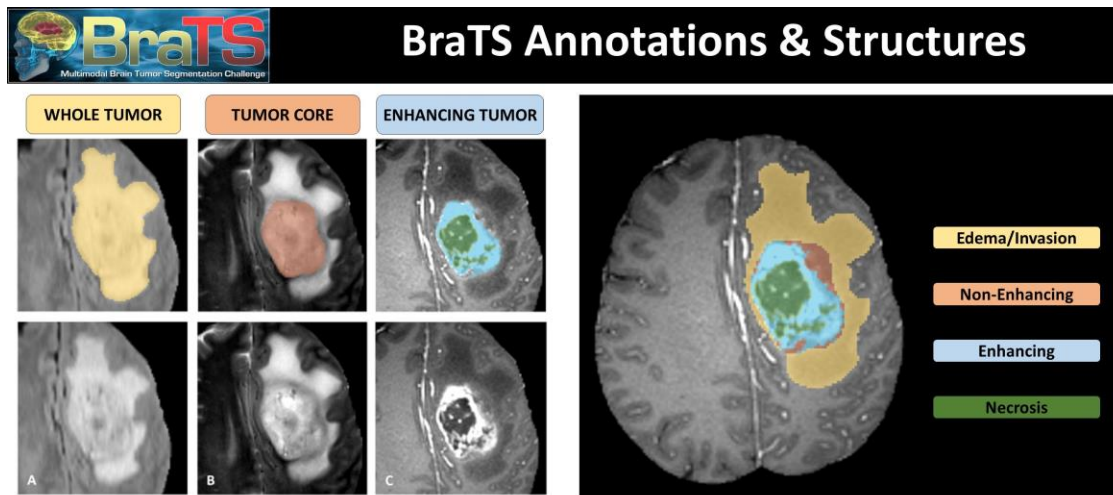


## Brain Tumor Segmentation BraTS 2020 Dataset



### Problem definition

#### Segmentation of gliomas in pre-operative MRI scans.

*Each pixel on image must be labeled:*

- Pixel is part of a tumor area (1 or 2 or 3) -> can be one of multiple classes / sub-regions
- Anything else -> pixel is not on a tumor region (0)

The sub-regions of tumor considered for evaluation are: 1) the "enhancing tumor" (ET), 2) the "tumor core" (TC), and 3) the "whole tumor" (WT). The provided segmentation labels have values of 1 for NCR & NET, 2 for ED, 4 for ET, and 0 for everything else.

```
import os
import cv2
import glob
import PIL
import shutil
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
from skimage import data
from skimage.util import montage
import skimage.transform as skTrans
from skimage.transform import rotate
from skimage.transform import resize
from PIL import Image, ImageOps
```

```

import nilearn as nl
import nibabel as nib
import nilearn.plotting as nlplt
!pip install git+https://github.com/miykael/gif_your_nifti
import gif_your_nifti.core as gif2nif

```

```

import keras
import keras.backend as K
from keras.callbacks import CSVLogger
import tensorflow as tf
from tensorflow.keras.utils import plot_model
from sklearn.preprocessing import MinMaxScaler
from sklearn.model_selection import train_test_split
from sklearn.metrics import classification_report
from tensorflow.keras.models import *
from tensorflow.keras.layers import *
from tensorflow.keras.optimizers import *
from tensorflow.keras.callbacks import ModelCheckpoint, ReduceLROnPlateau,
EarlyStopping, TensorBoard
from tensorflow.keras.layers.experimental import preprocessing

```

```

np.set_printoptions(precision=3, suppress=True)

```

Collecting git+https://github.com/miykael/gif\_your\_nifti

Cloning https://github.com/miykael/gif\_your\_nifti to /tmp/pip-req-build-aw647g58

Running command git clone -q https://github.com/miykael/gif\_your\_nifti /tmp/pip-req-build-aw647g58

Requirement already satisfied: numpy in /opt/conda/lib/python3.7/site-packages (from gif-your-nifti==0.2.2) (1.19.5)

Requirement already satisfied: nibabel in /opt/conda/lib/python3.7/site-packages (from gif-your-nifti==0.2.2) (3.2.1)

Requirement already satisfied: imageio<3 in /opt/conda/lib/python3.7/site-packages (from gif-your-nifti==0.2.2) (2.9.0)

Requirement already satisfied: matplotlib in /opt/conda/lib/python3.7/site-packages (from gif-your-nifti==0.2.2) (3.3.3)

Requirement already satisfied: scikit-image in /opt/conda/lib/python3.7/site-packages (from gif-your-nifti==0.2.2) (0.18.1)

Requirement already satisfied: pillow in /opt/conda/lib/python3.7/site-packages (from imageio<3->gif-your-nifti==0.2.2) (7.2.0)

Requirement already satisfied: cycler>=0.10 in /opt/conda/lib/python3.7/site-packages (from matplotlib->gif-your-nifti==0.2.2) (0.10.0)

Requirement already satisfied: kiwisolver>=1.0.1 in /opt/conda/lib/python3.7/site-packages (from matplotlib->gif-your-nifti==0.2.2) (1.3.1)

Requirement already satisfied: python-dateutil>=2.1 in /opt/conda/lib/python3.7/site-packages (from matplotlib->gif-your-

```

nifti==0.2.2) (2.8.1)
Requirement already satisfied: pyparsing!=2.0.4,!=2.1.2,!=2.1.6,>=2.0.3 in
/opt/conda/lib/python3.7/site-packages (from matplotlib->gif-your-
nifti==0.2.2) (2.4.7)
Requirement already satisfied: six in /opt/conda/lib/python3.7/site-packages
(from cyciler>=0.10->matplotlib->gif-your-nifti==0.2.2) (1.15.0)
Requirement already satisfied: packaging>=14.3 in
/opt/conda/lib/python3.7/site-packages (from nibabel->gif-your-nifti==0.2.2)
(20.8)
Requirement already satisfied: networkx>=2.0 in
/opt/conda/lib/python3.7/site-packages (from scikit-image->gif-your-
nifti==0.2.2) (2.5)
Requirement already satisfied: PyWavelets>=1.1.1 in
/opt/conda/lib/python3.7/site-packages (from scikit-image->gif-your-
nifti==0.2.2) (1.1.1)
Requirement already satisfied: scipy>=1.0.1 in /opt/conda/lib/python3.7/site-
packages (from scikit-image->gif-your-nifti==0.2.2) (1.5.4)
Requirement already satisfied: tifffile>=2019.7.26 in
/opt/conda/lib/python3.7/site-packages (from scikit-image->gif-your-
nifti==0.2.2) (2021.2.1)
Requirement already satisfied: decorator>=4.3.0 in
/opt/conda/lib/python3.7/site-packages (from networkx>=2.0->scikit-image-
>gif-your-nifti==0.2.2) (4.4.2)
Building wheels for collected packages: gif-your-nifti
  Building wheel for gif-your-nifti (setup.py) ... e=gif_your_nifti-0.2.2-
py3-none-any.whl size=6634
sha256=c43944de372984e36d22f1d46e123143283e2888842fdd6a58782c71ca4b75aa
  Stored in directory: /tmp/pip-ephem-wheel-cache-
2mz6u469/wheels/4a/8c/d1/b228c3b67231f7459e8f70d73f4dadaf65cd90692d41f43e88
Successfully built gif-your-nifti
Installing collected packages: gif-your-nifti
Successfully installed gif-your-nifti-0.2.2
WARNING: You are using pip version 21.0.1; however, version 24.0 is
available.
You should consider upgrading via the '/opt/conda/bin/python3.7 -m pip
install --upgrade pip' command.

```

```

SEGMENT_CLASSES = {
    0 : 'NOT tumor',
    1 : 'NECROTIC/CORE',
    2 : 'EDEMA',
    3 : 'ENHANCING'
}

```

```

VOLUME_SLICES = 100
VOLUME_START_AT = 22

```

## Image data descriptions

All BraTS multimodal scans are available as NIfTI files (.nii.gz) -> commonly used medical imaging format to store brain imaging data obtained using MRI and describe different MRI settings

1. **T1:** T1-weighted, native image, sagittal or axial 2D acquisitions, with 1–6 mm slice thickness.
1. **T1c:** T1-weighted, contrast-enhanced (Gadolinium) image, with 3D acquisition and 1 mm isotropic voxel size for most patients.
2. **T2:** T2-weighted image, axial 2D acquisition, with 2–6 mm slice thickness.
3. **FLAIR:** T2-weighted FLAIR image, axial, coronal, or sagittal 2D acquisitions, 2–6 mm slice thickness.

Data were acquired with different clinical protocols and various scanners from multiple (n=19) institutions.

