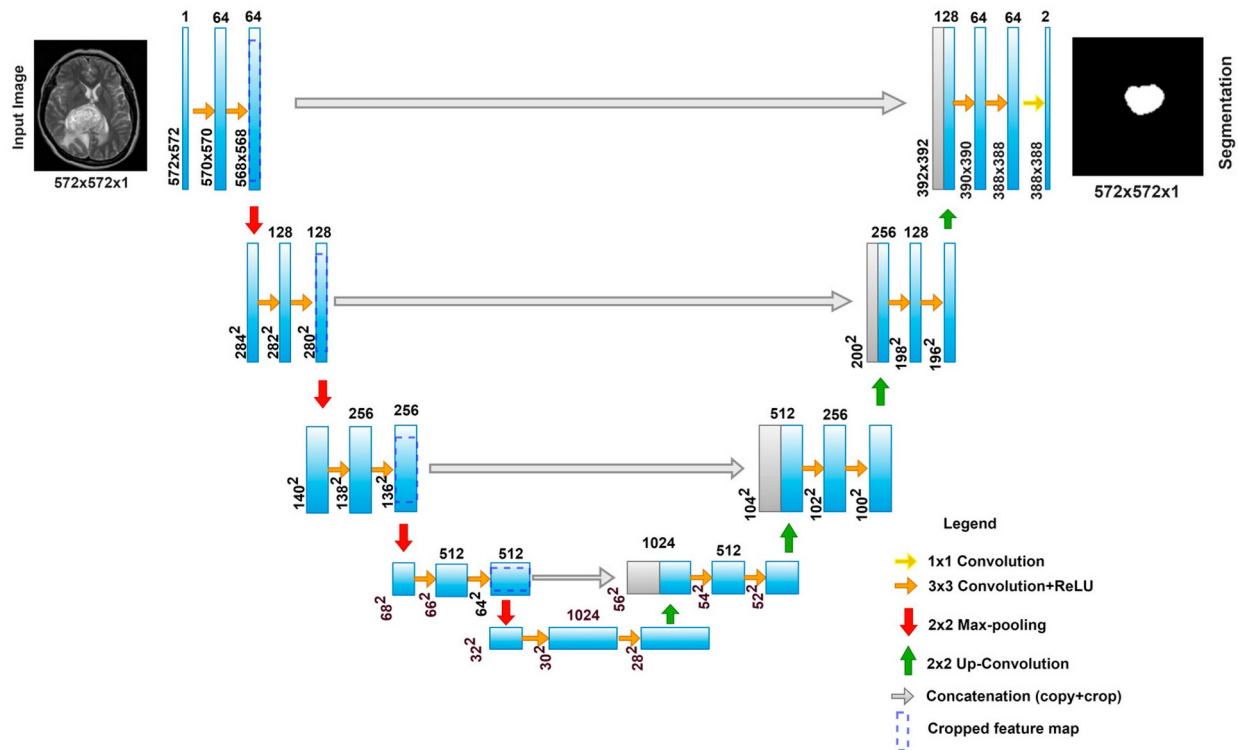


# Brain tumor segmentation with UNET



Steps to perform brain tumor segmentation using the U-Net architecture:

1. **Data Collection and Preparation:**
  - Gather a dataset of brain MRI images along with corresponding ground truth masks indicating tumor regions.
  - Preprocess the images (e.g., resizing, normalization) to ensure uniformity and enhance model performance.
2. **Data Augmentation:**
  - Augment the dataset to increase its size and diversity, which helps in improving the robustness and generalization of the model.
  - Common augmentation techniques include rotation, flipping, scaling, and adding noise to the images.
3. **Splitting the Dataset:**
  - Divide the dataset into training, validation, and testing sets. The training set is used to train the model, the validation set is used to tune hyperparameters and monitor training progress, and the testing set is used to evaluate the model's performance.
4. **Building the U-Net Model:**
  - Implement the U-Net architecture, which consists of an encoder (downsampling path) and a decoder (upsampling path) connected by skip connections.

- Use convolutional layers with appropriate activation functions (e.g., ReLU), pooling layers (e.g., max pooling), and upsampling layers (e.g., transposed convolution) to build the network.
5. **Loss Function Selection:**
    - Choose an appropriate loss function for training the model. For binary segmentation tasks like tumor segmentation, commonly used loss functions include binary cross-entropy loss, Dice loss, and Jaccard loss.
  6. **Training the Model:**
    - Train the U-Net model on the training dataset using the selected loss function and an optimization algorithm (e.g., Adam optimizer).
    - Monitor the model's performance on the validation set and adjust hyperparameters if necessary (e.g., learning rate, batch size) to improve performance and prevent overfitting.
    - Train the model until convergence or until the validation loss stops decreasing.
  7. **Evaluation:**
    - Evaluate the trained model on the testing set to assess its performance in segmenting brain tumors.
    - Compute evaluation metrics such as Dice coefficient, Jaccard index, sensitivity, specificity, and accuracy to quantify the model's performance.
  8. **Post-processing:**
    - Perform post-processing techniques (e.g., morphological operations like dilation and erosion) to refine the segmentation masks and improve the segmentation results.
  9. **Visualization:**
    - Visualize the segmented tumor regions overlaid on the original MRI images to visually inspect the quality of the segmentation results.
  10. **Deployment:**
    - Deploy the trained model for practical use, such as automating tumor segmentation in clinical settings or integrating it into medical image analysis software.

```
import os
import random
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
%matplotlib inline
from mpl_toolkits.axes_grid1 import ImageGrid

import cv2
from glob import glob

import tensorflow as tf
from tensorflow.keras import Input
from tensorflow.keras.models import Model, load_model, save_model
from tensorflow.keras.layers import Input, Activation,
BatchNormalization, Dropout, Lambda, Conv2D
from tensorflow.keras.layers import Conv2DTranspose, MaxPooling2D,
```

```

concatenate, AveragePooling2D, Dense, Flatten
from tensorflow.keras.optimizers import Adam
from tensorflow.keras.callbacks import EarlyStopping, ModelCheckpoint,
ReduceLROnPlateau

from tensorflow.keras import backend as K
from tensorflow.keras.preprocessing.image import ImageDataGenerator

# Set parameters
IMAGE_SIZE = (256, 256)

```

## Create DataFrame

```

mask_files = glob('../input/lgg-mri-segmentation/kaggle_3m/*/*_mask*')
train_files = [file.replace('_mask', '') for file in mask_files]

def diagnosis(mask_path):
    value = np.max(cv2.imread(mask_path))
    return '1' if value > 0 else '0'
df = pd.DataFrame({"image_path": train_files,
                   "mask_path": mask_files,
                   "diagnosis": [diagnosis(x) for x in mask_files]})
df.head()

```

```

                                image_path \
0  ../input/lgg-mri-segmentation/kaggle_3m/TCGA_D...
1  ../input/lgg-mri-segmentation/kaggle_3m/TCGA_D...
2  ../input/lgg-mri-segmentation/kaggle_3m/TCGA_D...
3  ../input/lgg-mri-segmentation/kaggle_3m/TCGA_D...
4  ../input/lgg-mri-segmentation/kaggle_3m/TCGA_D...

```

```

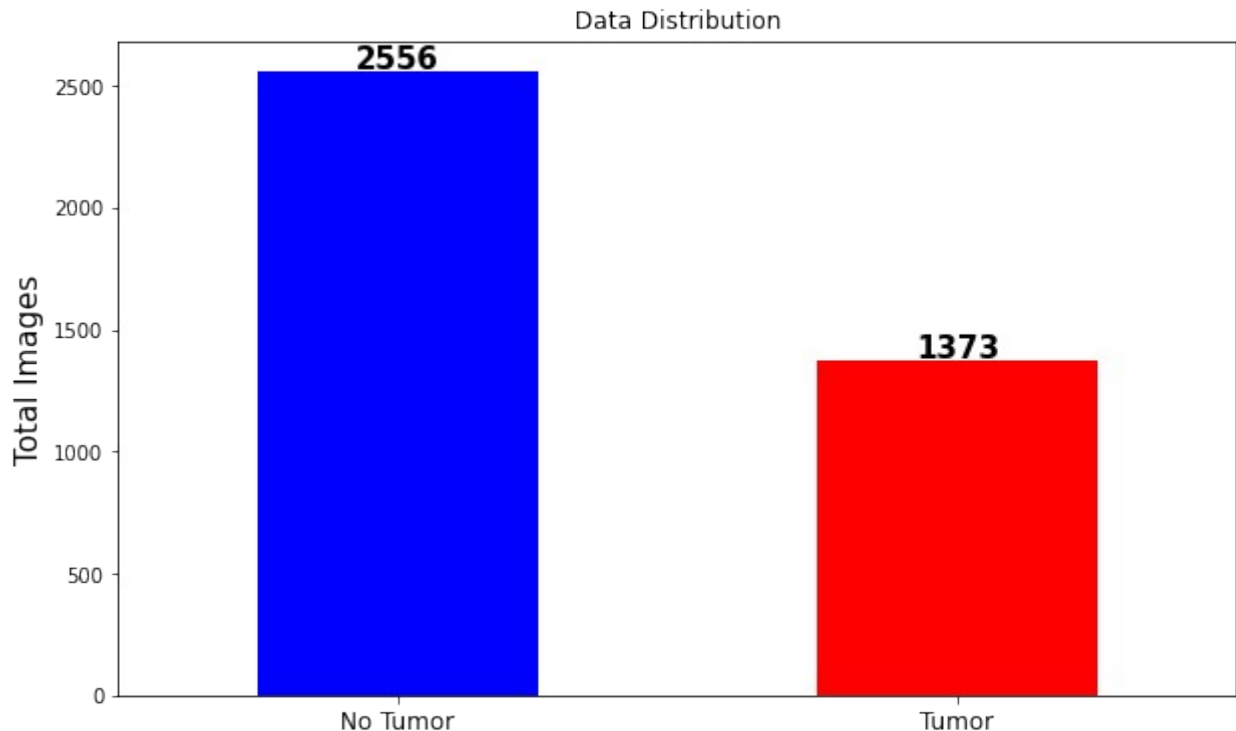
                                mask_path diagnosis
0  ../input/lgg-mri-segmentation/kaggle_3m/TCGA_D...      0
1  ../input/lgg-mri-segmentation/kaggle_3m/TCGA_D...      0
2  ../input/lgg-mri-segmentation/kaggle_3m/TCGA_D...      1
3  ../input/lgg-mri-segmentation/kaggle_3m/TCGA_D...      1
4  ../input/lgg-mri-segmentation/kaggle_3m/TCGA_D...      1

```

```

ax = df['diagnosis'].value_counts().plot(kind='bar', stacked=True,
figsize=(10,6), color=['blue', 'red'])
ax.set_title('Data Distribution')
ax.set_ylabel('Total Images', fontsize=15)
ax.set_xticklabels(['No Tumor', 'Tumor'], fontsize=12, rotation=0)
for i, rows in enumerate(df['diagnosis'].value_counts().values):
    ax.annotate(int(rows), xy=(i, rows+12), ha='center',
fontweight='bold', fontsize=15)

```



## Visualize MRI with Mask

```
df_positive = df[df['diagnosis']=='1'].sample(5).values
df_negative = df[df['diagnosis']=='0'].sample(5).values

def show_data(df, positive=True):
    images = []
    masks = []
    for data in df:
        img = cv2.imread(data[0])
        mask = cv2.imread(data[1])
        images.append(img)
        masks.append(mask)
    images = np.hstack(np.array(images))
    masks = np.hstack(np.array(masks))

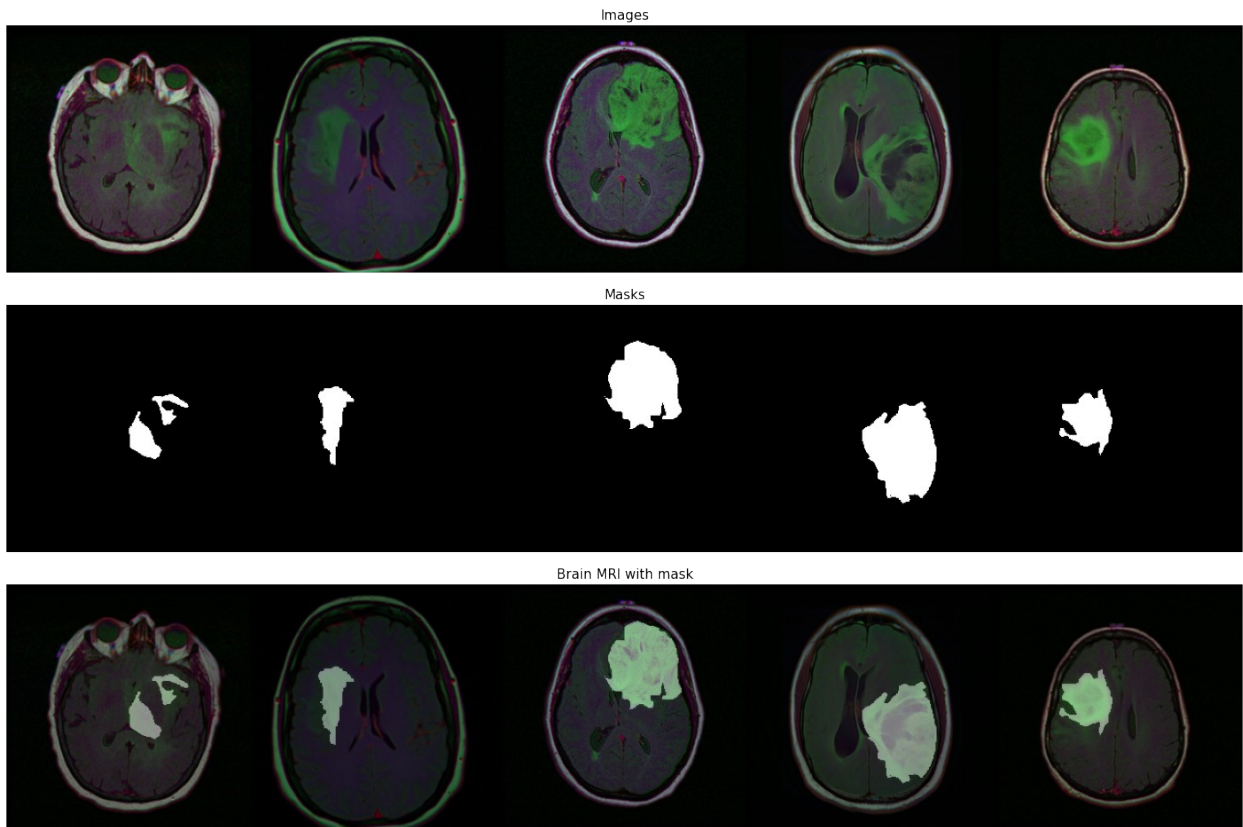
    fig = plt.figure(figsize=(25,25))
    if positive:
        grid = ImageGrid(fig, 111, nrows_ncols=(3,1), axes_pad=0.5)
    else:
        grid = ImageGrid(fig, 111, nrows_ncols=(2,1), axes_pad=0.5)
    grid[0].imshow(images)
    grid[0].set_title('Images', fontsize=15)
    grid[0].axis('off')
    grid[1].imshow(masks)
    grid[1].set_title('Masks', fontsize=15)
    grid[1].axis('off')
```

