = model.fit_generator(
    train_generator,
    steps_per_epoch=6,
    epochs=EPOCHS,
    validation_data=validation_generator,
    validation_steps=3,
    callbacks=[es]
)
```

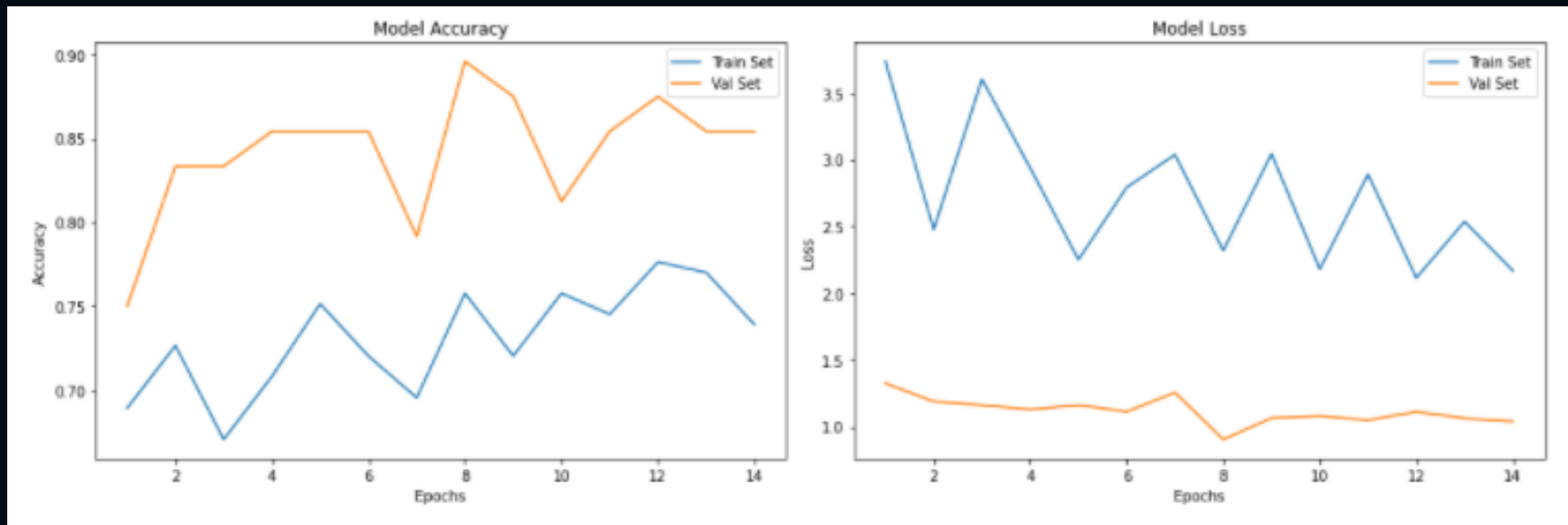
```
Epoch 1/30
6/6 [=====] - 62s 10s/step - loss: 3.7390 - accuracy: 0.6894 - val_loss: 1.3249 - val_accuracy: 0.7500
Epoch 2/30
6/6 [=====] - 69s 12s/step - loss: 2.4777 - accuracy: 0.7267 - val_loss: 1.1888 - val_accuracy: 0.8333
Epoch 3/30
6/6 [=====] - 72s 12s/step - loss: 3.6052 - accuracy: 0.6708 - val_loss: 1.1622 - val_accuracy: 0.8333
Epoch 4/30
6/6 [=====] - 82s 14s/step - loss: 2.9390 - accuracy: 0.7083 - val_loss: 1.1284 - val_accuracy: 0.8542
Epoch 5/30
6/6 [=====] - 67s 11s/step - loss: 2.2526 - accuracy: 0.7516 - val_loss: 1.1618 - val_accuracy: 0.8542
Epoch 6/30
6/6 [=====] - 71s 12s/step - loss: 2.7945 - accuracy: 0.7205 - val_loss: 1.1129 - val_accuracy: 0.8542
Epoch 7/30
6/6 [=====] - 69s 12s/step - loss: 3.0394 - accuracy: 0.6957 - val_loss: 1.2556 - val_accuracy: 0.7917
Epoch 8/30