All the imaging datasets have been segmented manually, by one to four raters, following the same annotation protocol, and their annotations were approved by experienced neuro-radiologists. Annotations comprise the GD-enhancing tumor (ET — label 4), the peritumoral edema (ED — label 2), and the necrotic and non-enhancing tumor core (NCR/NET — label 1), as described both in the BraTS 2012-2013 TMI paper and in the latest BraTS summarizing paper. The provided data are distributed after their pre-processing, i.e., co-registered to the same anatomical template, interpolated to the same resolution (1 mm<sup>3</sup>) and skull-stripped.

```
TRAIN_DATASET_PATH = '../input/brats20-dataset-training-validation/BraTS2020_TrainingData/MICCAI_BraTS2020_TrainingData/'
VALIDATION_DATASET_PATH = '../input/brats20-dataset-training-validation/BraTS2020_ValidationData/MICCAI_BraTS2020_ValidationData'
```

```
test_image_flair=nib.load(TRAIN_DATASET_PATH +
'BraTS20_Training_001/BraTS20_Training_001_flair.nii').get_fdata()
test_image_t1=nib.load(TRAIN_DATASET_PATH +
'BraTS20_Training_001/BraTS20_Training_001_t1.nii').get_fdata()
test_image_t1ce=nib.load(TRAIN_DATASET_PATH +
'BraTS20_Training_001/BraTS20_Training_001_t1ce.nii').get_fdata()
test_image_t2=nib.load(TRAIN_DATASET_PATH +
'BraTS20_Training_001/BraTS20_Training_001_t2.nii').get_fdata()
test_mask=nib.load(TRAIN_DATASET_PATH +
'BraTS20_Training_001/BraTS20_Training_001_seg.nii').get_fdata()
```

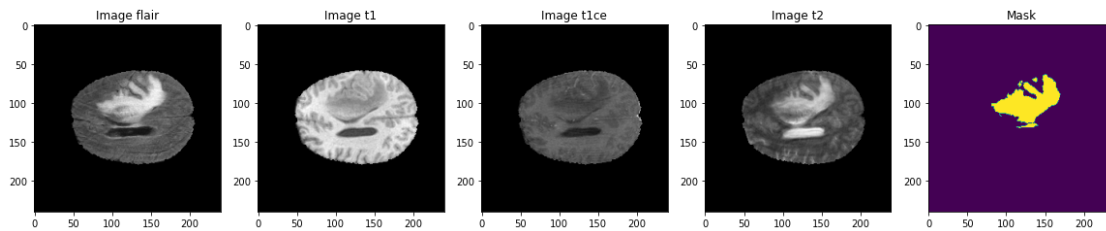
```
fig, (ax1, ax2, ax3, ax4, ax5) = plt.subplots(1,5, figsize = (20, 10))
slice_w = 25
ax1.imshow(test_image_flair[:, :, test_image_flair.shape[0]//2-slice_w], cmap =
'gray')
ax1.set_title('Image flair')
```

```

ax2.imshow(test_image_t1[:, :, test_image_t1.shape[0]//2-slice_w], cmap =
'gray')
ax2.set_title('Image t1')
ax3.imshow(test_image_t1ce[:, :, test_image_t1ce.shape[0]//2-slice_w], cmap =
'gray')
ax3.set_title('Image t1ce')
ax4.imshow(test_image_t2[:, :, test_image_t2.shape[0]//2-slice_w], cmap =
'gray')
ax4.set_title('Image t2')
ax5.imshow(test_mask[:, :, test_mask.shape[0]//2-slice_w])
ax5.set_title('Mask')

Text(0.5, 1.0, 'Mask')

```

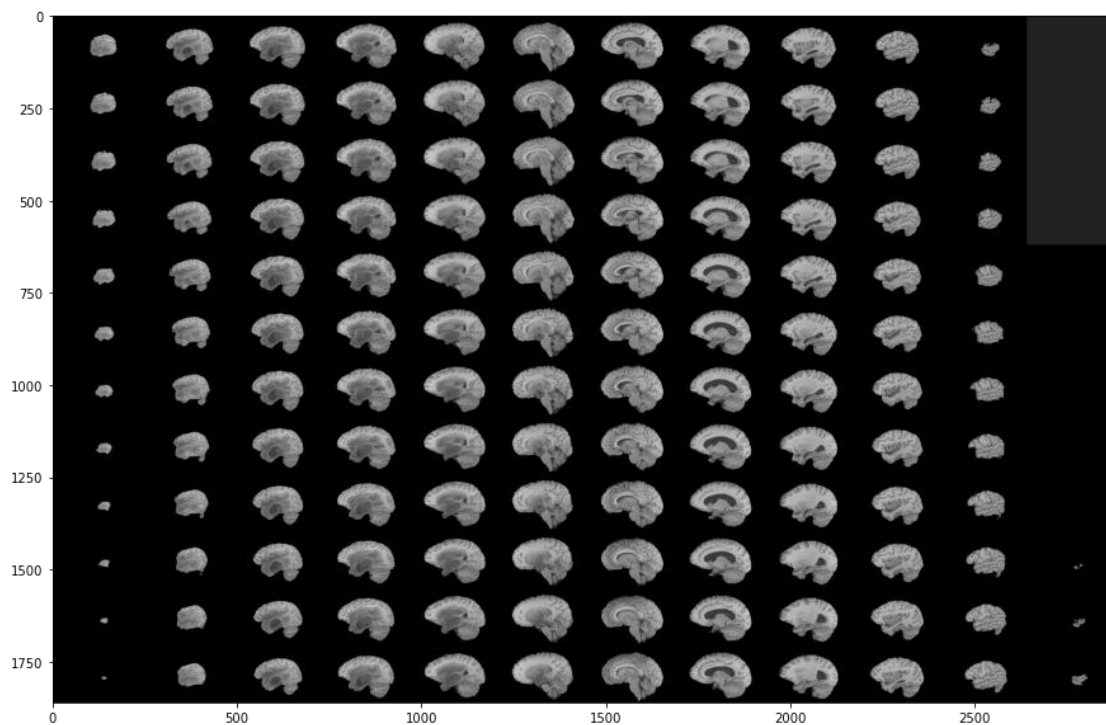


```

fig, ax1 = plt.subplots(1, 1, figsize = (15,15))
ax1.imshow(rotate(montage(test_image_t1[50:-50,:,:]), 90, resize=True), cmap
='gray')

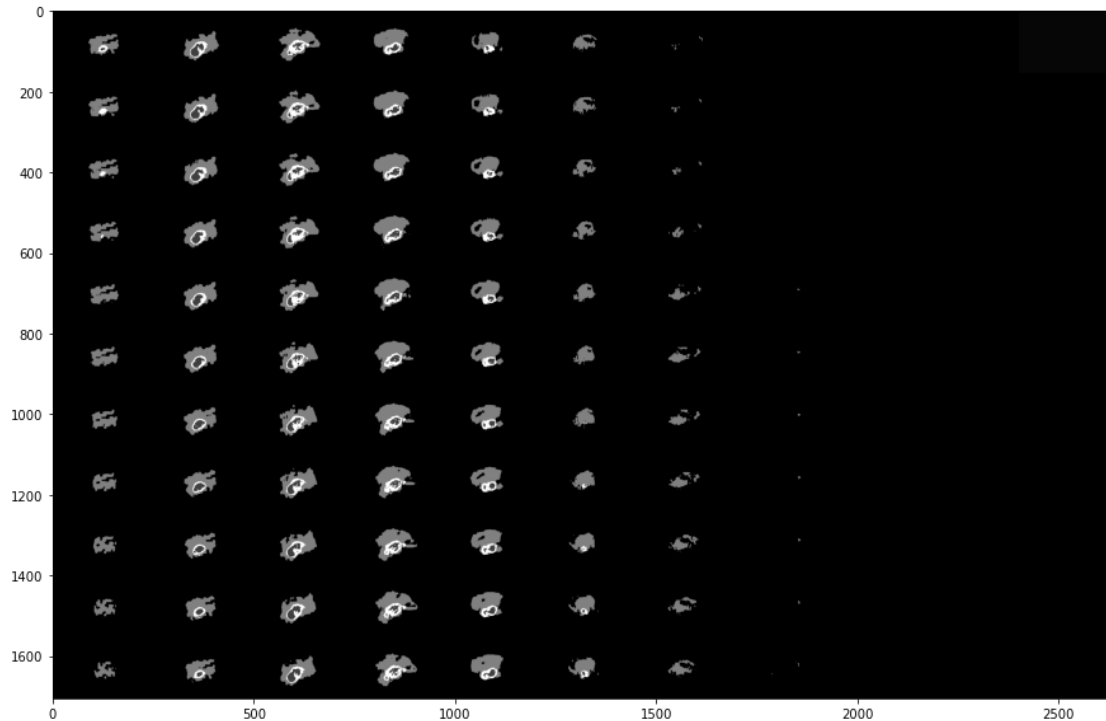
```

<matplotlib.image.AxesImage at 0x7a7b9a3732d0>



```
fig, ax1 = plt.subplots(1, 1, figsize = (15,15))
ax1.imshow(rotate(montage(test_mask[60:-60,:,:]), 90, resize=True), cmap
='gray')
```

<matplotlib.image.AxesImage at 0x7a7bdb744250>



```
shutil.copy2(TRAIN_DATASET_PATH +
'BraTS20_Training_001/BraTS20_Training_001_flair.nii',
'./test_gif_BraTS20_Training_001_flair.nii')
gif2nif.write_gif_normal('./test_gif_BraTS20_Training_001_flair.nii')
```

```
niimg = nl.image.load_img(TRAIN_DATASET_PATH +
'BraTS20_Training_001/BraTS20_Training_001_flair.nii')
nimask = nl.image.load_img(TRAIN_DATASET_PATH +
'BraTS20_Training_001/BraTS20_Training_001_seg.nii')
```

```
fig, axes = plt.subplots(nrows=4, figsize=(30, 40))
```

```
nlplt.plot_anat(niimg,
                title='BraTS20_Training_001_flair.nii plot_anat',
                axes=axes[0])
```