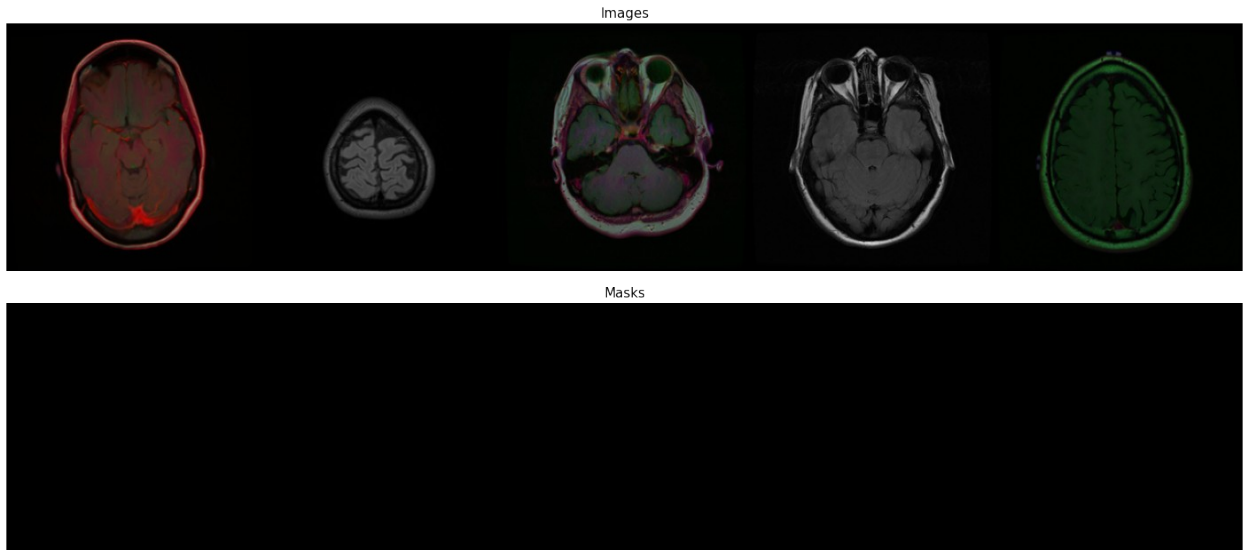
```

if positive:
    grid[2].imshow(images)
    grid[2].imshow(masks, alpha=0.4)
    grid[2].set_title('Brain MRI with mask', fontsize=15)
    grid[2].axis('off')

show_data(df_positive)
show_data(df_negative, positive=False)

```





## Split data into Train, Validation and Test Set

```
from sklearn.model_selection import train_test_split
df_train, df_test = train_test_split(df, test_size=0.15)
df_train, df_val = train_test_split(df_train, test_size=0.15)
print(df_train.values.shape)
print(df_val.values.shape)
print(df_test.values.shape)

(2838, 3)
(501, 3)
(590, 3)
```

## U-Net Model

### Data Generator, Data Augmentation and Adjust Data

```
def train_generator(data_frame, batch_size, aug_dict,
                    image_color_mode="rgb",
                    mask_color_mode="grayscale",
                    image_save_prefix="image",
                    mask_save_prefix="mask",
                    save_to_dir=None,
                    target_size=(256,256),
                    seed=1):

    image_datagen = ImageDataGenerator(**aug_dict)
    mask_datagen = ImageDataGenerator(**aug_dict)

    image_generator = image_datagen.flow_from_dataframe(
        data_frame,
        x_col = "image_path",
        class_mode = None,
```

```

        color_mode = image_color_mode,
        target_size = target_size,
        batch_size = batch_size,
        save_to_dir = save_to_dir,
        save_prefix = image_save_prefix,
        seed = seed)

mask_generator = mask_datagen.flow_from_dataframe(
    data_frame,
    x_col = "mask_path",
    class_mode = None,
    color_mode = mask_color_mode,
    target_size = target_size,
    batch_size = batch_size,
    save_to_dir = save_to_dir,
    save_prefix = mask_save_prefix,
    seed = seed)

train_gen = zip(image_generator, mask_generator)

for (img, mask) in train_gen:
    img, mask = adjust_data(img, mask)
    yield (img, mask)

def adjust_data(img, mask):
    img = img / 255.
    mask = mask / 255.
    mask[mask > 0.5] = 1
    mask[mask <= 0.5] = 0

    return (img, mask)

smooth=1.

def dice_coef(y_true, y_pred):
    y_true = K.flatten(y_true)

    y_pred = K.flatten(y_pred)
    intersection = K.sum(y_true * y_pred)
    union = K.sum(y_true) + K.sum(y_pred)
    return (2.0 * intersection + smooth) / (union + smooth)

def dice_coef_loss(y_true, y_pred):
    return 1 - dice_coef(y_true, y_pred)

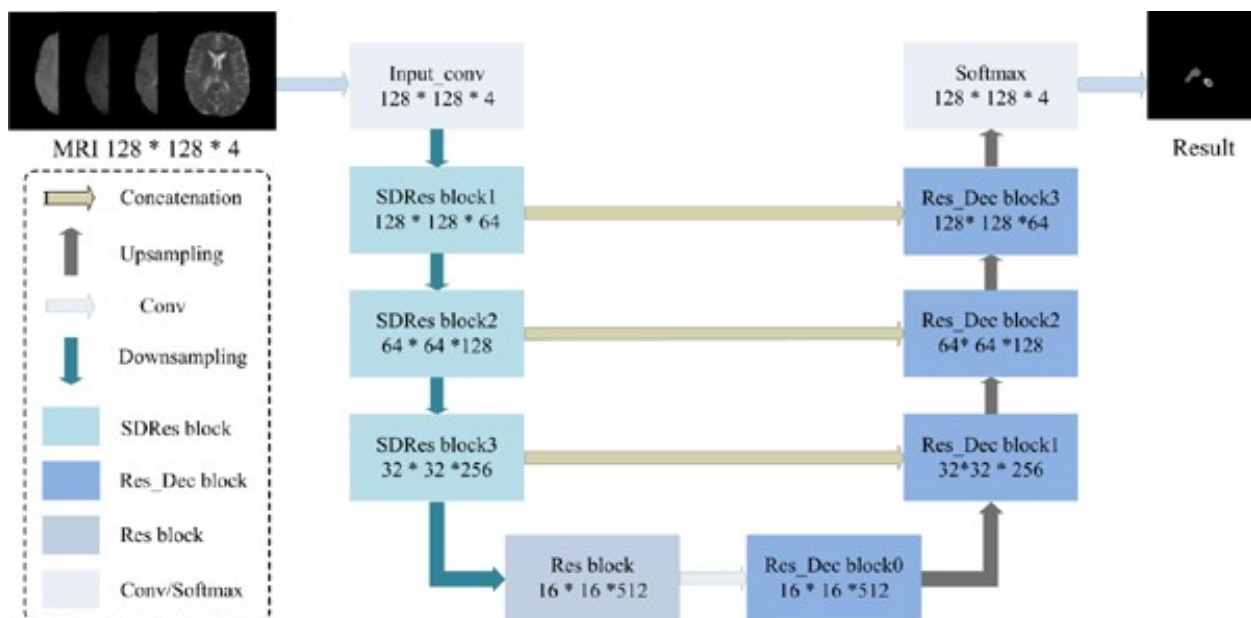
def bce_dice_loss(y_true, y_pred):
    bce = tf.keras.losses.BinaryCrossentropy(from_logits=True)
    return dice_coef_loss(y_true, y_pred) + bce(y_true, y_pred)

```

```
def iou(y_true, y_pred):
    intersection = K.sum(y_true * y_pred)
    sum_ = K.sum(y_true + y_pred)
    jac = (intersection + smooth) / (sum_ - intersection + smooth)
    return jac
```

## UNet Model Architecture

### Working and Flow Diagram of UNet





```

def unet(input_size=(256,256,3)):
    inputs = Input(input_size)

    conv1 = Conv2D(64, (3, 3), padding='same')(inputs)
    bn1 = Activation('relu')(conv1)
    conv1 = Conv2D(64, (3, 3), padding='same')(bn1)
    bn1 = BatchNormalization(axis=3)(conv1)
    bn1 = Activation('relu')(bn1)
    pool1 = MaxPooling2D(pool_size=(2, 2))(bn1)

    conv2 = Conv2D(128, (3, 3), padding='same')(pool1)
    bn2 = Activation('relu')(conv2)
    conv2 = Conv2D(128, (3, 3), padding='same')(bn2)
    bn2 = BatchNormalization(axis=3)(conv2)
    bn2 = Activation('relu')(bn2)
    pool2 = MaxPooling2D(pool_size=(2, 2))(bn2)

    conv3 = Conv2D(256, (3, 3), padding='same')(pool2)
    bn3 = Activation('relu')(conv3)
    conv3 = Conv2D(256, (3, 3), padding='same')(bn3)
    bn3 = BatchNormalization(axis=3)(conv3)
    bn3 = Activation('relu')(bn3)
    pool3 = MaxPooling2D(pool_size=(2, 2))(bn3)

    conv4 = Conv2D(512, (3, 3), padding='same')(pool3)
    bn4 = Activation('relu')(conv4)
    conv4 = Conv2D(512, (3, 3), padding='same')(bn4)
    bn4 = BatchNormalization(axis=3)(conv4)
    bn4 = Activation('relu')(bn4)
    pool4 = MaxPooling2D(pool_size=(2, 2))(bn4)

    conv5 = Conv2D(1024, (3, 3), padding='same')(pool4)
    bn5 = Activation('relu')(conv5)
    conv5 = Conv2D(1024, (3, 3), padding='same')(bn5)
    bn5 = BatchNormalization(axis=3)(conv5)
    bn5 = Activation('relu')(bn5)

    up6 = concatenate([Conv2DTranspose(512, (2, 2), strides=(2, 2),
padding='same')(bn5), conv4], axis=3)
    conv6 = Conv2D(512, (3, 3), padding='same')(up6)
    bn6 = Activation('relu')(conv6)
    conv6 = Conv2D(512, (3, 3), padding='same')(bn6)
    bn6 = BatchNormalization(axis=3)(conv6)
    bn6 = Activation('relu')(bn6)

    up7 = concatenate([Conv2DTranspose(256, (2, 2), strides=(2, 2),
padding='same')(bn6), conv3], axis=3)
    conv7 = Conv2D(256, (3, 3), padding='same')(up7)
    bn7 = Activation('relu')(conv7)
    conv7 = Conv2D(256, (3, 3), padding='same')(bn7)

```

```

bn7 = BatchNormalization(axis=3)(conv7)
bn7 = Activation('relu')(bn7)

up8 = concatenate([Conv2DTranspose(128, (2, 2), strides=(2, 2),
padding='same')(bn7), conv2], axis=3)
conv8 = Conv2D(128, (3, 3), padding='same')(up8)
bn8 = Activation('relu')(conv8)
conv8 = Conv2D(128, (3, 3), padding='same')(bn8)
bn8 = BatchNormalization(axis=3)(conv8)
bn8 = Activation('relu')(bn8)

up9 = concatenate([Conv2DTranspose(64, (2, 2), strides=(2, 2),
padding='same')(bn8), conv1], axis=3)
conv9 = Conv2D(64, (3, 3), padding='same')(up9)
bn9 = Activation('relu')(conv9)
conv9 = Conv2D(64, (3, 3), padding='same')(bn9)
bn9 = BatchNormalization(axis=3)(conv9)
bn9 = Activation('relu')(bn9)

conv10 = Conv2D(1, (1, 1), activation='sigmoid')(bn9)

return Model(inputs=[inputs], outputs=[conv10])

```

## Training

```

# Set parameters
EPOCHS = 10
BATCH_SIZE = 32
learning_rate = 1e-4

train_generator_args = dict(rotation_range=0.1,
                             width_shift_range=0.05,
                             height_shift_range=0.05,
                             shear_range=0.05,
                             zoom_range=0.05,
                             horizontal_flip=True,
                             vertical_flip=True,
                             fill_mode='nearest')

train_gen = train_generator(df_train, BATCH_SIZE,
                             train_generator_args,
                             target_size=IMAGE_SIZE)

val_gen = train_generator(df_val, BATCH_SIZE,
                           dict(),
                           target_size=IMAGE_SIZE)

model = unet(input_size=(IMAGE_SIZE[0], IMAGE_SIZE[1], 3))

```

```
opt = Adam(lr=learning_rate, beta_1=0.9, beta_2=0.999, epsilon=None,
amsgrad=False)
model.compile(optimizer=opt, loss=bce_dice_loss, metrics=[iou,
dice_coef])
```

```
callbacks = [ModelCheckpoint('unet_brainMRI_seg.hdf5', verbose=0,
save_best_only=True),
             ReduceLROnPlateau(monitor='val_loss', factor=0.1,
patience=5, verbose=1, min_lr=1e-11),
             EarlyStopping(monitor='val_loss',
restore_best_weights=True, patience=15)]
```

```
history = model.fit(train_gen,
                    steps_per_epoch=len(df_train) / BATCH_SIZE,
                    epochs=EPOCHS,
                    callbacks=callbacks,
                    validation_data = val_gen,
                    validation_steps=len(df_val) / BATCH_SIZE)
```

Found 2838 validated image filenames.

Found 2838 validated image filenames.

Epoch 1/10

89/88 [=====] - ETA: 0s - loss: 1.1530 - iou: 0.0374 - dice\_coef: 0.0708

Found 501 validated image filenames.

Found 501 validated image filenames.