6/6 [=====] - 61s 10s/step - loss: 2.3205 - accuracy: 0.7578 - val_loss: 0.9040 - val_accuracy: 0.8958
Epoch 9/30
6/6 [=====] - 61s 10s/step - loss: 3.0458 - accuracy: 0.7205 - val_loss: 1.0647 - val_accuracy: 0.8750
Epoch 10/30
6/6 [=====] - 61s 10s/step - loss: 2.1817 - accuracy: 0.7578 - val_loss: 1.0796 - val_accuracy: 0.8125
Epoch 11/30
6/6 [=====] - 63s 12s/step - loss: 2.8925 - accuracy: 0.7453 - val_loss: 1.0494 - val_accuracy: 0.8542
Epoch 12/30
6/6 [=====] - 63s 11s/step - loss: 2.1157 - accuracy: 0.7764 - val_loss: 1.1112 - val_accuracy: 0.8750
Epoch 13/30
6/6 [=====] - 66s 11s/step - loss: 2.5382 - accuracy: 0.7702 - val_loss: 1.0636 - val_accuracy: 0.8542
Epoch 14/30
6/6 [=====] - 63s 11s/step - loss: 2.1697 - accuracy: 0.7391 - val_loss: 1.0387 - val_accuracy: 0.8542
```

5. Results

- In this phase,
 - We have checked the performance of the model based on Model Accuracy and Loss.
 - Then, we checked from confusion matrix on validation and test set.

5. Results- cont.

- Accuracy & Loss Graphs-

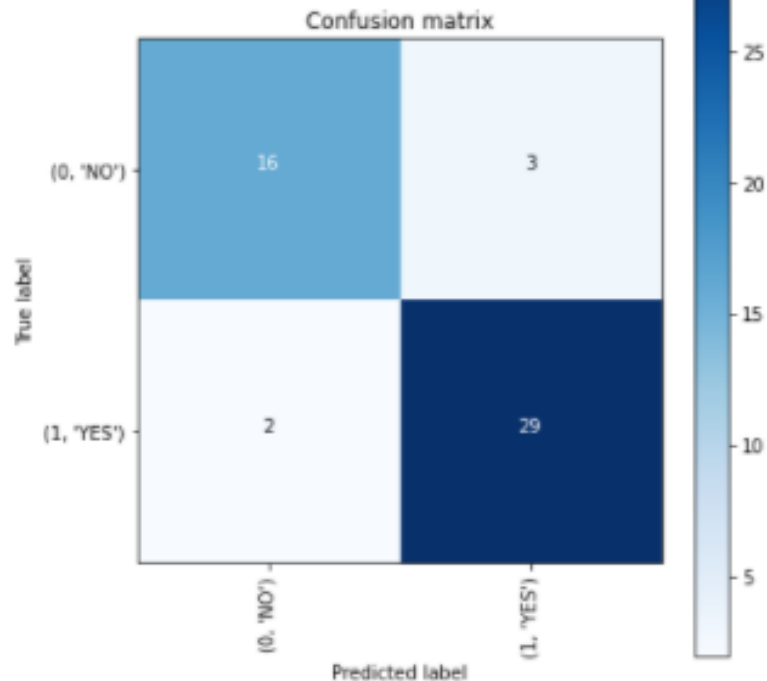