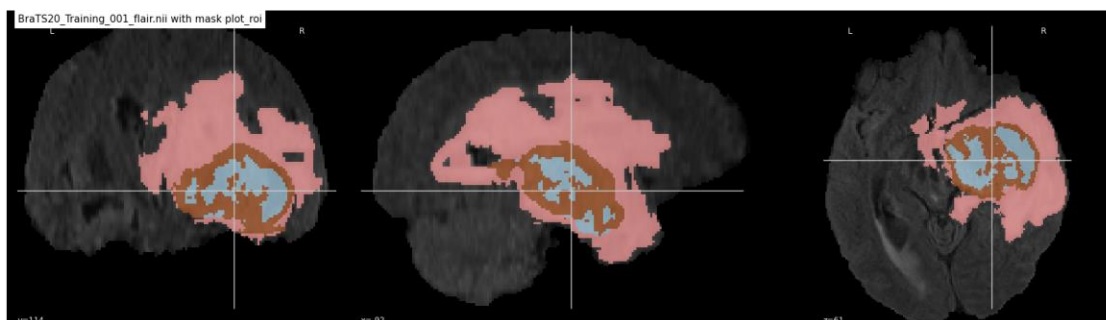
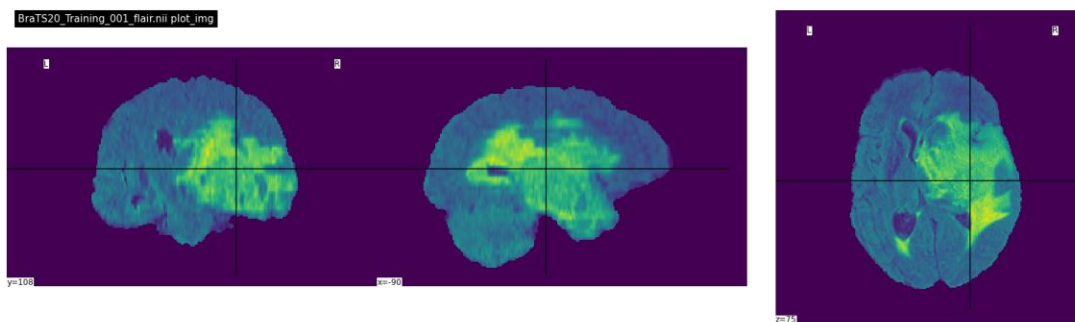
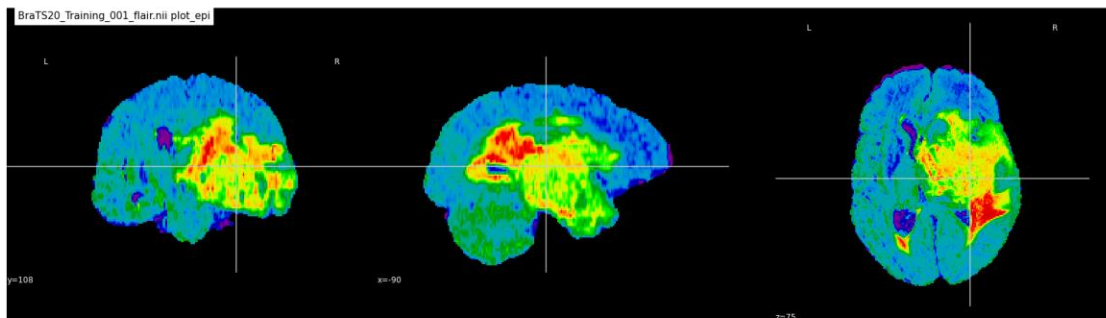
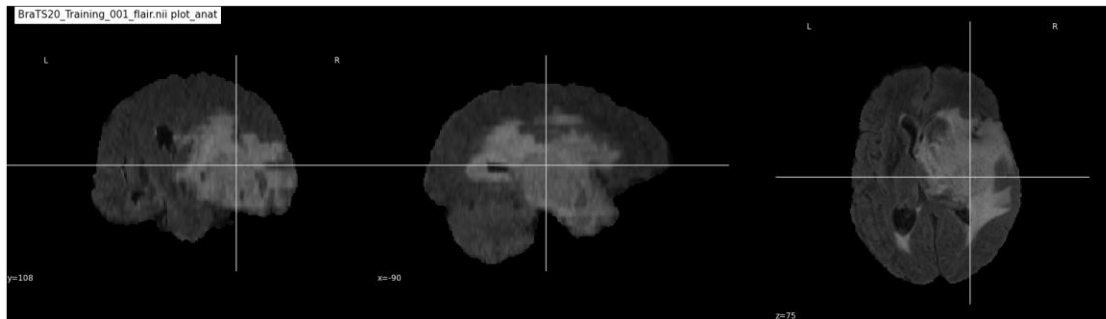
```
nlplt.plot_epi(niimg,
               title='BraTS20_Training_001_flair.nii plot_epi',
               axes=axes[1])
```

```
nlplt.plot_img(niimg,
```

```
        title='BraTS20_Training_001_flair.nii plot_img',
        axes=axes[2])

nlplt.plot_roi(nimask,
               title='BraTS20_Training_001_flair.nii with mask plot_roi',
               bg_img=niimg,
               axes=axes[3], cmap='Paired')

plt.show()
```



```
def dice_coef(y_true, y_pred, smooth=1.0):
    class_num = 4
    for i in range(class_num):
        y_true_f = K.flatten(y_true[:, :, :, i])
        y_pred_f = K.flatten(y_pred[:, :, :, i])
        intersection = K.sum(y_true_f * y_pred_f)
        loss = ((2. * intersection + smooth) / (K.sum(y_true_f) +
```



```

K.sum(y_pred_f) + smooth))
    if i == 0:
        total_loss = loss
    else:
        total_loss = total_loss + loss
total_loss = total_loss / class_num
return total_loss

def dice_coef_necrotic(y_true, y_pred, epsilon=1e-6):
    intersection = K.sum(K.abs(y_true[:, :, :, 1] * y_pred[:, :, :, 1]))
    return (2. * intersection) / (K.sum(K.square(y_true[:, :, :, 1])) +
K.sum(K.square(y_pred[:, :, :, 1])) + epsilon)

def dice_coef_edema(y_true, y_pred, epsilon=1e-6):
    intersection = K.sum(K.abs(y_true[:, :, :, 2] * y_pred[:, :, :, 2]))
    return (2. * intersection) / (K.sum(K.square(y_true[:, :, :, 2])) +
K.sum(K.square(y_pred[:, :, :, 2])) + epsilon)

def dice_coef_enhancing(y_true, y_pred, epsilon=1e-6):
    intersection = K.sum(K.abs(y_true[:, :, :, 3] * y_pred[:, :, :, 3]))
    return (2. * intersection) / (K.sum(K.square(y_true[:, :, :, 3])) +
K.sum(K.square(y_pred[:, :, :, 3])) + epsilon)

def precision(y_true, y_pred):
    true_positives = K.sum(K.round(K.clip(y_true * y_pred, 0, 1)))
    predicted_positives = K.sum(K.round(K.clip(y_pred, 0, 1)))
    precision = true_positives / (predicted_positives + K.epsilon())
    return precision

def sensitivity(y_true, y_pred):
    true_positives = K.sum(K.round(K.clip(y_true * y_pred, 0, 1)))
    possible_positives = K.sum(K.round(K.clip(y_true, 0, 1)))
    return true_positives / (possible_positives + K.epsilon())

def specificity(y_true, y_pred):
    true_negatives = K.sum(K.round(K.clip((1-y_true) * (1-y_pred), 0, 1)))
    possible_negatives = K.sum(K.round(K.clip(1-y_true, 0, 1)))
    return true_negatives / (possible_negatives + K.epsilon())

IMG_SIZE=128

from tensorflow.keras import layers, models, Input
from tensorflow.keras.layers import Conv2D, MaxPooling2D, UpSampling2D,
concatenate, Dropout, BatchNormalization

def build_improved_unet(inputs, ker_init, dropout):
    def conv_block(input_tensor, num_filters):
        conv = Conv2D(num_filters, 3, activation='relu', padding='same',
kernel_initializer=ker_init)(input_tensor)
        conv = BatchNormalization()(conv)