88/88 [=====] - 109s 1s/step - loss: 1.1517 - iou: 0.0377 - dice\_coef: 0.0713 - val\_loss: 1.4798 - val\_iou: 0.0104 - val\_dice\_coef: 0.0206

Epoch 2/10

88/88 [=====] - 70s 791ms/step - loss: 0.8467 - iou: 0.1279 - dice\_coef: 0.2246 - val\_loss: 1.1900 - val\_iou: 0.0100 - val\_dice\_coef: 0.0199

Epoch 3/10

88/88 [=====] - 69s 783ms/step - loss: 0.7537 - iou: 0.1780 - dice\_coef: 0.3000 - val\_loss: 1.0648 - val\_iou: 0.0100 - val\_dice\_coef: 0.0197

Epoch 4/10

88/88 [=====] - 69s 773ms/step - loss: 0.7083 - iou: 0.2065 - dice\_coef: 0.3386 - val\_loss: 0.9609 - val\_iou: 0.0515 - val\_dice\_coef: 0.0966

Epoch 5/10

88/88 [=====] - 69s 783ms/step - loss: 0.6353 - iou: 0.2540 - dice\_coef: 0.4003 - val\_loss: 0.6831 - val\_iou: 0.2139 - val\_dice\_coef: 0.3493

Epoch 6/10

88/88 [=====] - 69s 780ms/step - loss: 0.5807 - iou: 0.2964 - dice\_coef: 0.4510 - val\_loss: 0.5329 - val\_iou: 0.3403 - val\_dice\_coef: 0.5042

Epoch 7/10

88/88 [=====] - 70s 785ms/step - loss: 0.5229

```

- iou: 0.3484 - dice_coef: 0.5086 - val_loss: 0.5051 - val_iou: 0.3645
- val_dice_coef: 0.5298
Epoch 8/10
88/88 [=====] - 70s 792ms/step - loss: 0.4598
- iou: 0.3994 - dice_coef: 0.5660 - val_loss: 0.4120 - val_iou: 0.4446
- val_dice_coef: 0.6128
Epoch 9/10
88/88 [=====] - 69s 784ms/step - loss: 0.4599
- iou: 0.4095 - dice_coef: 0.5721 - val_loss: 0.4312 - val_iou: 0.4236
- val_dice_coef: 0.5892
Epoch 10/10
88/88 [=====] - 69s 774ms/step - loss: 0.4012
- iou: 0.4604 - dice_coef: 0.6230 - val_loss: 0.3164 - val_iou: 0.5466
- val_dice_coef: 0.7032

```

## Visualize the model performance

```

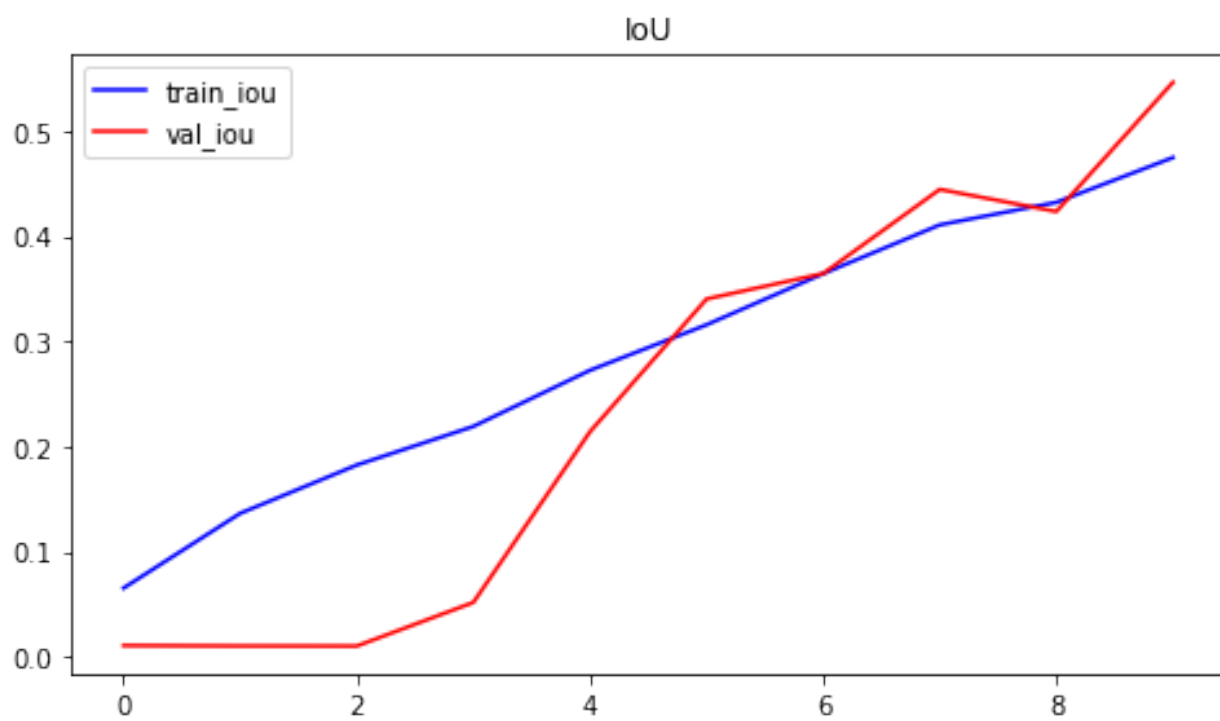
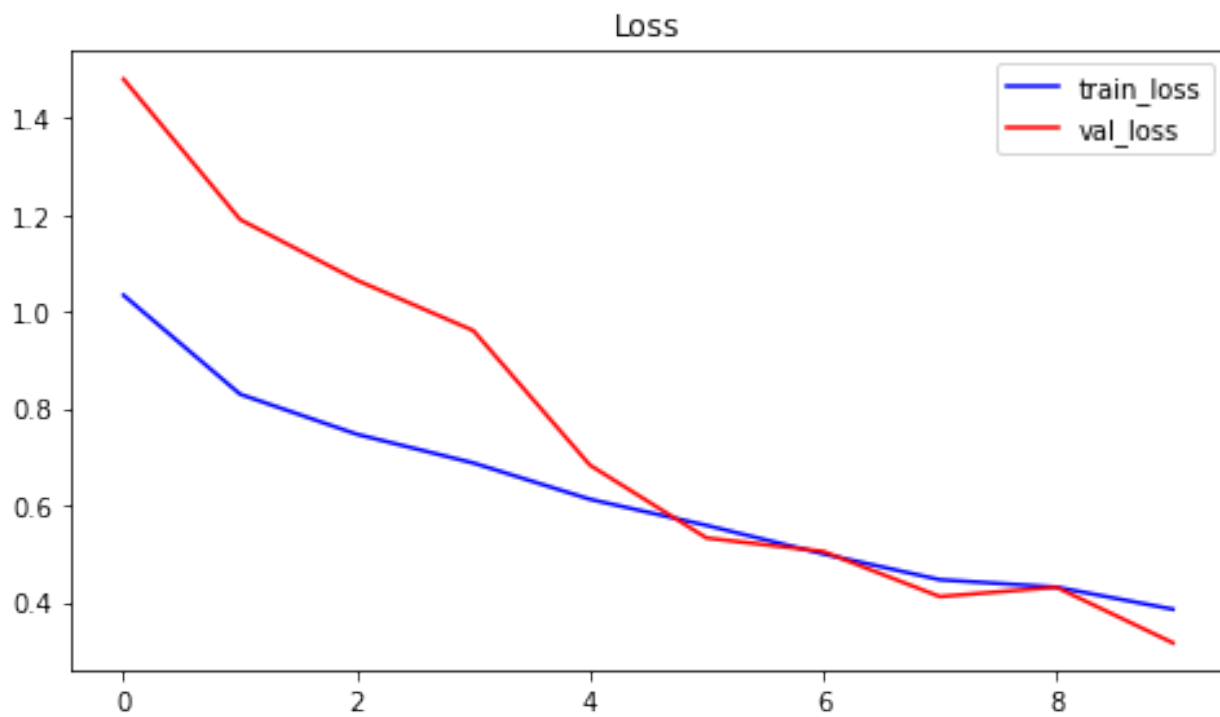
plt.figure(figsize=(8,15))
plt.subplot(3,1,1)
plt.plot(model.history.history['loss'], 'b-', label='train_loss')
plt.plot(model.history.history['val_loss'], 'r-', label='val_loss')
plt.legend(loc='best')
plt.title('Loss')

plt.subplot(3,1,2)
plt.plot(model.history.history['iou'], 'b-', label='train_iou')
plt.plot(model.history.history['val_iou'], 'r-', label='val_iou')
plt.legend(loc='best')
plt.title('IoU')

plt.subplot(3,1,3)
plt.plot(model.history.history['dice_coef'], 'b-',
label='train_dice_coef')
plt.plot(model.history.history['val_dice_coef'], 'r-',
label='val_dice_coef')
plt.legend(loc='best')
plt.title('Dice Coef')

Text(0.5, 1.0, 'Dice Coef')

```



```

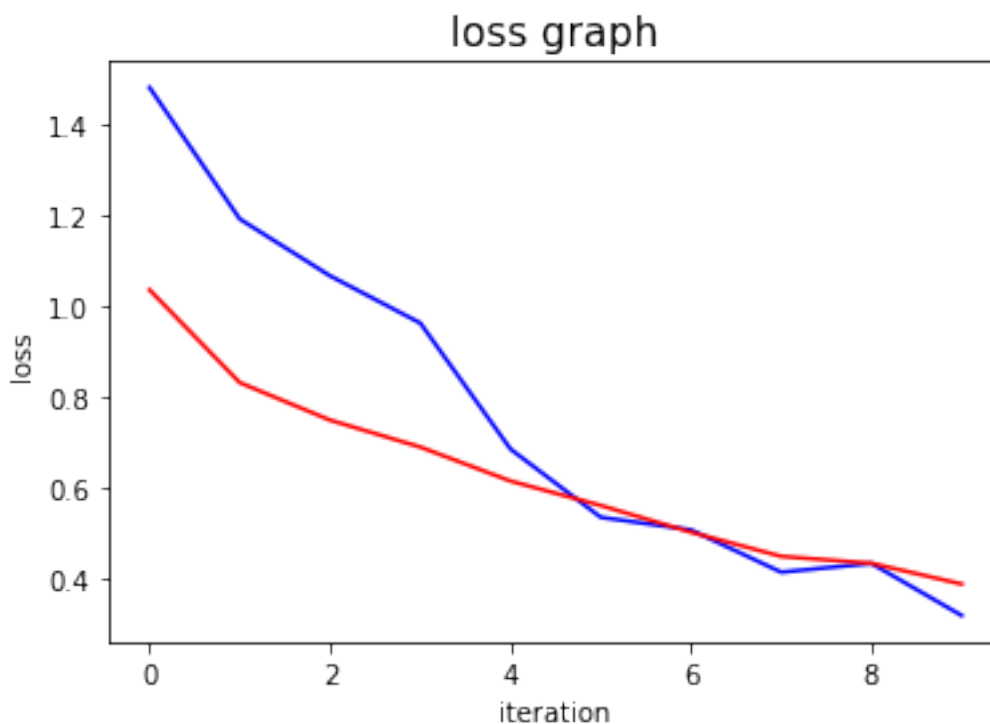
a = history.history

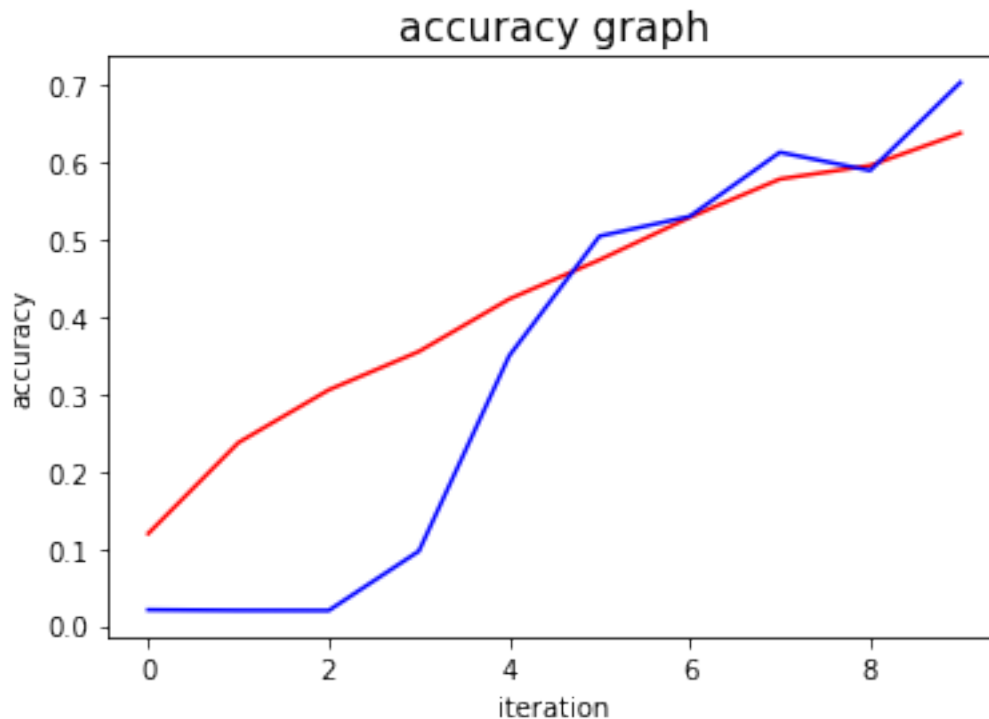
list_traindice = a['dice_coef']
list_testdice = a['val_dice_coef']

list_trainjaccard = a['iou']
list_testjaccard = a['val_iou']

list_trainloss = a['loss']
list_testloss = a['val_loss']
plt.figure(1)
plt.plot(list_testloss, 'b-')
plt.plot(list_trainloss, 'r-')
plt.xlabel('iteration')
plt.ylabel('loss')
plt.title('loss graph', fontsize = 15)
plt.figure(2)
plt.plot(list_traindice, 'r-')
plt.plot(list_testdice, 'b-')
plt.xlabel('iteration')
plt.ylabel('accuracy')
plt.title('accuracy graph', fontsize = 15)
plt.show()

```





## Evaluate the model

```
test_gen = train_generator(df_test, BATCH_SIZE,
                           dict(),
                           target_size=IMAGE_SIZE)
results = model.evaluate(test_gen, steps=len(df_test) / BATCH_SIZE)
print("Test IOU: ", results[1])
print("Test Dice Coefficient: ", results[2])
```

Found 590 validated image filenames.  
Found 590 validated image filenames.  
18/18 [=====] - 9s 487ms/step - loss: 0.3254  
- iou: 0.5431 - dice\_coef: 0.7001  
Test IOU: 0.5431138277053833  
Test Dice Coefficient: 0.7001265287399292