5. Results- cont.

- **Confusion Matrixes-**

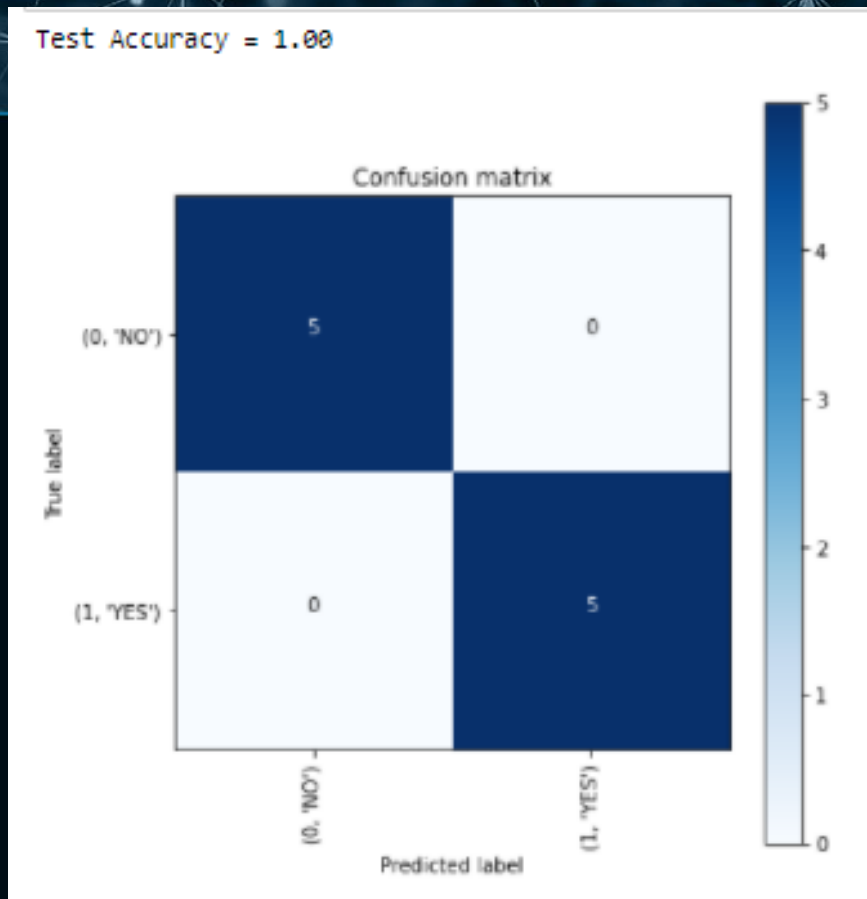
- Validate on validation set-
Validation Accuracy = 90%
- 50 Samples
- FP=3, FN=2, TN=16 and TP=29.

Val Accuracy = 0.90



5. Results- cont.

- **Confusion Matrixes- cont.**
 - Validate on test set- Test Accuracy = 100%
 - 10 Samples
 - FP and FN are 0. TN and TP are classified as 5.



5. Results- cont.

- **Checking misclassification in test set**
 - Test Accuracy is 100%. So, there is no misclassification.

```
In [51]: ind_list = np.argwhere((y_test == predictions) == False)[:, -1]
if ind_list.size == 0:
    print('There are no missclassified images.')
else:
    for i in ind_list:
        plt.figure()
        plt.imshow(X_test_crop[i])
        plt.xticks([])
        plt.yticks([])
        plt.title(f'Actual class: {y_val[i]}\nPredicted class: {predictions[i]}')
        plt.show()
```

There are no missclassified images.

5. Results- cont.

- **Saving the Model**

- We can reuse this model from these weights.

```
In [52]: # save the model  
model.save('2022-03-07-1pm_VGG_model.h5')
```

5. Results- cont.

- **Vital Findings from these Results**

- Accuracy of Validation higher than training data.
- Loss of Validation lesser than training data.
- Validation Accuracy is 90%. But, test accuracy is 100%. So, Unseen data can predict more accurate.
- When, we automate this model with MRI Scan images it will predict most probably right decision which classify correctly the patient is cancer or not.
- This model will improve business (Hospital) value by reducing the wasting time for manual process and don't need human intervention.

6. Measuring of effeteness

- This Model is more generalized to unseen data than seen data.
- We can integrate this model with MRI scan and when the patient taking the Scan itself it will predict the automatically, which the patient has cancer or not.
- Then, Patients(Customers) satisfaction will increase towards the hospital(Business). They don't need to waste another time for get report.

6. Measuring of effeteness

- Patients can save their time and cost.
- If it is urgent(cancer), They can immediately take any actions rather than waiting many days for reports. Then, we can save many lives.
- Instead of Doctors are wasting their precious time on viewing Scans as a manual process; they can spend their time for other human involving treatments. This model can automate this task.
- This model will improve business (Hospital) value by reducing the time and cost of physicians and patients. So, Hospitals will attract more patients.
- Code GitHub Repository- https://github.com/Farha-Imthiyaz/Tumor_Classification

THANK
YOU