```

```

        conv = Conv2D(num_filters, 3, activation='relu', padding='same',
kernel_initializer=ker_init)(conv)
        conv = BatchNormalization()(conv)
        return conv

conv1 = conv_block(inputs, 32)
pool1 = MaxPooling2D(pool_size=(2, 2))(conv1)

conv2 = conv_block(pool1, 64)
pool2 = MaxPooling2D(pool_size=(2, 2))(conv2)

conv3 = conv_block(pool2, 128)
pool3 = MaxPooling2D(pool_size=(2, 2))(conv3)

conv4 = conv_block(pool3, 256)
pool4 = MaxPooling2D(pool_size=(2, 2))(conv4)

conv5 = conv_block(pool4, 512)
drop5 = Dropout(dropout)(conv5)

up6 = Conv2D(256, 2, activation='relu', padding='same',
kernel_initializer=ker_init)(UpSampling2D(size=(2, 2))(drop5))
merge6 = concatenate([conv4, up6], axis=3)
conv6 = conv_block(merge6, 256)

up7 = Conv2D(128, 2, activation='relu', padding='same',
kernel_initializer=ker_init)(UpSampling2D(size=(2, 2))(conv6))
merge7 = concatenate([conv3, up7], axis=3)
conv7 = conv_block(merge7, 128)

up8 = Conv2D(64, 2, activation='relu', padding='same',
kernel_initializer=ker_init)(UpSampling2D(size=(2, 2))(conv7))
merge8 = concatenate([conv2, up8], axis=3)
conv8 = conv_block(merge8, 64)

up9 = Conv2D(32, 2, activation='relu', padding='same',
kernel_initializer=ker_init)(UpSampling2D(size=(2, 2))(conv8))
merge9 = concatenate([conv1, up9], axis=3)
conv9 = conv_block(merge9, 32)

conv10 = Conv2D(4, (1, 1), activation='softmax')(conv9)

return models.Model(inputs=inputs, outputs=conv10)

input_layer = Input((IMG_SIZE, IMG_SIZE, 2))
improved_model = build_improved_unet(input_layer, 'he_normal', 0.2)
improved_model.compile(loss="categorical_crossentropy",
optimizer=keras.optimizers.Adam(learning_rate=0.001), metrics=['accuracy',

```

```
tf.keras.metrics.MeanIoU(num_classes=4), dice_coef, precision, sensitivity,  
specificity, dice_coef_necrotic, dice_coef_edema, dice_coef_enhancing])
```

```
improved_model.summary()
```

```
Model: "model"
```

Layer (type)	Output Shape	Param #	Connected to
=====			
input_1 (InputLayer)	[(None, 128, 128, 2)]	0	
=====			
conv2d (Conv2D) input_1[0][0]	(None, 128, 128, 32)	608	
=====			
batch_normalization (BatchNorma	(None, 128, 128, 32)	128	conv2d[0][0]
=====			
conv2d_1 (Conv2D) batch_normalization[0][0]	(None, 128, 128, 32)	9248	
=====			
batch_normalization_1 (BatchNor	(None, 128, 128, 32)	128	conv2d_1[0][0]
=====			
max_pooling2d (MaxPooling2D) batch_normalization_1[0][0]	(None, 64, 64, 32)	0	
=====			
conv2d_2 (Conv2D) max_pooling2d[0][0]	(None, 64, 64, 64)	18496	
=====			
batch_normalization_2 (BatchNor	(None, 64, 64, 64)	256	conv2d_2[0][0]
=====			
conv2d_3 (Conv2D) batch_normalization_2[0][0]	(None, 64, 64, 64)	36928	
=====			
batch_normalization_3 (BatchNor	(None, 64, 64, 64)	256	conv2d_3[0][0]
=====			
max_pooling2d_1 (MaxPooling2D)	(None, 32, 32, 64)	0	

batch\_normalization\_3[0][0]

---

conv2d_4 (Conv2D)	(None, 32, 32, 128)	73856
-------------------	---------------------	-------

---

max\_pooling2d\_1[0][0]

---

batch_normalization_4 (BatchNor	(None, 32, 32, 128)	512
---------------------------------	---------------------	-----

---

conv2d\_4[0][0]

---

conv2d_5 (Conv2D)	(None, 32, 32, 128)	147584
-------------------	---------------------	--------

---

batch\_normalization\_4[0][0]

---

batch_normalization_5 (BatchNor	(None, 32, 32, 128)	512
---------------------------------	---------------------	-----

---

conv2d\_5[0][0]

---

max_pooling2d_2 (MaxPooling2D)	(None, 16, 16, 128)	0
--------------------------------	---------------------	---

---

batch\_normalization\_5[0][0]

---

conv2d_6 (Conv2D)	(None, 16, 16, 256)	295168
-------------------	---------------------	--------

---

max\_pooling2d\_2[0][0]

---

batch_normalization_6 (BatchNor	(None, 16, 16, 256)	1024
---------------------------------	---------------------	------

---

conv2d\_6[0][0]

---

conv2d_7 (Conv2D)	(None, 16, 16, 256)	590080
-------------------	---------------------	--------

---

batch\_normalization\_6[0][0]

---

batch_normalization_7 (BatchNor	(None, 16, 16, 256)	1024
---------------------------------	---------------------	------

---

conv2d\_7[0][0]

---

max_pooling2d_3 (MaxPooling2D)	(None, 8, 8, 256)	0
--------------------------------	-------------------	---

---

batch\_normalization\_7[0][0]

---

conv2d_8 (Conv2D)	(None, 8, 8, 512)	1180160
-------------------	-------------------	---------

---

max\_pooling2d\_3[0][0]

---

batch_normalization_8 (BatchNor	(None, 8, 8, 512)	2048
---------------------------------	-------------------	------

---

conv2d\_8[0][0]

---

---

conv2d_9 (Conv2D) batch_normalization_8[0][0]	(None, 8, 8, 512)	2359808
--	-------------------	---------

---

batch_normalization_9 (BatchNor conv2d_9[0][0]	(None, 8, 8, 512)	2048
---	-------------------	------

---

dropout (Dropout) batch_normalization_9[0][0]	(None, 8, 8, 512)	0
--	-------------------	---

---

up_sampling2d (UpSampling2D) dropout[0][0]	(None, 16, 16, 512)	0
---	---------------------	---

---

conv2d_10 (Conv2D) up_sampling2d[0][0]	(None, 16, 16, 256)	524544
---	---------------------	--------

---

concatenate (Concatenate) batch_normalization_7[0][0]	(None, 16, 16, 512)	0
--	---------------------	---

---

conv2d\_10[0][0]

---

conv2d_11 (Conv2D) concatenate[0][0]	(None, 16, 16, 256)	1179904
---	---------------------	---------

---

batch_normalization_10 (BatchNo conv2d_11[0][0]	(None, 16, 16, 256)	1024
--	---------------------	------

---

conv2d_12 (Conv2D) batch_normalization_10[0][0]	(None, 16, 16, 256)	590080
--	---------------------	--------

---

batch_normalization_11 (BatchNo conv2d_12[0][0]	(None, 16, 16, 256)	1024
--	---------------------	------

---

up_sampling2d_1 (UpSampling2D) batch_normalization_11[0][0]	(None, 32, 32, 256)	0
--	---------------------	---

---

conv2d_13 (Conv2D) up_sampling2d_1[0][0]	(None, 32, 32, 128)	131200
---	---------------------	--------

---

---

concatenate_1 (Concatenate)	(None, 32, 32, 256)	0
batch_normalization_5[0][0]		

conv2d\_13[0][0]

---

---

conv2d_14 (Conv2D)	(None, 32, 32, 128)	295040
concatenate_1[0][0]		

---

---

batch_normalization_12 (Batch Normalization)	(None, 32, 32, 128)	512
conv2d_14[0][0]		

---

---

conv2d_15 (Conv2D)	(None, 32, 32, 128)	147584
batch_normalization_12[0][0]		

---

---

batch_normalization_13 (Batch Normalization)	(None, 32, 32, 128)	512
conv2d_15[0][0]		

---

---

up_sampling2d_2 (UpSampling2D)	(None, 64, 64, 128)	0
batch_normalization_13[0][0]		

---

---

conv2d_16 (Conv2D)	(None, 64, 64, 64)	32832
up_sampling2d_2[0][0]		

---

---

concatenate_2 (Concatenate)	(None, 64, 64, 128)	0
batch_normalization_3[0][0]		

conv2d\_16[0][0]

---

---

conv2d_17 (Conv2D)	(None, 64, 64, 64)	73792
concatenate_2[0][0]		

---

---

batch_normalization_14 (Batch Normalization)	(None, 64, 64, 64)	256
conv2d_17[0][0]		

---

---

conv2d_18 (Conv2D)	(None, 64, 64, 64)	36928
batch_normalization_14[0][0]		

---

---

batch_normalization_15 (Batch Normalization)	(None, 64, 64, 64)	256
--	--------------------	-----

conv2d\_18[0][0]

---

up\_sampling2d\_3 (UpSampling2D) (None, 128, 128, 64) 0  
batch\_normalization\_15[0][0]

---

conv2d\_19 (Conv2D) (None, 128, 128, 32) 8224  
up\_sampling2d\_3[0][0]

---

concatenate\_3 (Concatenate) (None, 128, 128, 64) 0  
batch\_normalization\_1[0][0]

---

conv2d\_19[0][0]

---

conv2d\_20 (Conv2D) (None, 128, 128, 32) 18464  
concatenate\_3[0][0]

---

batch\_normalization\_16 (BatchNormaliz (None, 128, 128, 32) 128  
conv2d\_20[0][0]

---

conv2d\_21 (Conv2D) (None, 128, 128, 32) 9248  
batch\_normalization\_16[0][0]

---

batch\_normalization\_17 (BatchNormaliz (None, 128, 128, 32) 128  
conv2d\_21[0][0]

---

conv2d\_22 (Conv2D) (None, 128, 128, 4) 132  
batch\_normalization\_17[0][0]

---

=====  
Total params: 7,771,684  
Trainable params: 7,765,796  
Non-trainable params: 5,888