## Visualize the Result

```
for i in range(30):
    index=np.random.randint(1,len(df_test.index))
    img = cv2.imread(df_test['image_path'].iloc[index])
    img = cv2.resize(img ,IMAGE_SIZE)
    img = img / 255
    img = img[np.newaxis, :, :, :]
    pred=model.predict(img)

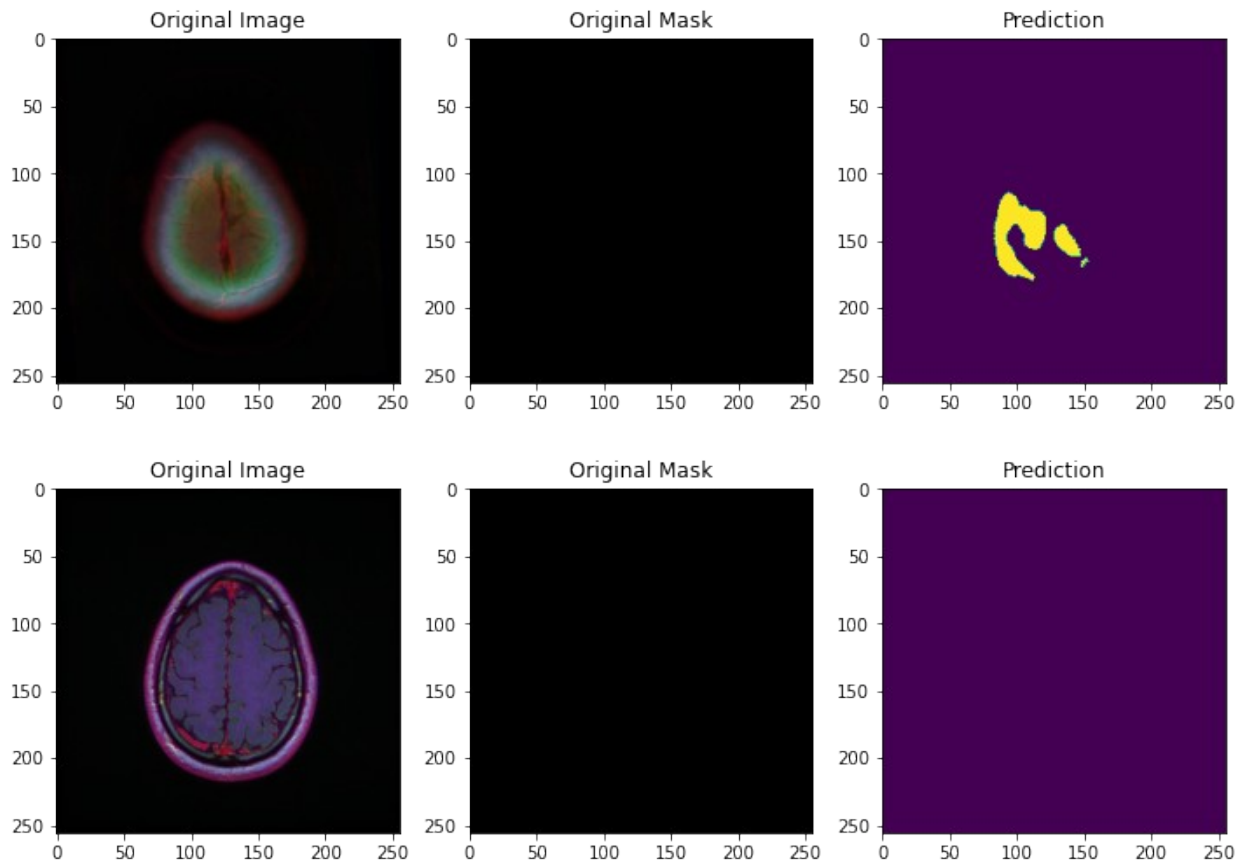
    plt.figure(figsize=(12,12))
```

```

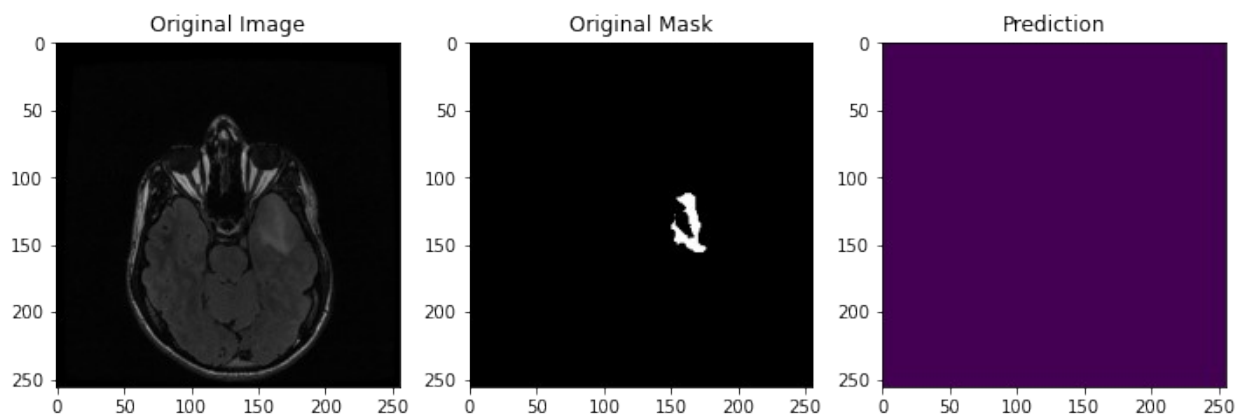
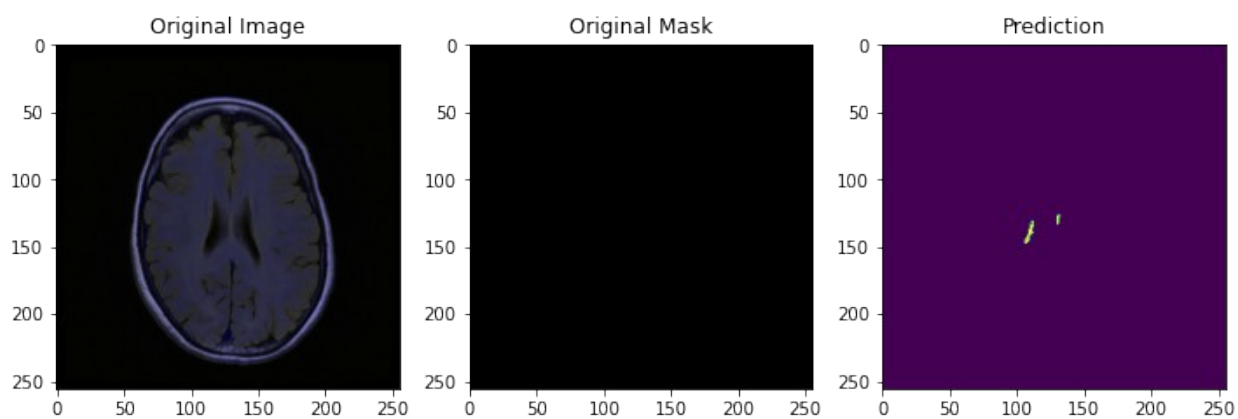
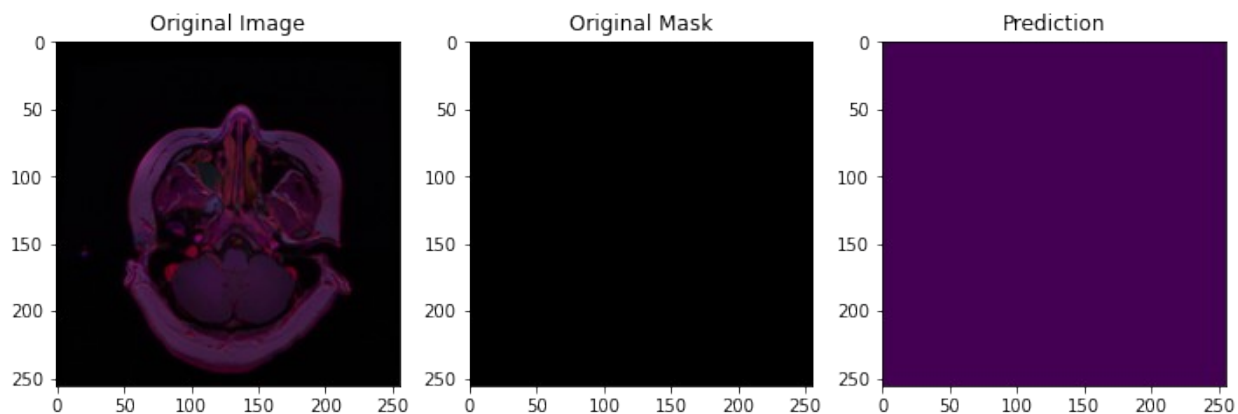
plt.subplot(1,3,1)
plt.imshow(np.squeeze(img))
plt.title('Original Image')
plt.subplot(1,3,2)

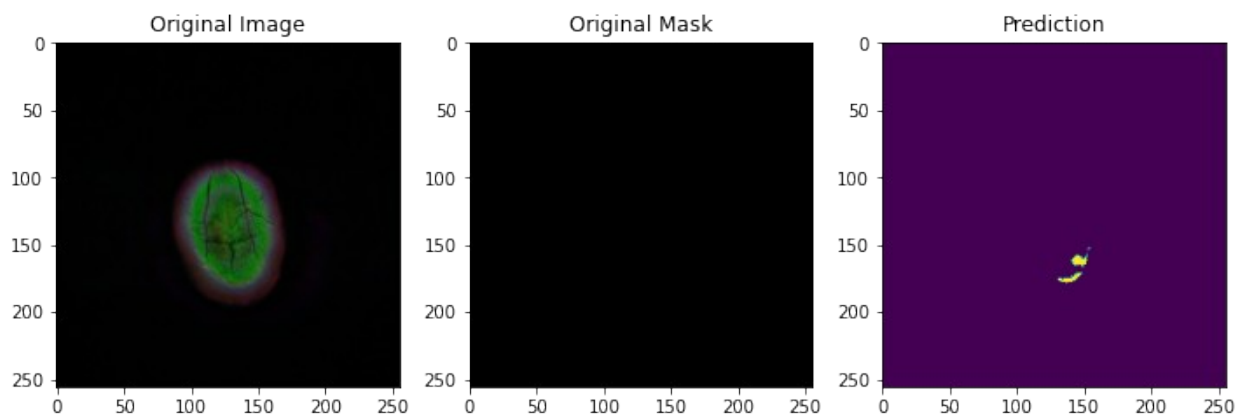
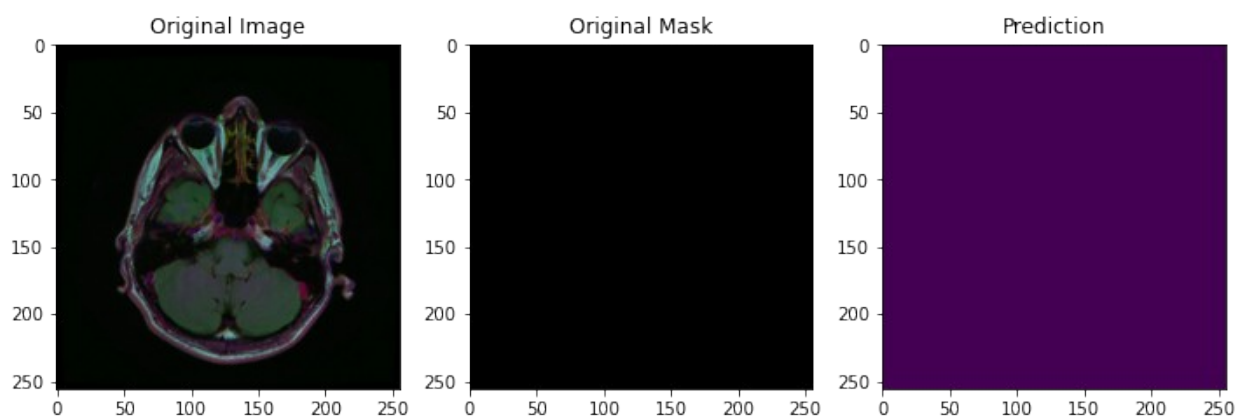
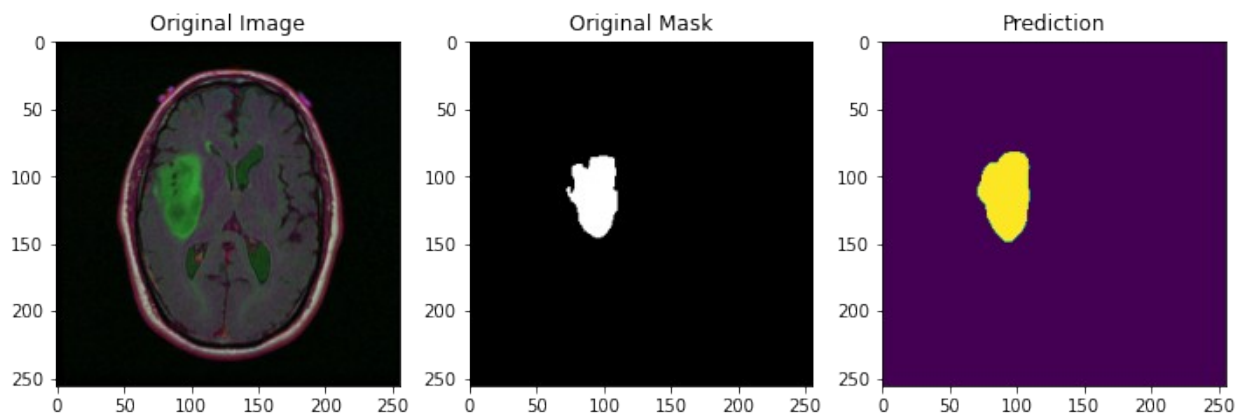
plt.imshow(np.squeeze(cv2.imread(df_test['mask_path'].iloc[index])))
plt.title('Original Mask')
plt.subplot(1,3,3)
plt.imshow(np.squeeze(pred) > .5)
plt.title('Prediction')
plt.show()

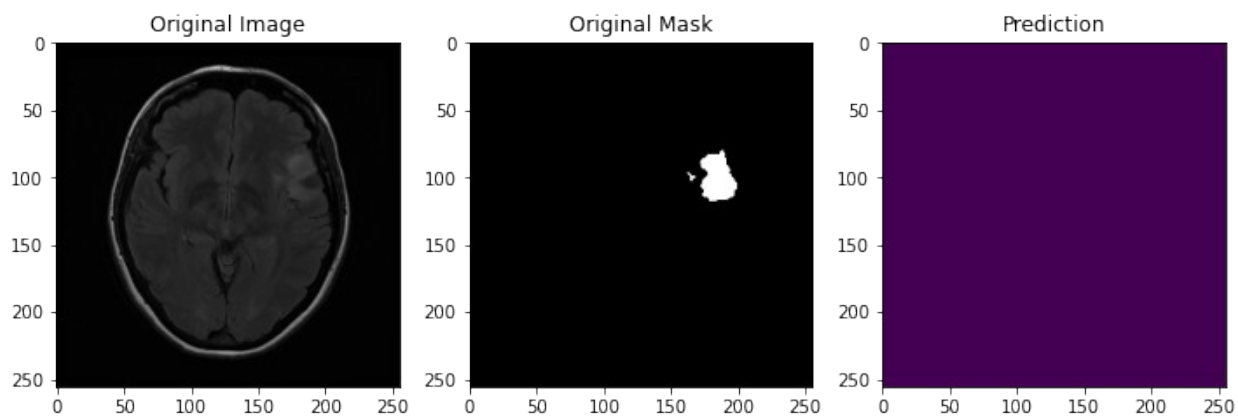
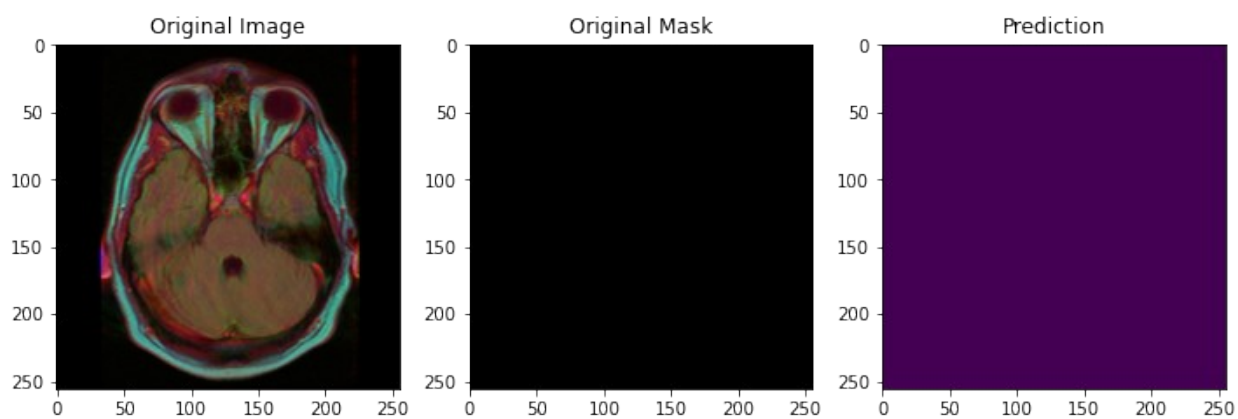
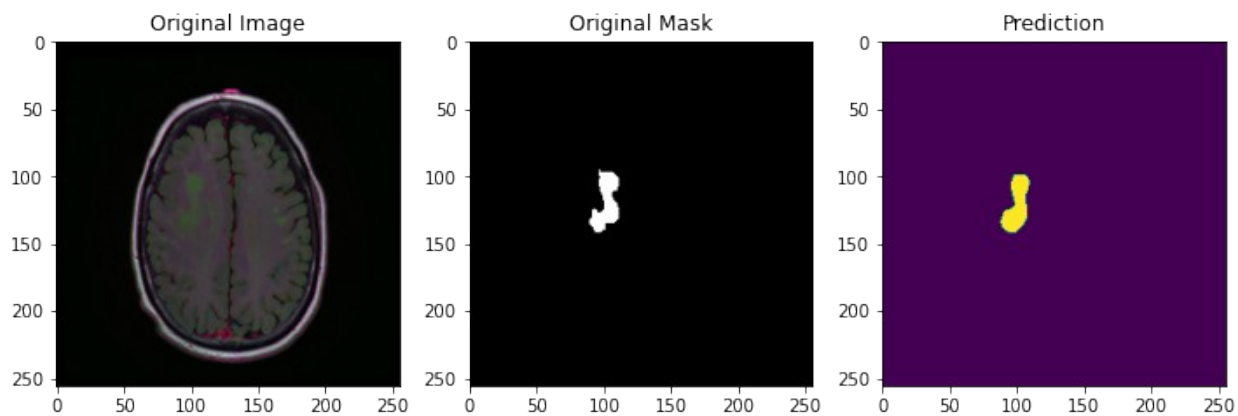
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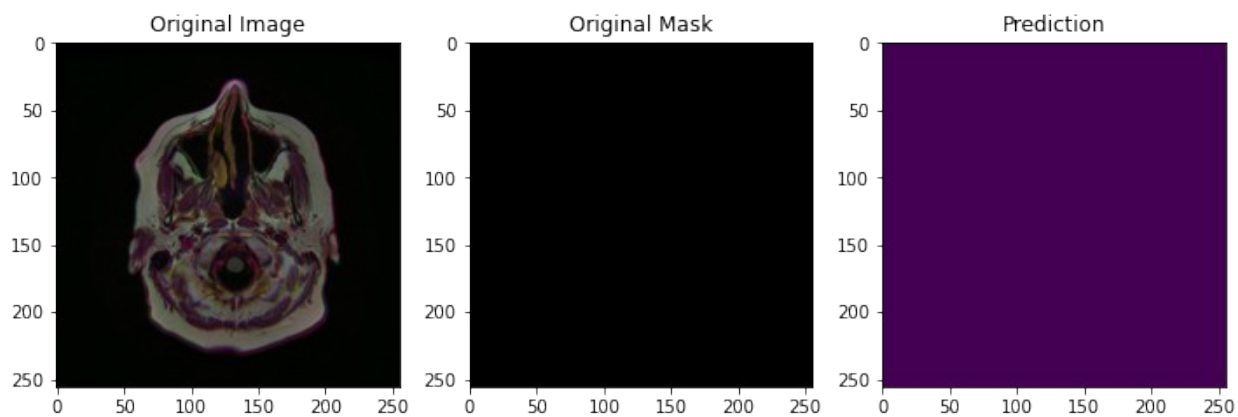
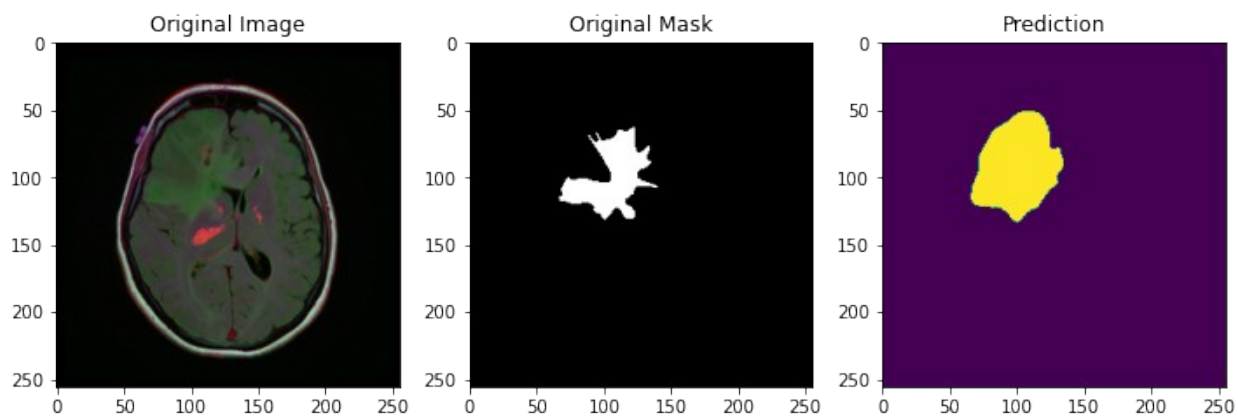
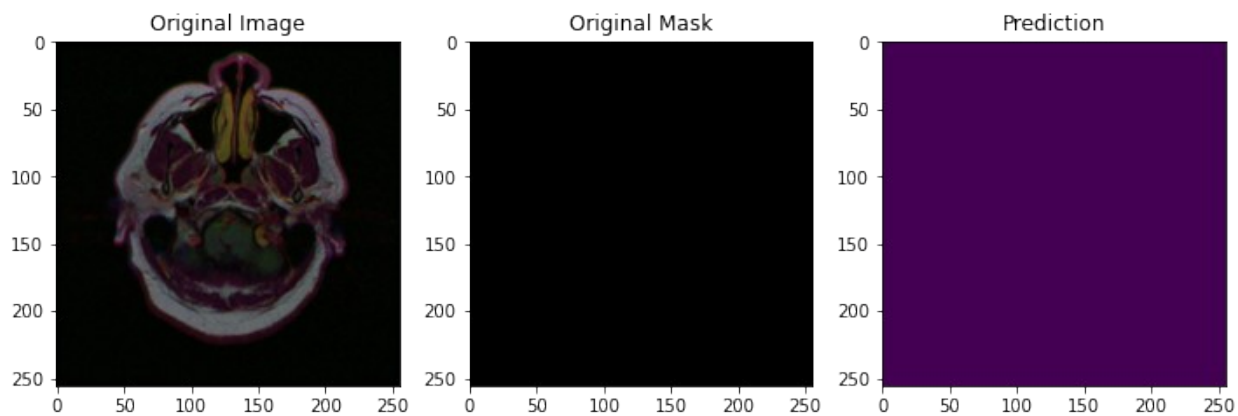


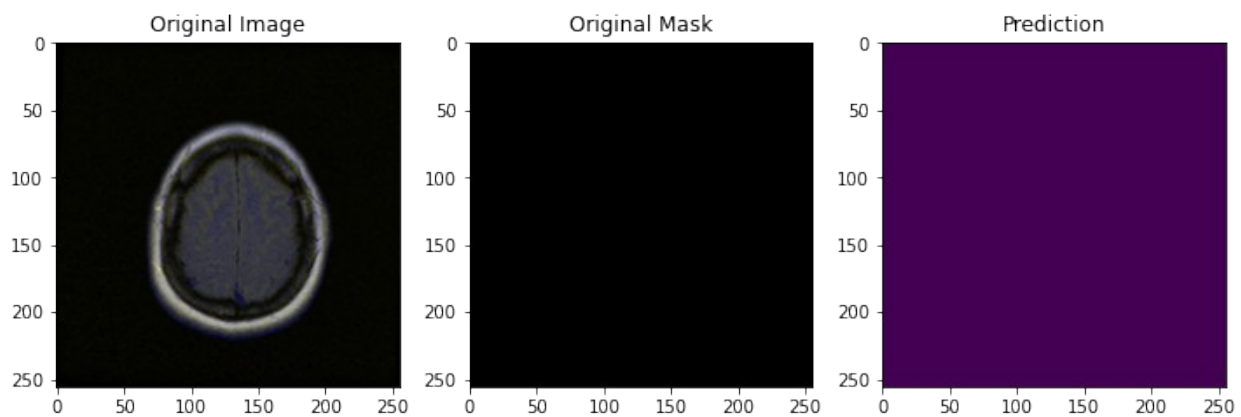