---

```
def build_unet(inputs, ker_init, dropout):  
    conv1 = Conv2D(32, 3, activation = 'relu', padding = 'same',  
kernel_initializer = ker_init)(inputs)  
    conv1 = Conv2D(32, 3, activation = 'relu', padding = 'same',  
kernel_initializer = ker_init)(conv1)  
  
    pool = MaxPooling2D(pool_size=(2, 2))(conv1)
```

```

conv = Conv2D(64, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(pool)
conv = Conv2D(64, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(conv)

pool1 = MaxPooling2D(pool_size=(2, 2))(conv)
conv2 = Conv2D(128, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(pool1)
conv2 = Conv2D(128, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(conv2)

pool2 = MaxPooling2D(pool_size=(2, 2))(conv2)
conv3 = Conv2D(256, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(pool2)
conv3 = Conv2D(256, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(conv3)

pool4 = MaxPooling2D(pool_size=(2, 2))(conv3)
conv5 = Conv2D(512, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(pool4)
conv5 = Conv2D(512, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(conv5)
drop5 = Dropout(dropout)(conv5)

up7 = Conv2D(256, 2, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(UpSampling2D(size = (2,2))(drop5))
merge7 = concatenate([conv3,up7], axis = 3)
conv7 = Conv2D(256, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(merge7)
conv7 = Conv2D(256, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(conv7)

up8 = Conv2D(128, 2, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(UpSampling2D(size = (2,2))(conv7))
merge8 = concatenate([conv2,up8], axis = 3)
conv8 = Conv2D(128, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(merge8)
conv8 = Conv2D(128, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(conv8)

up9 = Conv2D(64, 2, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(UpSampling2D(size = (2,2))(conv8))
merge9 = concatenate([conv,up9], axis = 3)
conv9 = Conv2D(64, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(merge9)
conv9 = Conv2D(64, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(conv9)

```



```

        up = Conv2D(32, 2, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(UpSampling2D(size = (2,2))(conv9))
        merge = concatenate([conv1,up], axis = 3)
        conv = Conv2D(32, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(merge)
        conv = Conv2D(32, 3, activation = 'relu', padding = 'same',
kernel_initializer = ker_init)(conv)

        conv10 = Conv2D(4, (1,1), activation = 'softmax')(conv)

        return Model(inputs = inputs, outputs = conv10)

input_layer = Input((IMG_SIZE, IMG_SIZE, 2))

model = build_unet(input_layer, 'he_normal', 0.2)
model.compile(loss="categorical_crossentropy",
optimizer=keras.optimizers.Adam(learning_rate=0.001), metrics =
['accuracy',tf.keras.metrics.MeanIoU(num_classes=4), dice_coef, precision,
sensitivity, specificity, dice_coef_necrotic, dice_coef_edema
,dice_coef_enhancing] )

plot_model(improved_model,
            show_shapes = True,
            show_dtype=False,
            show_layer_names = True,
            rankdir = 'TB',
            expand_nested = False,
            dpi = 70)

```

```

train_and_val_directories = [f.path for f in os.scandir(TRAIN_DATASET_PATH)
if f.is_dir()]

```

```

train_and_val_directories.remove(TRAIN_DATASET_PATH+'BraTS20_Training_355')

```

```

def pathListIntoIds(dirList):
    x = []
    for i in range(0,len(dirList)):
        x.append(dirList[i][dirList[i].rfind('/')+1:])
    return x

```

```

train_and_test_ids = pathListIntoIds(train_and_val_directories);

```

```

train_test_ids, val_ids = train_test_split(train_and_test_ids,test_size=0.2)
train_ids, test_ids = train_test_split(train_test_ids,test_size=0.15)

```

```

class DataGenerator(keras.utils.Sequence):
    'Generates data for Keras'
    def __init__(self, list_IDS, dim=(IMG_SIZE,IMG_SIZE), batch_size = 1,
n_channels = 2, shuffle=True):
        'Initialization'
        self.dim = dim
        self.batch_size = batch_size
        self.list_IDS = list_IDS
        self.n_channels = n_channels
        self.shuffle = shuffle
        self.on_epoch_end()

    def __len__(self):
        'Denotes the number of batches per epoch'
        return int(np.floor(len(self.list_IDS) / self.batch_size))

    def __getitem__(self, index):
        'Generate one batch of data'
        indexes =
self.indexes[index*self.batch_size:(index+1)*self.batch_size]
        Batch_ids = [self.list_IDS[k] for k in indexes]
        X, y = self.__data_generation(Batch_ids)

        return X, y

    def on_epoch_end(self):
        'Updates indexes after each epoch'
        self.indexes = np.arange(len(self.list_IDS))
        if self.shuffle == True:
            np.random.shuffle(self.indexes)

    def __data_generation(self, Batch_ids):
        'Generates data containing batch_size samples'
        X = np.zeros((self.batch_size*VOLUME_SLICES, *self.dim,
self.n_channels))
        y = np.zeros((self.batch_size*VOLUME_SLICES, 240, 240))
        Y = np.zeros((self.batch_size*VOLUME_SLICES, *self.dim, 4))

        for c, i in enumerate(Batch_ids):
            case_path = os.path.join(TRAIN_DATASET_PATH, i)

            data_path = os.path.join(case_path, f'{i}_flair.nii');
            flair = nib.load(data_path).get_fdata()

            data_path = os.path.join(case_path, f'{i}_t1ce.nii');
            ce = nib.load(data_path).get_fdata()

            data_path = os.path.join(case_path, f'{i}_seg.nii');

```

```

        seg = nib.load(data_path).get_fdata()

        for j in range(VOLUME_SLICES):
            X[j + VOLUME_SLICES*c, :, :, 0] =
cv2.resize(flair[:, :, j+VOLUME_START_AT], (IMG_SIZE, IMG_SIZE));
            X[j + VOLUME_SLICES*c, :, :, 1] =
cv2.resize(ce[:, :, j+VOLUME_START_AT], (IMG_SIZE, IMG_SIZE));

            y[j + VOLUME_SLICES*c] = seg[:, :, j+VOLUME_START_AT];

        y[y==4] = 3;
        mask = tf.one_hot(y, 4);
        Y = tf.image.resize(mask, (IMG_SIZE, IMG_SIZE));
        return X/np.max(X), Y

training_generator = DataGenerator(train_ids)
valid_generator = DataGenerator(val_ids)
test_generator = DataGenerator(test_ids)

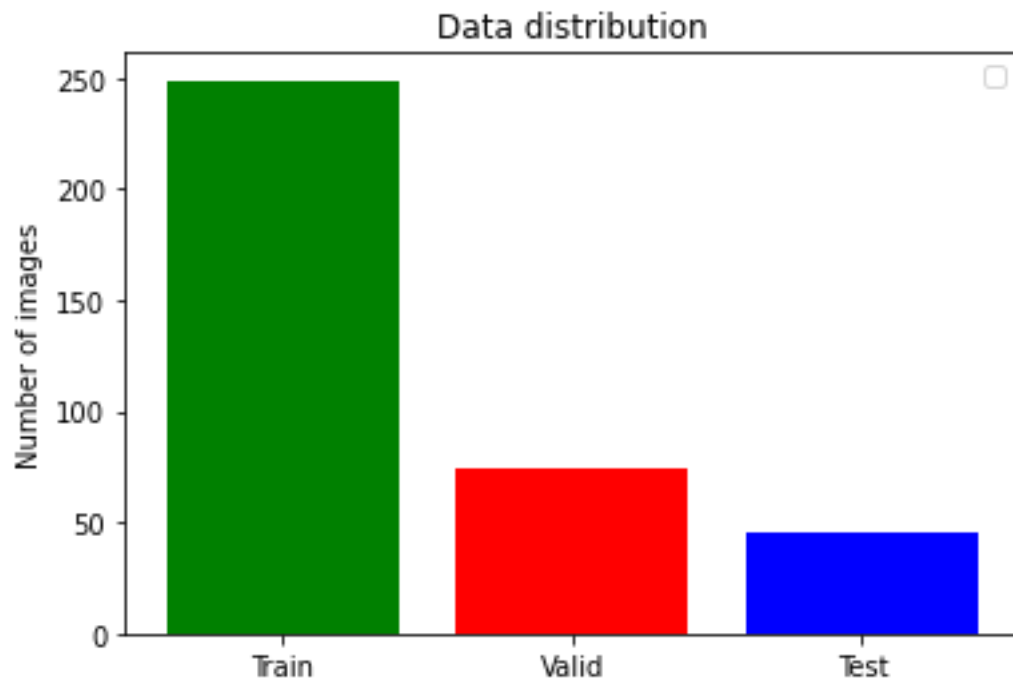
def showDataLayout():
    plt.bar(["Train", "Valid", "Test"],
            [len(train_ids), len(val_ids), len(test_ids)], align='center', color=[
'green', 'red', 'blue'])
    plt.legend()

    plt.ylabel('Number of images')
    plt.title('Data distribution')

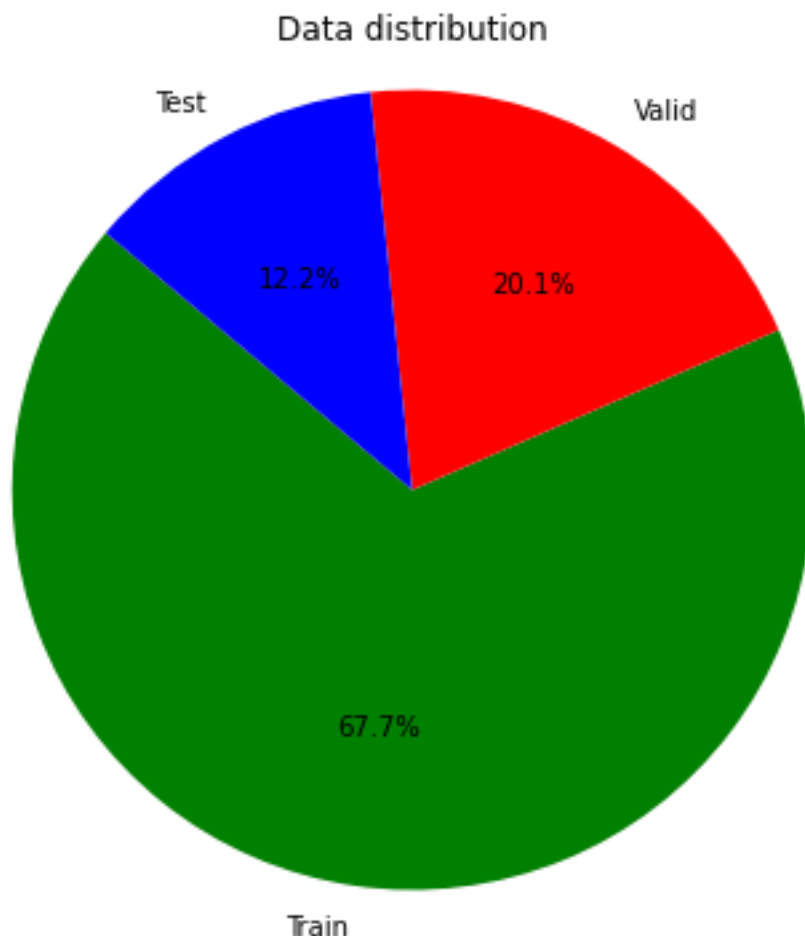
    plt.show()

showDataLayout()

```



```
def showDataLayout():  
    labels = ["Train", "Valid", "Test"]  
    sizes = [len(train_ids), len(val_ids), len(test_ids)]  
    colors = ['green', 'red', 'blue']  
  
    plt.figure(figsize=(6, 6))  
    plt.pie(sizes, labels=labels, colors=colors, autopct='%1.1f%%',  
startangle=140)  
    plt.title('Data distribution')  
    plt.axis('equal')  
  
    plt.show()  
  
showDataLayout()
```



```
csv_logger = CSVLogger('training.log', separator=',', append=False)
callbacks = [
    keras.callbacks.ReduceLROnPlateau(monitor='val_loss', factor=0.2,
                                       patience=2, min_lr=0.000001, verbose=1),
    csv_logger
]
```

```
K.clear_session()
history = improved_model.fit(training_generator,
                             epochs=5,
                             steps_per_epoch=len(train_ids),
                             callbacks= callbacks,
                             validation_data = valid_generator
                             )
```

Epoch 1/5

249/249 [=====] - 216s 833ms/step - loss: 0.8919 - accuracy: 0.8354 - mean\_io\_u: 0.3756 - dice\_coef: 0.1800 - precision: 0.7696 - sensitivity: 0.4265 - specificity: 0.9878 - dice\_coef\_necrotic: 0.0579 - dice\_coef\_edema: 0.2102 - dice\_coef\_enhancing: 0.0956 - val\_loss: 3.9397 - val\_accuracy: 0.6902 - val\_mean\_io\_u: 0.3756 - val\_dice\_coef: 0.2108 - val\_precision: 0.8034 - val\_sensitivity: 0.6487 - val\_specificity: 0.9470 -

```

val_dice_coef_necrotic: 0.0473 - val_dice_coef_edema: 0.1420 -
val_dice_coef_enhancing: 0.0503
Epoch 2/5
249/249 [=====] - 118s 472ms/step - loss: 0.0831 -
accuracy: 0.9882 - mean_io_u: 0.3756 - dice_coef: 0.3458 - precision: 0.9905
- sensitivity: 0.9856 - specificity: 0.9968 - dice_coef_necrotic: 0.2357 -
dice_coef_edema: 0.5405 - dice_coef_enhancing: 0.4251 - val_loss: 1.1279 -
val_accuracy: 0.9465 - val_mean_io_u: 0.3756 - val_dice_coef: 0.2988 -
val_precision: 0.9523 - val_sensitivity: 0.9425 - val_specificity: 0.9844 -
val_dice_coef_necrotic: 0.0863 - val_dice_coef_edema: 0.2563 -
val_dice_coef_enhancing: 0.2466
Epoch 3/5
249/249 [=====] - 117s 469ms/step - loss: 0.0461 -
accuracy: 0.9894 - mean_io_u: 0.3756 - dice_coef: 0.4208 - precision: 0.9910
- sensitivity: 0.9872 - specificity: 0.9970 - dice_coef_necrotic: 0.2761 -
dice_coef_edema: 0.5894 - dice_coef_enhancing: 0.5515 - val_loss: 2.3090 -
val_accuracy: 0.9433 - val_mean_io_u: 0.3756 - val_dice_coef: 0.3214 -
val_precision: 0.9510 - val_sensitivity: 0.9400 - val_specificity: 0.9841 -
val_dice_coef_necrotic: 0.1185 - val_dice_coef_edema: 0.1589 -
val_dice_coef_enhancing: 0.3189
Epoch 4/5
249/249 [=====] - 117s 470ms/step - loss: 0.0377 -
accuracy: 0.9899 - mean_io_u: 0.3756 - dice_coef: 0.4502 - precision: 0.9913
- sensitivity: 0.9879 - specificity: 0.9971 - dice_coef_necrotic: 0.3002 -
dice_coef_edema: 0.5732 - dice_coef_enhancing: 0.5718 - val_loss: 0.5554 -
val_accuracy: 0.9547 - val_mean_io_u: 0.3756 - val_dice_coef: 0.3183 -
val_precision: 0.9627 - val_sensitivity: 0.9502 - val_specificity: 0.9878 -
val_dice_coef_necrotic: 0.0779 - val_dice_coef_edema: 0.1482 -
val_dice_coef_enhancing: 0.2835
Epoch 5/5
249/249 [=====] - 117s 470ms/step - loss: 0.0315 -
accuracy: 0.9905 - mean_io_u: 0.3756 - dice_coef: 0.4872 - precision: 0.9918
- sensitivity: 0.9886 - specificity: 0.9972 - dice_coef_necrotic: 0.3310 -
dice_coef_edema: 0.6118 - dice_coef_enhancing: 0.6151 - val_loss: 0.6838 -
val_accuracy: 0.9528 - val_mean_io_u: 0.3756 - val_dice_coef: 0.3379 -
val_precision: 0.9588 - val_sensitivity: 0.9500 - val_specificity: 0.9866 -
val_dice_coef_necrotic: 0.1131 - val_dice_coef_edema: 0.1659 -
val_dice_coef_enhancing: 0.2944

```

```

-----
NameError                                Traceback (most recent call last)
<ipython-input-18-08b3f0c01a12> in <module>
      6                 validation_data = valid_generator
      7             )
----> 8 model.save("3D_MRI_Brain_tumor_segmentation.h5")

```

NameError: name 'model' is not defined

```

improved_model.save('3D_MRI_Brain_tumor_segmentation.h5')
print("Model saved successfully as 3D_MRI_Brain_tumor_segmentation.h5")