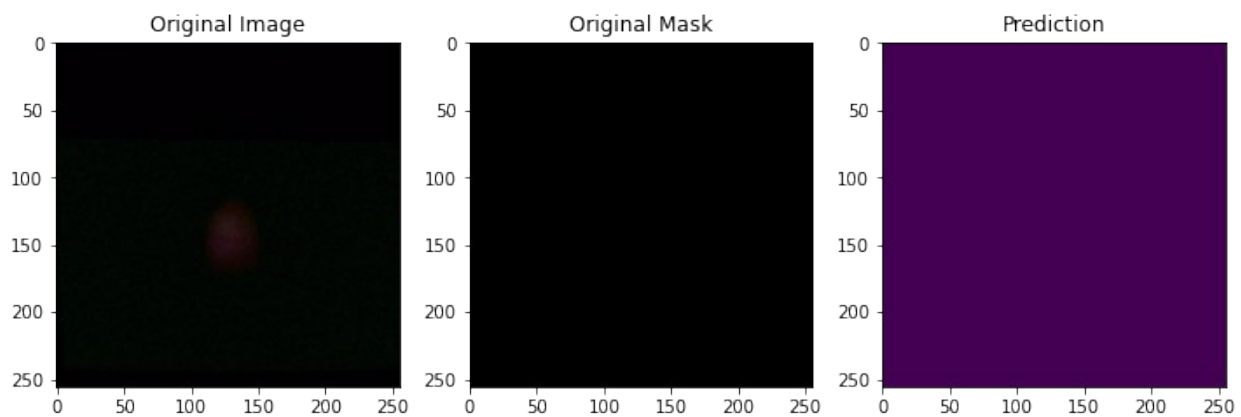
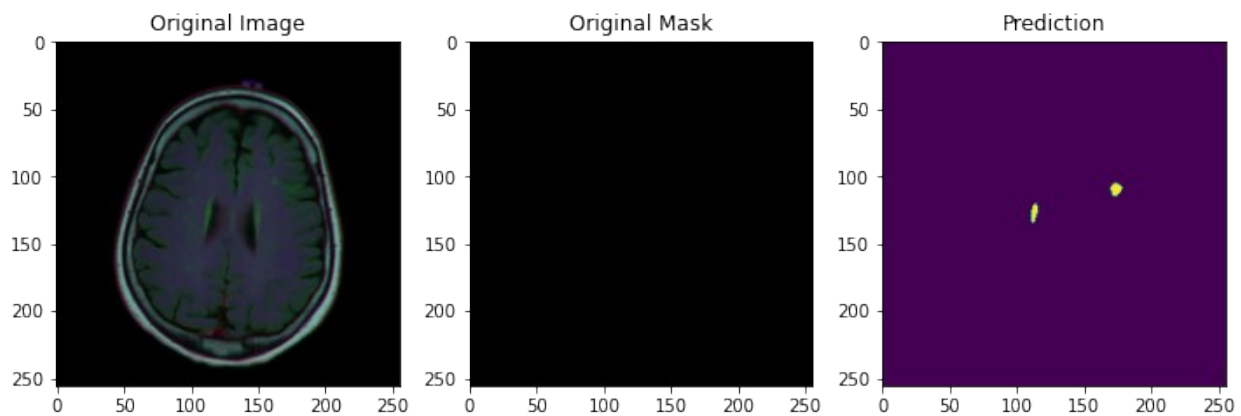


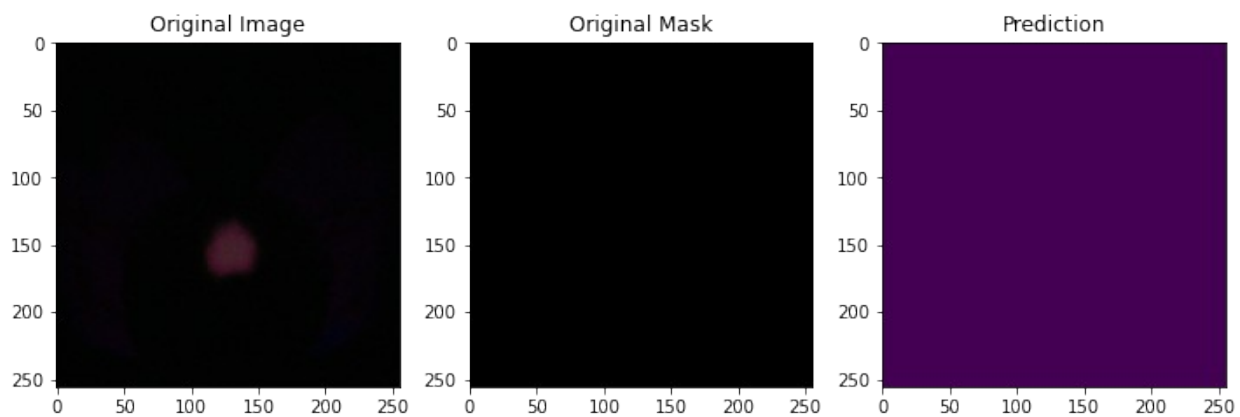
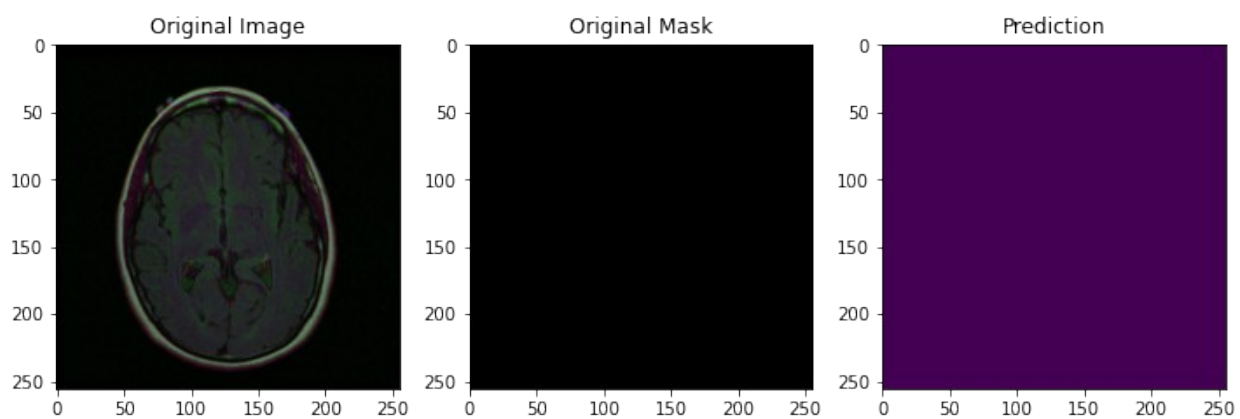
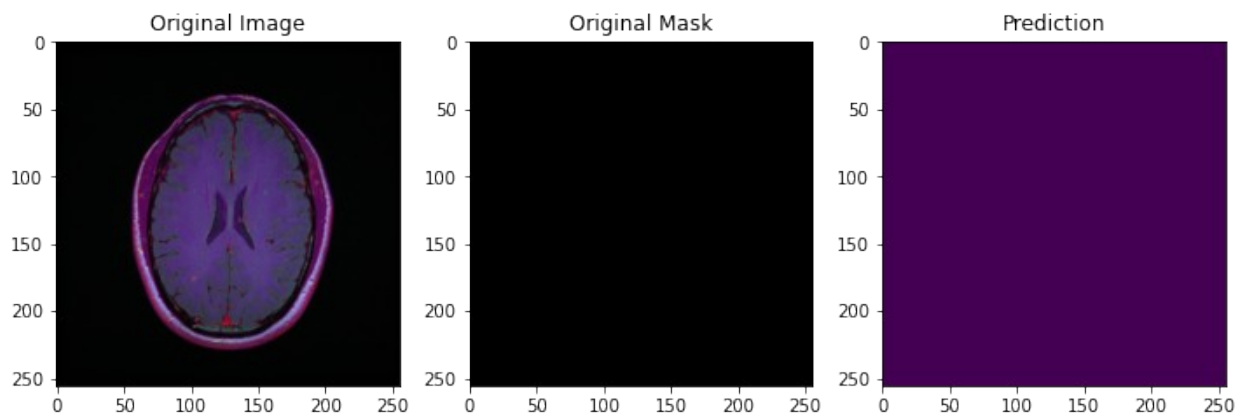


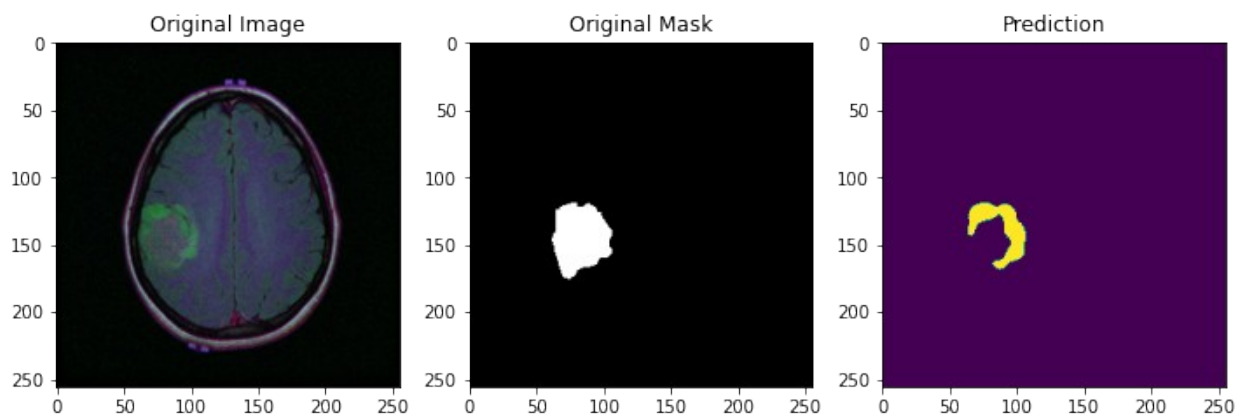
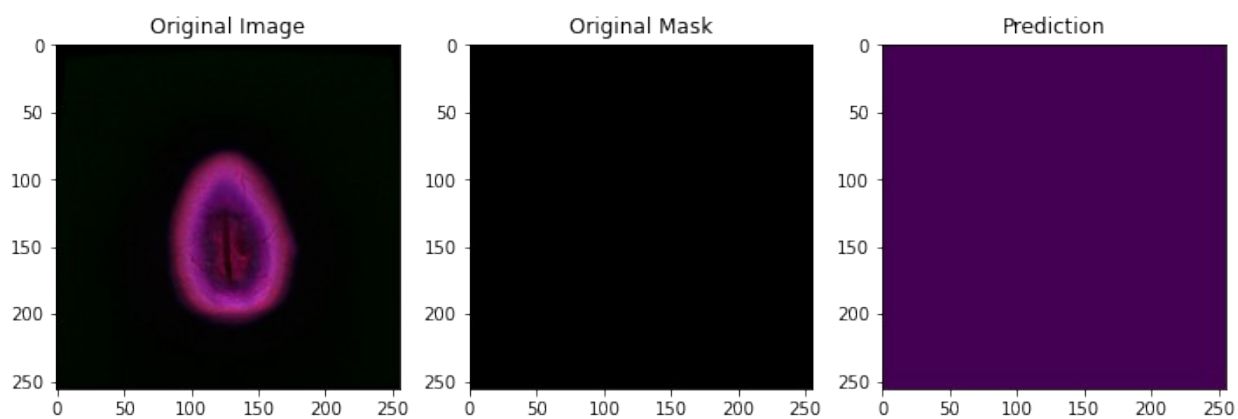
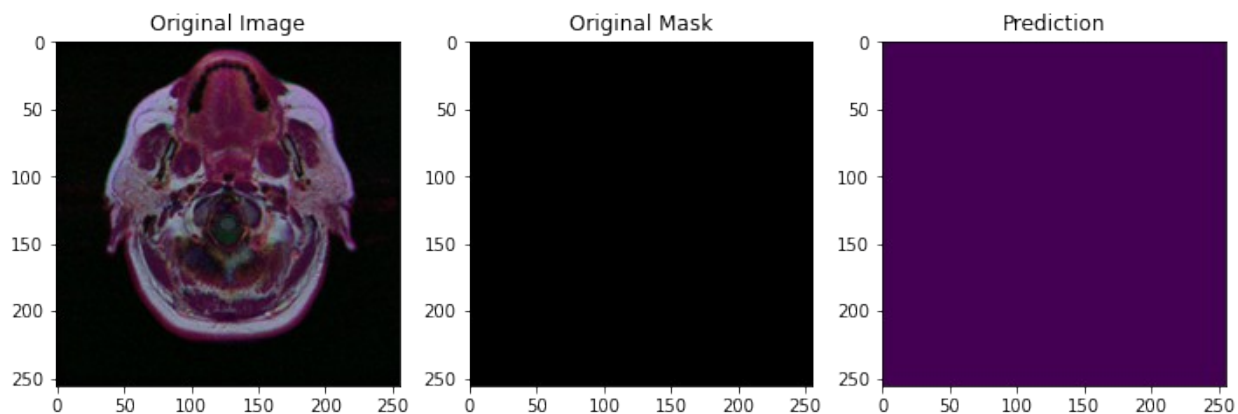


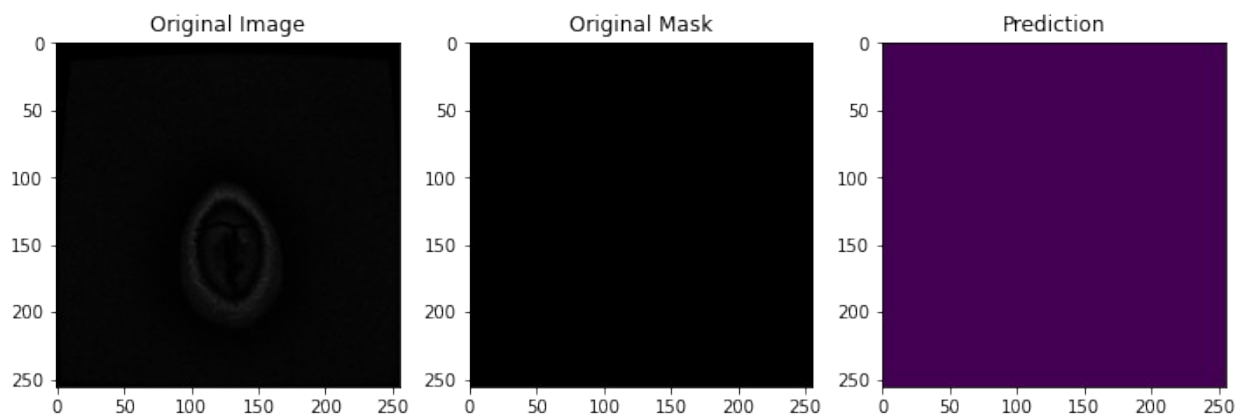
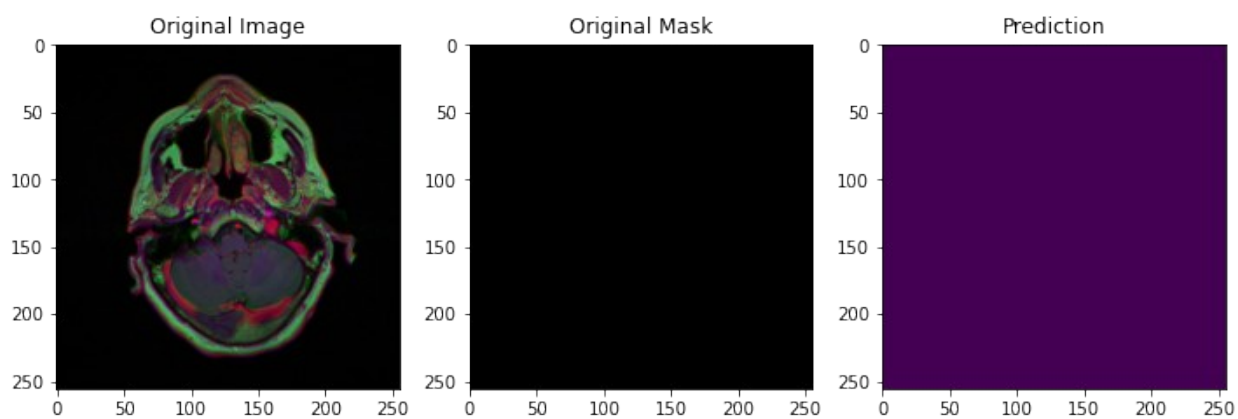
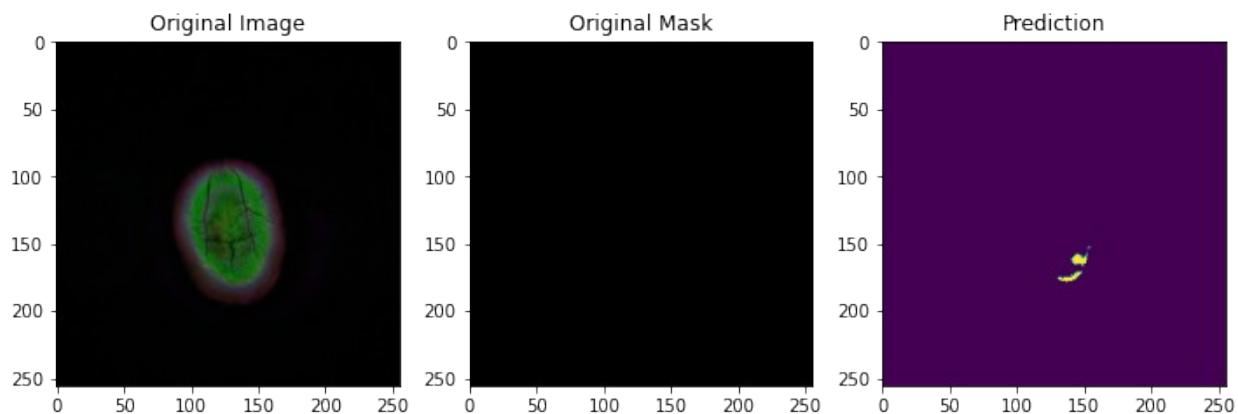




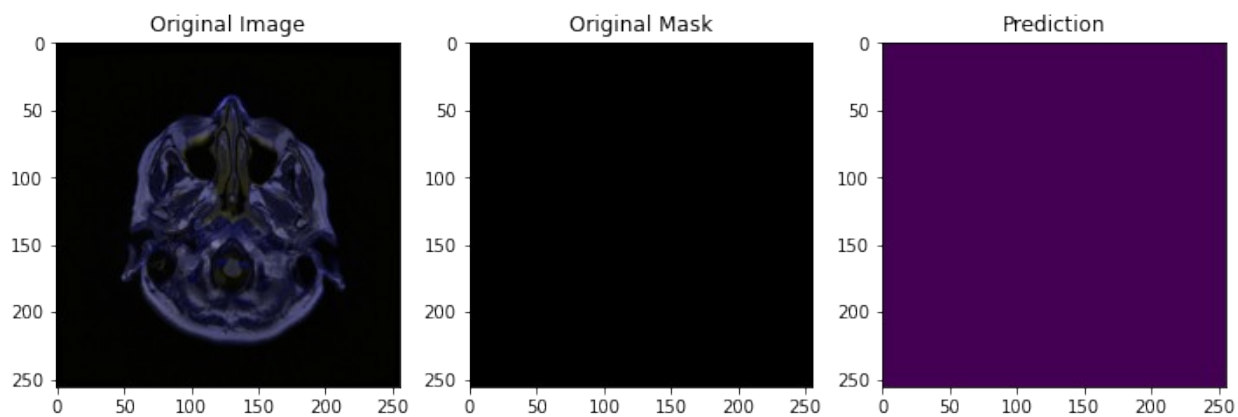
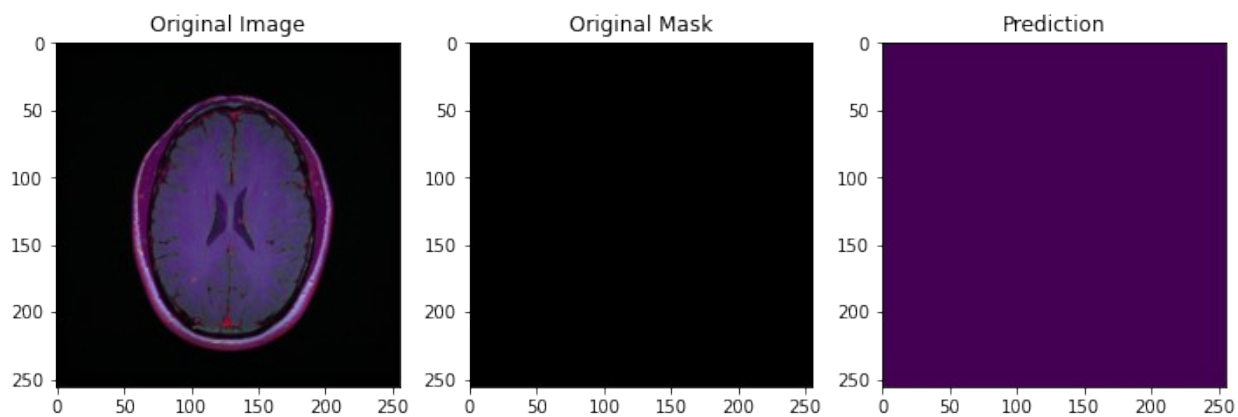
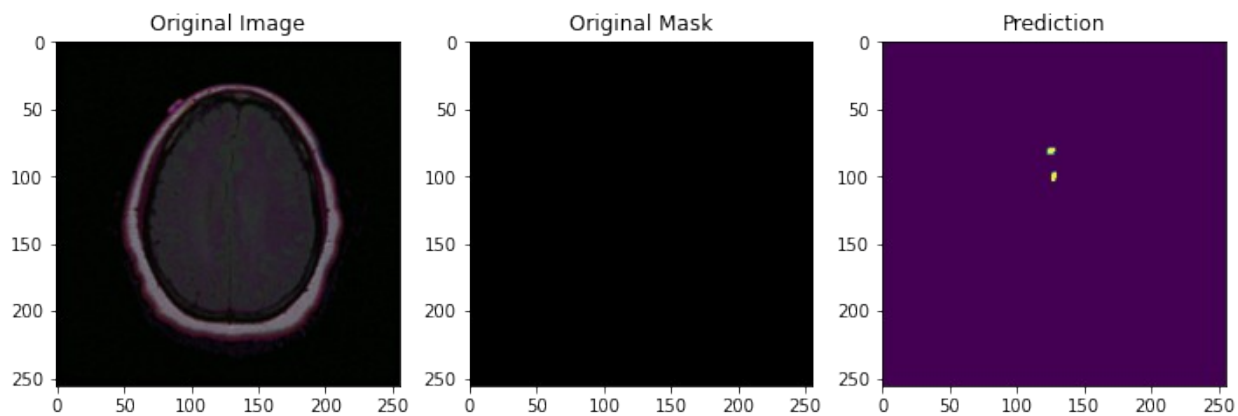


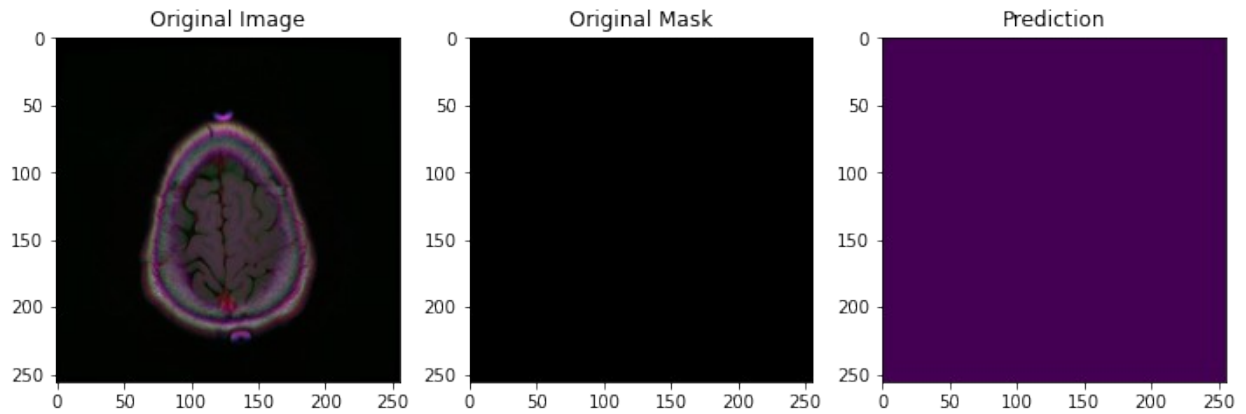