```

Model saved successfully as 3D\_MRI\_Brain\_tumor\_segmentation.h5

```
import tensorflow as tf
import numpy as np
import nibabel as nib
import cv2
```

```
model = tf.keras.models.load_model("3D_MRI_Brain_tumor_segmentation.h5",
custom_objects={
    'dice_coef': dice_coef,
    'precision': precision,
    'sensitivity': sensitivity,
    'specificity': specificity,
    'dice_coef_necrotic': dice_coef_necrotic,
    'dice_coef_edema': dice_coef_edema,
    'dice_coef_enhancing': dice_coef_enhancing
})
```

```
def preprocess_image(image_file, slice_index=None):
```

```
    img = nib.load(image_file).get_fdata()
```

```
    if slice_index is not None:
        img = img[:, :, slice_index]
```

```
    img_resized = cv2.resize(img, (IMG_SIZE, IMG_SIZE))
```

```
    img_resized = img_resized / np.max(img_resized)
```

```
    return img_resized
```

```
def predict(image_paths, slice_index):
```

```
    X = np.zeros((1, IMG_SIZE, IMG_SIZE, 2))
```

```
    X[0, :, :, 0] = preprocess_image(image_paths[0], slice_index)
```

```
    X[0, :, :, 1] = preprocess_image(image_paths[1], slice_index)
```

```
    pred = model.predict(X)
```

```
    return np.argmax(pred[0], axis=-1)
```

```
image_paths = [
```

```
    '/kaggle/input/brats20-dataset-training-
validation/BraTS2020_ValidationData/MICCAI_BraTS2020_ValidationData/BraTS20_V
alidation_010/BraTS20_Validation_010_flair.nii',
```

```
    '/kaggle/input/brats20-dataset-training-
validation/BraTS2020_ValidationData/MICCAI_BraTS2020_ValidationData/BraTS20_V
alidation_010/BraTS20_Validation_010_t1ce.nii'
```

```
]
```

```

slice_index = 75
prediction = predict(image_paths, slice_index)

import numpy as np

def get_classification(pred):
    class_predictions = np.argmax(pred, axis=-1)
    return class_predictions

get_classification(prediction)

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,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,
        94,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,
         0,  0,  0, 79,  0,  0,  0, 38, 36, 35, 37, 38, 38, 38, 39, 73, 73,
        72, 71, 71, 75, 75, 76, 77, 76, 75, 75, 54, 55, 55, 56, 58, 61, 75,
        75, 75, 75, 76, 76, 76, 77, 78,  0,  0,  0,  0,  0,  0,  0,  0,  0,
         0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,  0,
         0,  0,  0,  0,  0,  0,  0,  0,  0])

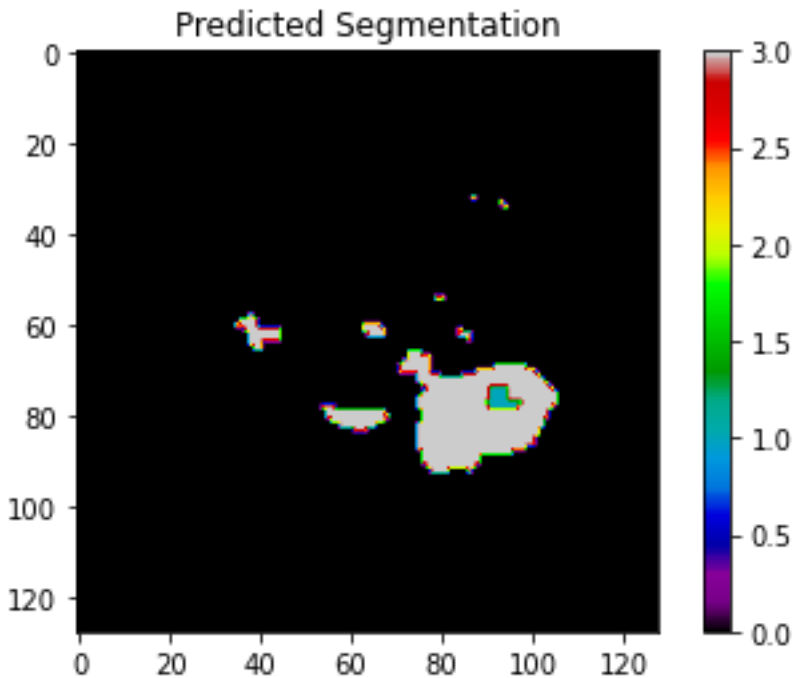
import matplotlib.pyplot as plt

def visualize_prediction(prediction):
    plt.imshow(prediction, cmap='nipy_spectral')
    plt.title('Predicted Segmentation')
    plt.colorbar()
    plt.show()

visualize_prediction(prediction)

```





```

model =
keras.models.load_model('../input/modelperclasseval/model_per_class.h5',
                        custom_objects={ 'accuracy' :
tf.keras.metrics.MeanIoU(num_classes=4),

                                "dice_coef": dice_coef,
                                "precision": precision,
                                "sensitivity":sensitivity,
                                "specificity":specificity,
                                "dice_coef_necrotic":

dice_coef_necrotic,

                                "dice_coef_edema":

dice_coef_edema,

                                "dice_coef_enhancing":

dice_coef_enhancing

                                }, compile=False)

history = pd.read_csv('../input/modelperclasseval/training_per_class.log',
sep=',', engine='python')

hist=history

acc=hist['accuracy']
val_acc=hist['val_accuracy']

epoch=range(len(acc))

loss=hist['loss']
val_loss=hist['val_loss']

```

```

train_dice=hist['dice_coef']
val_dice=hist['val_dice_coef']

f,ax=plt.subplots(1,4,figsize=(16,8))

ax[0].plot(epoch,acc,'b',label='Training Accuracy')
ax[0].plot(epoch,val_acc,'r',label='Validation Accuracy')
ax[0].legend()

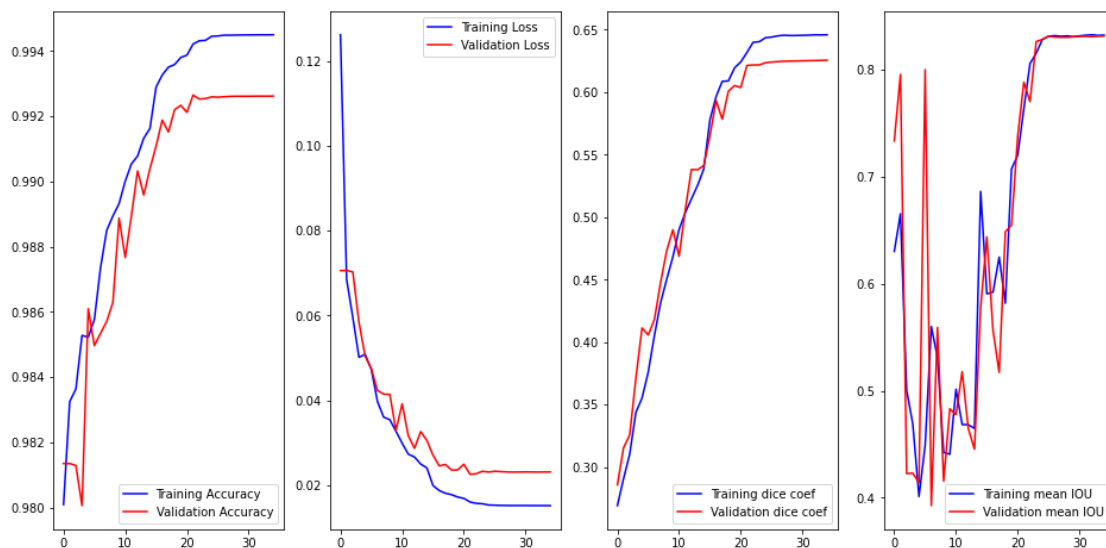
ax[1].plot(epoch,loss,'b',label='Training Loss')
ax[1].plot(epoch,val_loss,'r',label='Validation Loss')
ax[1].legend()

ax[2].plot(epoch,train_dice,'b',label='Training dice coef')
ax[2].plot(epoch,val_dice,'r',label='Validation dice coef')
ax[2].legend()

ax[3].plot(epoch,hist['mean_io_u'],'b',label='Training mean IOU')
ax[3].plot(epoch,hist['val_mean_io_u'],'r',label='Validation mean IOU')
ax[3].legend()

plt.show()

```



```

def imageLoader(path):
    image = nib.load(path).get_fdata()
    X = np.zeros((self.batch_size*VOLUME_SLICES, *self.dim, self.n_channels))
    for j in range(VOLUME_SLICES):
        X[j +VOLUME_SLICES*c,:,:,:] =
cv2.resize(image[:,:,:j+VOLUME_START_AT], (IMG_SIZE, IMG_SIZE));
        X[j +VOLUME_SLICES*c,:,:,:] = cv2.resize(ce[:,:,:j+VOLUME_START_AT],
(IMG_SIZE, IMG_SIZE));

```

```

        y[j + VOLUME_SLICES*c] = seg[:, :, j + VOLUME_START_AT];
    return np.array(image)

def loadDataFromDir(path, list_of_files, mriType, n_images):
    scans = []
    masks = []
    for i in list_of_files[:n_images]:
        fullPath = glob.glob( i + '/*'+ mriType + '*')[0]
        currentScanVolume = imageLoader(fullPath)
        currentMaskVolume = imageLoader( glob.glob( i + '/*seg*')[0] )

        for j in range(0, currentScanVolume.shape[2]):
            scan_img = cv2.resize(currentScanVolume[:, :, j],
                dsize=(IMG_SIZE, IMG_SIZE), interpolation=cv2.INTER_AREA).astype('uint8')
            mask_img = cv2.resize(currentMaskVolume[:, :, j],
                dsize=(IMG_SIZE, IMG_SIZE), interpolation=cv2.INTER_AREA).astype('uint8')
            scans.append(scan_img[..., np.newaxis])
            masks.append(mask_img[..., np.newaxis])
    return np.array(scans, dtype='float32'), np.array(masks, dtype='float32')

def predictByPath(case_path, case):
    files = next(os.walk(case_path))[2]
    X = np.empty((VOLUME_SLICES, IMG_SIZE, IMG_SIZE, 2))

    vol_path = os.path.join(case_path, f'BraTS20_Training_{case}_flair.nii');
    flair=nib.load(vol_path).get_fdata()

    vol_path = os.path.join(case_path, f'BraTS20_Training_{case}_t1ce.nii');
    ce=nib.load(vol_path).get_fdata()

    for j in range(VOLUME_SLICES):
        X[j, :, :, 0] = cv2.resize(flair[:, :, j + VOLUME_START_AT],
            (IMG_SIZE, IMG_SIZE))
        X[j, :, :, 1] = cv2.resize(ce[:, :, j + VOLUME_START_AT],
            (IMG_SIZE, IMG_SIZE))

    return model.predict(X/np.max(X), verbose=1)

def showPredictsById(case, start_slice = 60):
    path = f"../input/brats20-dataset-training-validation/BraTS2020_TrainingData/MICCAI_BraTS2020_TrainingData/BraTS20_Training_{case}"
    gt = nib.load(os.path.join(path,
        f'BraTS20_Training_{case}_seg.nii')).get_fdata()
    origImage = nib.load(os.path.join(path,
        f'BraTS20_Training_{case}_flair.nii')).get_fdata()
    p = predictByPath(path, case)