```
pred = np.where(pred > 0.5 , 1 , 0)
pred
```

```
array([[[[0],
         [0],
         [0],
         ...,
         [0],
         [0],
         [0]],
        [[0],
         [0],
         [0],
         ...,
         [0],
         [0],
         [0]],
        [[0],
         [0],
         [0],
         ...,
         [0],
         [0],
         [0]],
        ...,
        [[0],
         [0],
         [0],
         ...,
         [0],
         [0],
         [0]]],
       dtype=object])
```

```

[[0],
 [0],
 [0],
 ...,
 [0],
 [0],
 [0]],

[[0],
 [0],
 [0],
 ...,
 [0],
 [0],
 [0]]])

pred = np.squeeze(pred) > .5
pred
array([[False, False, False, ..., False, False, False],
       [False, False, False, ..., False, False, False],
       [False, False, False, ..., False, False, False],
       ...,
       [False, False, False, ..., False, False, False],
       [False, False, False, ..., False, False, False],
       [False, False, False, ..., False, False, False]])

cor , incor = 0 , 0
for i in range(len(df_test.index)):
    index=i
    img = cv2.imread(df_test['image_path'].iloc[index])
    img = cv2.resize(img ,IMAGE_SIZE)
    img = img / 255
    img = img[np.newaxis, :, :, :]
    pred = (np.squeeze(model.predict(img)) > 0.5)
    mask = np.squeeze(cv2.imread(df_test['mask_path'].iloc[index]))
    print(f"Prediction {i}: " , pred.any() == mask.any())
    if pred.any() == mask.any():
        cor += 1
    else:
        incor += 1

Prediction 0: True
Prediction 1: True
Prediction 2: False
Prediction 3: False
Prediction 4: True
Prediction 5: True
Prediction 6: True
Prediction 7: True

```

Prediction 8: True  
Prediction 9: True  
Prediction 10: True  
Prediction 11: False  
Prediction 12: True  
Prediction 13: True  
Prediction 14: True  
Prediction 15: True  
Prediction 16: True  
Prediction 17: False  
Prediction 18: True  
Prediction 19: True  
Prediction 20: True  
Prediction 21: True  
Prediction 22: True  
Prediction 23: True  
Prediction 24: True  
Prediction 25: True  
Prediction 26: True  
Prediction 27: True  
Prediction 28: True  
Prediction 29: True  
Prediction 30: True  
Prediction 31: True  
Prediction 32: True  
Prediction 33: True  
Prediction 34: True  
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Prediction 37: True  
Prediction 38: True  
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Prediction 41: True  
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Prediction 48: True  
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Prediction 50: False  
Prediction 51: False  
Prediction 52: True  
Prediction 53: False  
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Prediction 55: True  
Prediction 56: False

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Prediction 61: True  
Prediction 62: True  
Prediction 63: True  
Prediction 64: True  
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Prediction	251:	True
Prediction	252:	True



Prediction	253:	False
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Prediction	255:	True
Prediction	256:	True
Prediction	257:	True
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Prediction 548: True
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Prediction 583: True
Prediction 584: True
Prediction 585: True
Prediction 586: True
Prediction 587: True
Prediction 588: True
Prediction 589: True
```

```
import cv2
import numpy as np
import matplotlib.pyplot as plt
```

```
# Initialize lists to store prediction results and correctness
prediction_results = []
correctness = []
```

```

# Assuming df_test is your DataFrame containing image paths and mask
paths
for i in range(len(df_test.index)):
    index = i
    img = cv2.imread(df_test['image_path'].iloc[index])
    img = cv2.resize(img, IMAGE_SIZE)
    img = img / 255
    img = img[np.newaxis, :, :, :]
    pred = (np.squeeze(model.predict(img)) > 0.5)
    mask = np.squeeze(cv2.imread(df_test['mask_path'].iloc[index]))

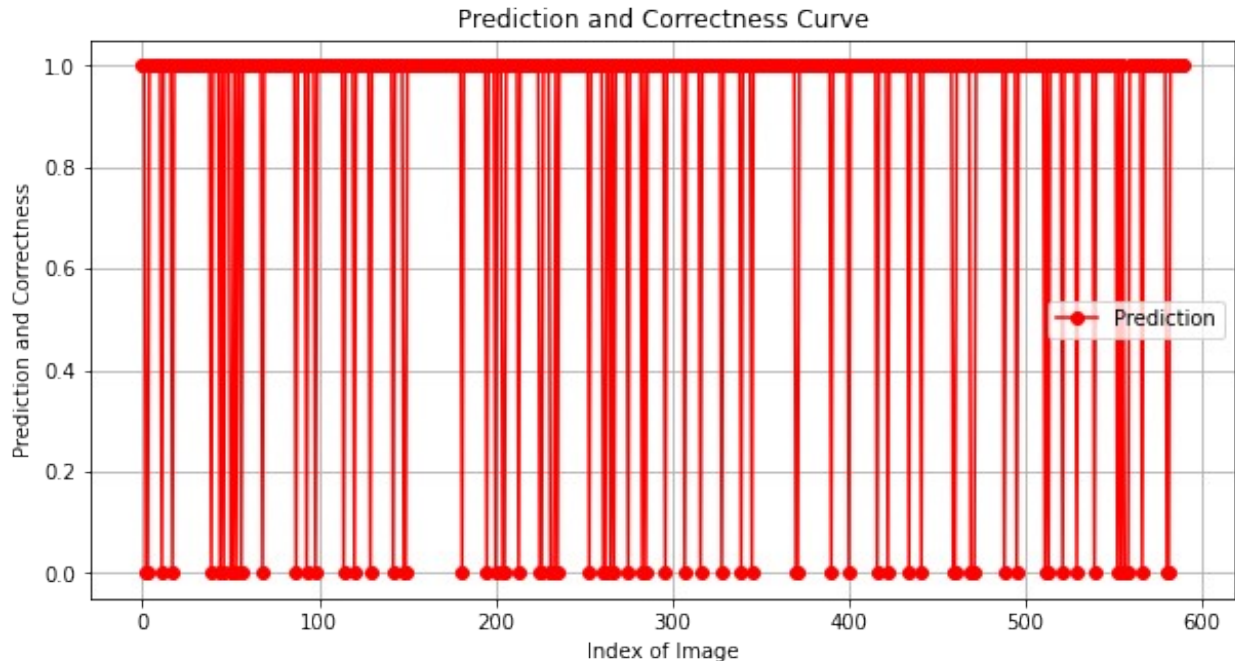
    # Check if prediction matches the ground truth mask
    prediction_correct = pred.any() == mask.any()

    # Append prediction result and correctness
    prediction_results.append(prediction_correct)
    correctness.append(1 if prediction_correct else 0)

# Plotting the curve
plt.figure(figsize=(10, 5))
plt.plot(prediction_results, label='Prediction', marker='o',
linestyle='-', color='red') # False predictions in red
#plt.plot(correctness, label='Correctness', marker='o', linestyle='-',
color='green') # Correct predictions in green
plt.xlabel('Index of Image')
plt.ylabel('Prediction and Correctness')
plt.title('Prediction and Correctness Curve')
plt.legend()
plt.grid(True)
plt.show()

```





```
import cv2
import numpy as np
import matplotlib.pyplot as plt

# Initialize lists to store prediction results and correctness
prediction_results = []
correctness = []

# Assuming df_test is your DataFrame containing image paths and mask paths
for i in range(len(df_test.index)):
    index = i
    img = cv2.imread(df_test['image_path'].iloc[index])
    img = cv2.resize(img, IMAGE_SIZE)
    img = img / 255
    img = img[np.newaxis, :, :, :]
    pred = (np.squeeze(model.predict(img)) > 0.5)
    mask = np.squeeze(cv2.imread(df_test['mask_path'].iloc[index]))

    # Check if prediction matches the ground truth mask
    prediction_correct = pred.any() == mask.any()

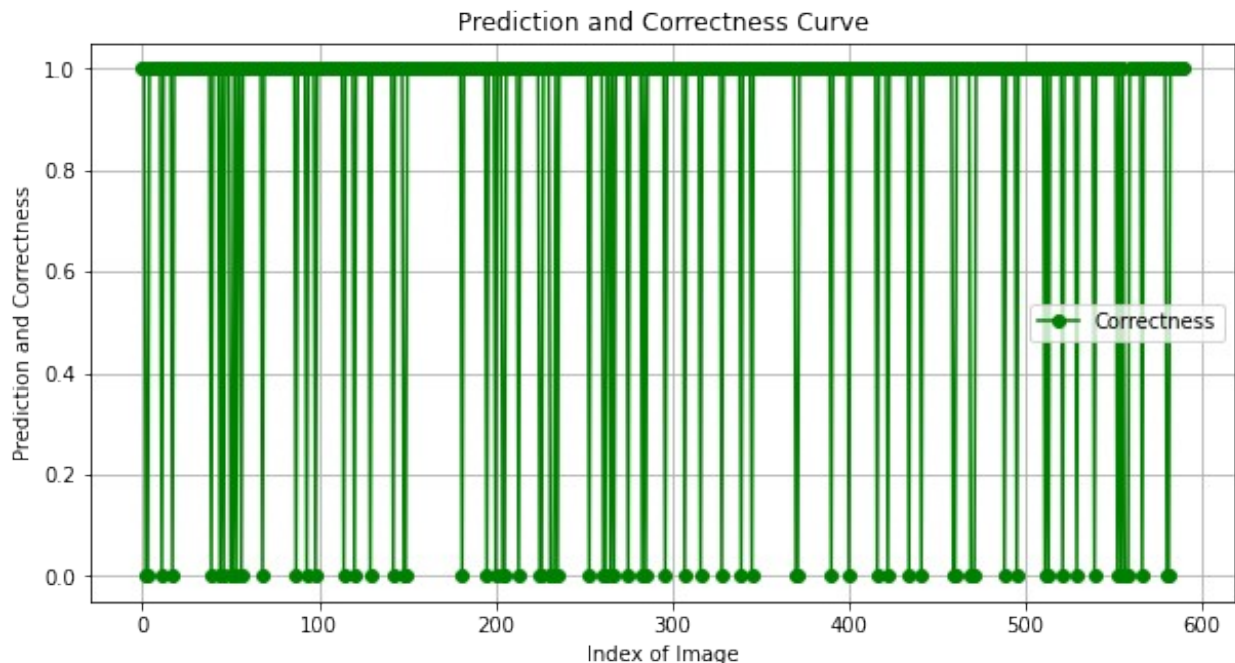
    # Append prediction result and correctness
    prediction_results.append(prediction_correct)
    correctness.append(1 if prediction_correct else 0)

# Plotting the curve
plt.figure(figsize=(10, 5))
# plt.plot(prediction_results, label='Prediction', marker='o',
```

```

linestyle='-', color='red') # False predictions in red
plt.plot(correctness, label='Correctness', marker='o', linestyle='-',
color='green') # Correct predictions in green
plt.xlabel('Index of Image')
plt.ylabel('Prediction and Correctness')
plt.title('Prediction and Correctness Curve')
plt.legend()
plt.grid(True)
plt.show()

```



```

import cv2
import numpy as np
import matplotlib.pyplot as plt

# Initialize lists to store prediction results and correctness
prediction_results = []
correctness = []

# Assuming df_test is your DataFrame containing image paths and mask
paths
for i in range(len(df_test.index)):
    index = i
    img = cv2.imread(df_test['image_path'].iloc[index])
    img = cv2.resize(img, IMAGE_SIZE)
    img = img / 255
    img = img[np.newaxis, :, :, :]
    pred = (np.squeeze(model.predict(img)) > 0.5)
    mask = np.squeeze(cv2.imread(df_test['mask_path'].iloc[index]))

```

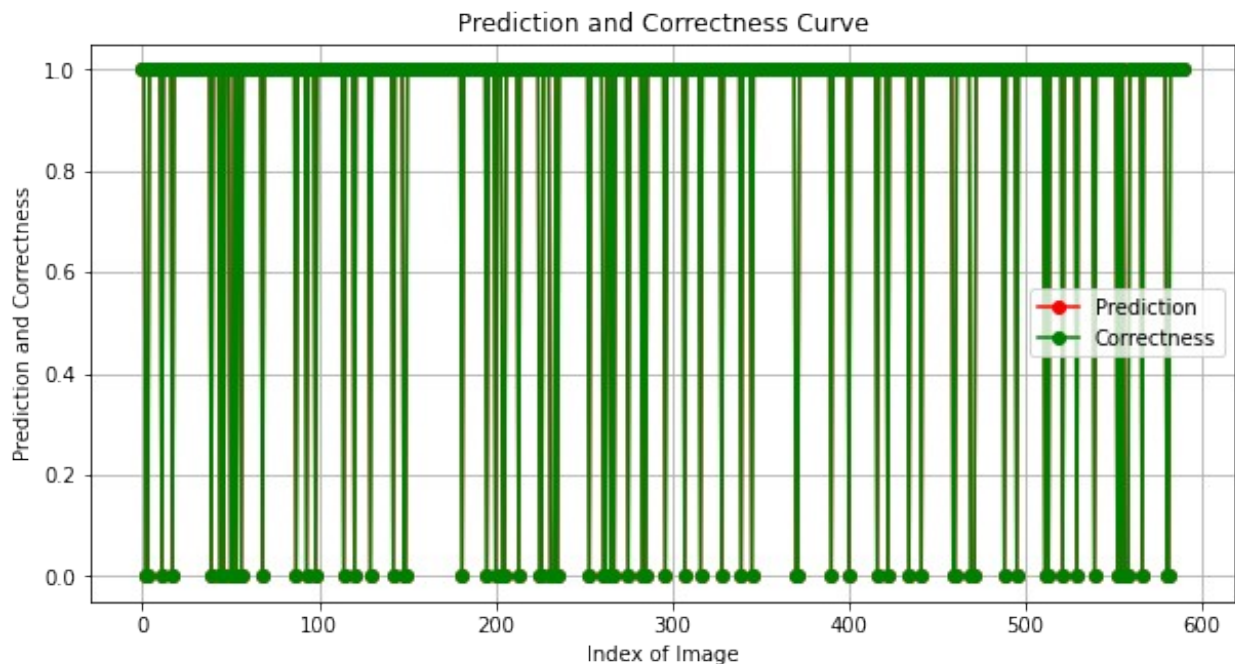
```

# Check if prediction matches the ground truth mask
prediction_correct = pred.any() == mask.any()

# Append prediction result and correctness
prediction_results.append(prediction_correct)
correctness.append(1 if prediction_correct else 0)

# Plotting the curve
plt.figure(figsize=(10, 5))
plt.plot(prediction_results, label='Prediction', marker='o',
linestyle='-', color='red') # False predictions in red
plt.plot(correctness, label='Correctness', marker='o', linestyle='-',
color='green') # Correct predictions in green
plt.xlabel('Index of Image')
plt.ylabel('Prediction and Correctness')
plt.title('Prediction and Correctness Curve')
plt.legend()
plt.grid(True)
plt.show()

```



```

print("Correct = " , cor , "Incorrect = " , incor )
print("Accuracy = " , (cor/(cor+incor)) * 100)

```

```

Correct = 515 Incorrect = 75
Accuracy = 87.28813559322035

```

```

import matplotlib.pyplot as plt

```

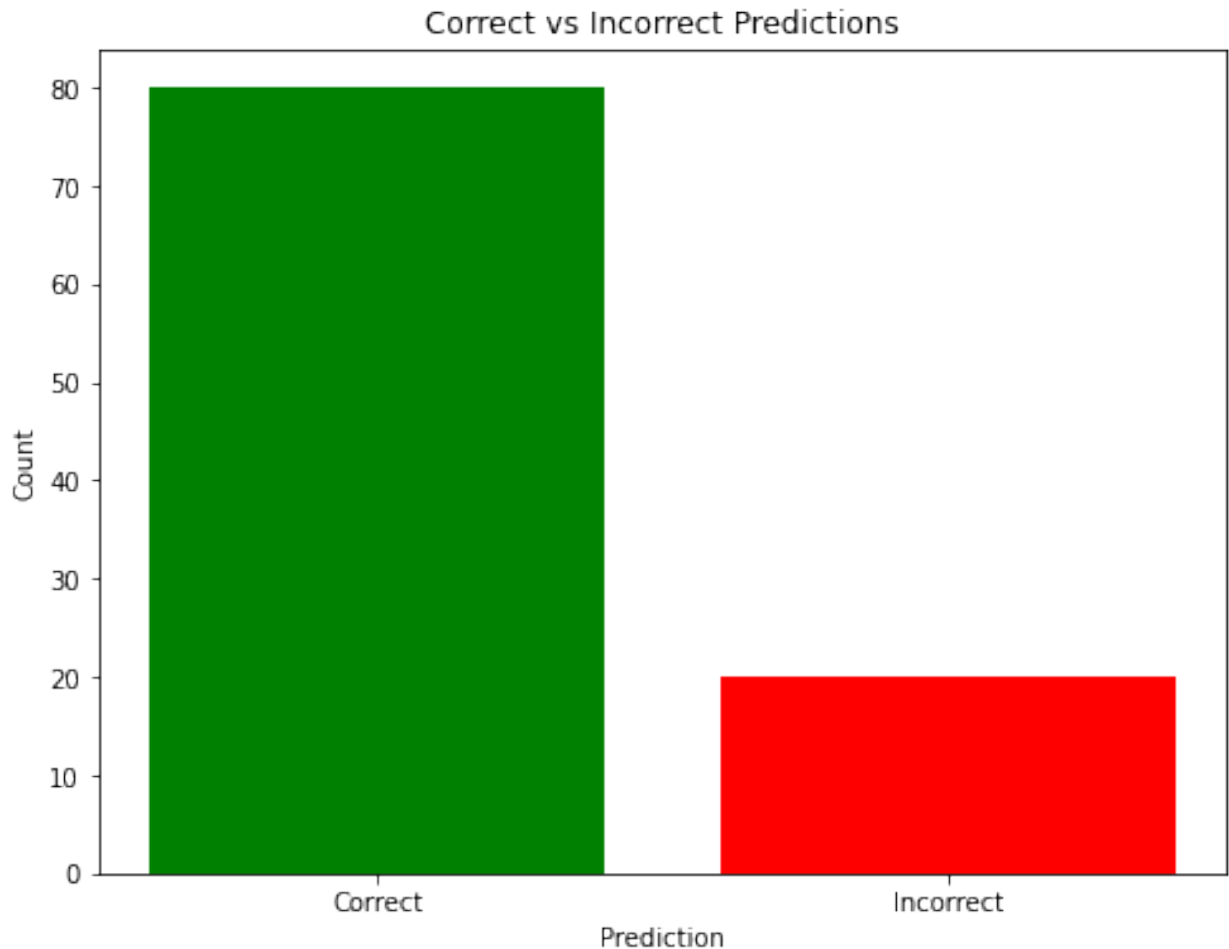
```
# Assuming cor and incor are the counts of correct and incorrect
predictions, respectively
cor = 80 # Example value
incor = 20 # Example value

# Calculate accuracy
accuracy = (cor / (cor + incor)) * 100

# Print correct and incorrect counts and accuracy
print("Correct =", cor, "Incorrect =", incor)
print("Accuracy =", accuracy)

# Plot a curve
# Here, you can plot any curve you want, for example, a bar plot
showing correct and incorrect counts
plt.figure(figsize=(8, 6))
plt.bar(["Correct", "Incorrect"], [cor, incor], color=['green',
'red'])
plt.xlabel('Prediction')
plt.ylabel('Count')
plt.title('Correct vs Incorrect Predictions')
plt.show()

Correct = 80 Incorrect = 20
Accuracy = 80.0
```

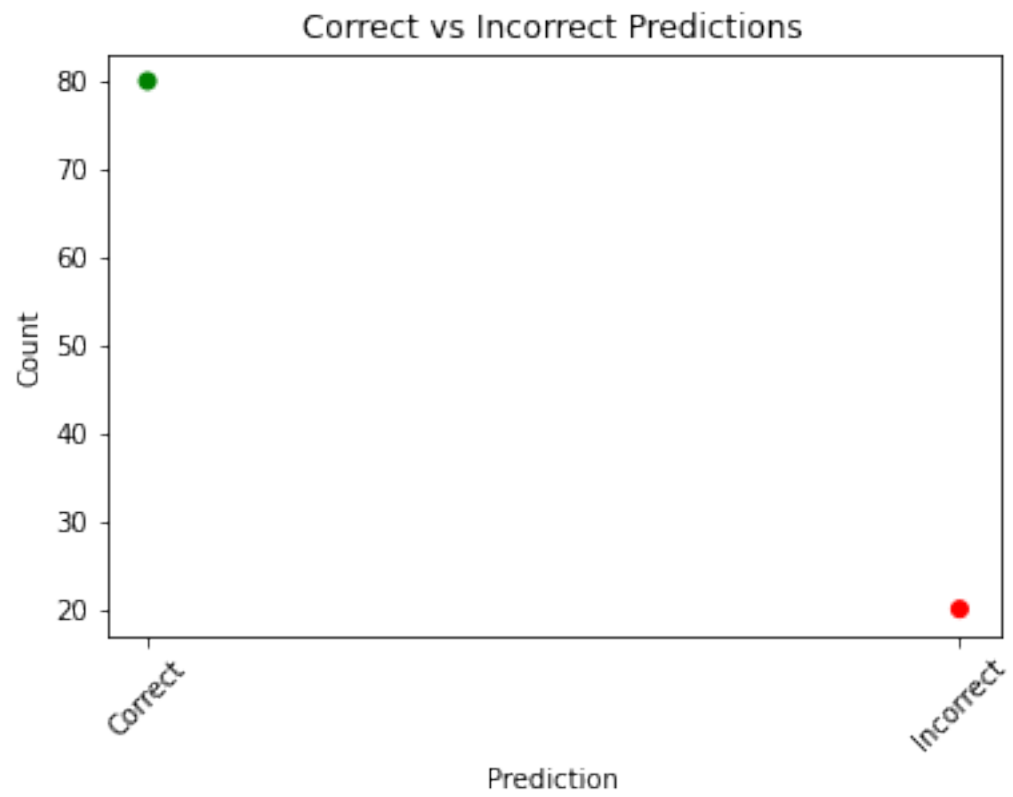


```
import matplotlib.pyplot as plt

# Assuming cor and incor are initialized earlier
cor, incor = 80, 20 # Example values

# Plotting the scatter plot
plt.figure(figsize=(6, 4))
plt.scatter(["Correct", "Incorrect"], [cor, incor], color=['green',
'red'])
plt.xlabel('Prediction')
plt.ylabel('Count')
plt.title('Correct vs Incorrect Predictions')
plt.xticks(rotation=45)
plt.show()

# Print correct and incorrect counts and accuracy
print("Correct =", cor, "Incorrect =", incor)
print("Accuracy =", (cor / (cor + incor)) * 100)
```



Correct = 80 Incorrect = 20  
Accuracy = 80.0