```

```

core = p[:, :, :, 1]
edema= p[:, :, :, 2]
enhancing = p[:, :, :, 3]

plt.figure(figsize=(18, 50))
f, axarr = plt.subplots(1,6, figsize = (18, 50))

for i in range(6):

axarr[i].imshow(cv2.resize(origImage[:, :, start_slice+VOLUME_START_AT],
(IMG_SIZE, IMG_SIZE)), cmap="gray", interpolation='none')

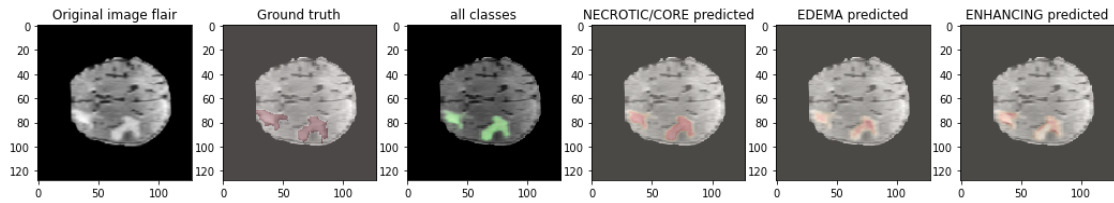
    axarr[0].imshow(cv2.resize(origImage[:, :, start_slice+VOLUME_START_AT],
(IMG_SIZE, IMG_SIZE)), cmap="gray")
    axarr[0].title.set_text('Original image flair')
    curr_gt=cv2.resize(gt[:, :, start_slice+VOLUME_START_AT], (IMG_SIZE,
IMG_SIZE), interpolation = cv2.INTER_NEAREST)
    axarr[1].imshow(curr_gt, cmap="Reds", interpolation='none', alpha=0.3) #
, alpha=0.3, cmap='Reds'
    axarr[1].title.set_text('Ground truth')
    axarr[2].imshow(p[start_slice, :, :, 1:4], cmap="Reds",
interpolation='none', alpha=0.3)
    axarr[2].title.set_text('all classes')
    axarr[3].imshow(edema[start_slice, :, :], cmap="OrRd",
interpolation='none', alpha=0.3)
    axarr[3].title.set_text(f'{SEGMENT_CLASSES[1]} predicted')
    axarr[4].imshow(core[start_slice, :, :], cmap="OrRd", interpolation='none',
alpha=0.3)
    axarr[4].title.set_text(f'{SEGMENT_CLASSES[2]} predicted')
    axarr[5].imshow(enhancing[start_slice, :, :], cmap="OrRd",
interpolation='none', alpha=0.3)
    axarr[5].title.set_text(f'{SEGMENT_CLASSES[3]} predicted')
    plt.show()

showPredictsById(case=test_ids[0][-3:])
showPredictsById(case=test_ids[1][-3:])
showPredictsById(case=test_ids[2][-3:])
showPredictsById(case=test_ids[3][-3:])
showPredictsById(case=test_ids[4][-3:])
showPredictsById(case=test_ids[5][-3:])
showPredictsById(case=test_ids[6][-3:])

4/4 [=====] - 1s 96ms/step

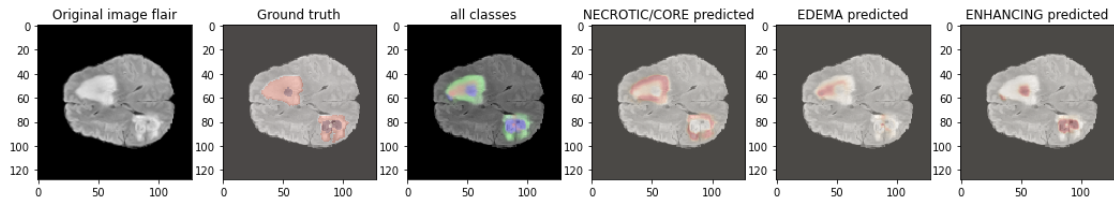
<Figure size 1296x3600 with 0 Axes>

```



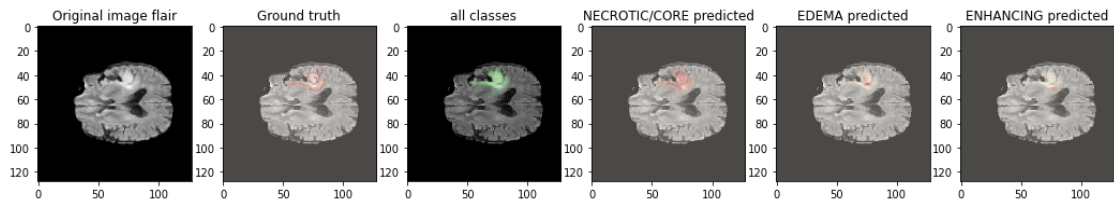
4/4 [=====] - 0s 33ms/step

<Figure size 1296x3600 with 0 Axes>



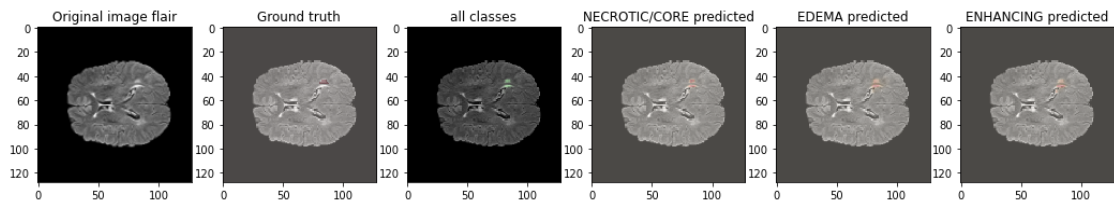
4/4 [=====] - 0s 33ms/step

<Figure size 1296x3600 with 0 Axes>



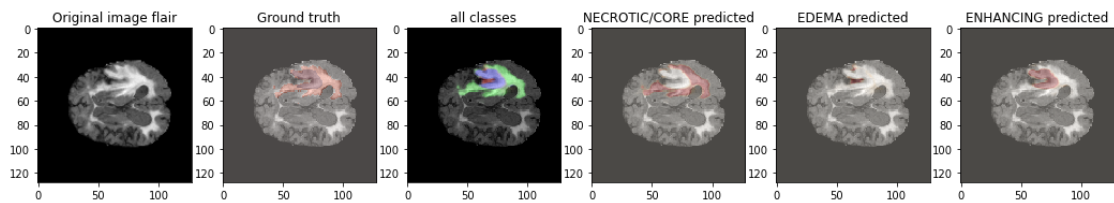
4/4 [=====] - 0s 33ms/step

<Figure size 1296x3600 with 0 Axes>



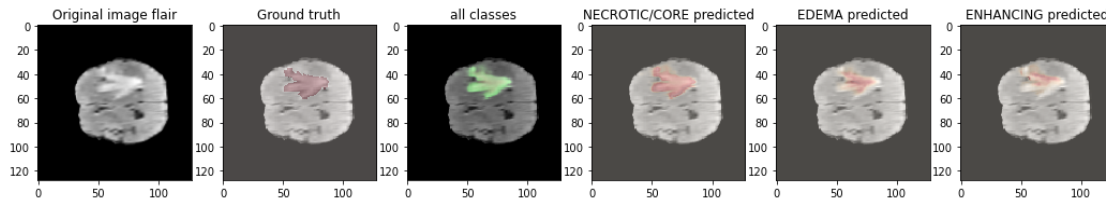
4/4 [=====] - 0s 33ms/step

<Figure size 1296x3600 with 0 Axes>



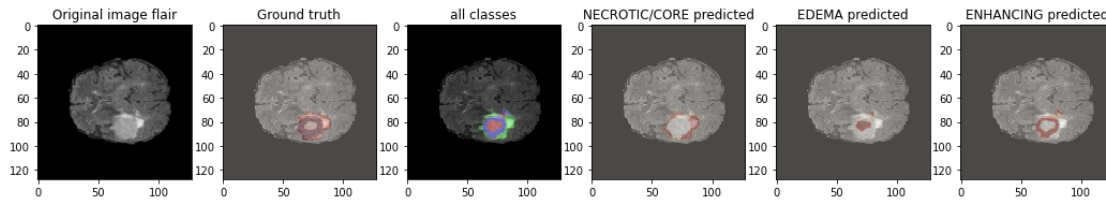
4/4 [=====] - 0s 33ms/step

<Figure size 1296x3600 with 0 Axes>



4/4 [=====] - 0s 33ms/step

<Figure size 1296x3600 with 0 Axes>



```
case = case=test_ids[3][-3:]
path = f"../input/brats20-dataset-training-validation/BraTS2020_TrainingData/MICCAI_BraTS2020_TrainingData/BraTS20_Training_{case}"
gt = nib.load(os.path.join(path,
f'BraTS20_Training_{case}_seg.nii')).get_fdata()
p = predictByPath(path,case)
```

```
core = p[:, :, :, 1]
edema= p[:, :, :, 2]
enhancing = p[:, :, :, 3]
```

```
i=40
eval_class = 2
```

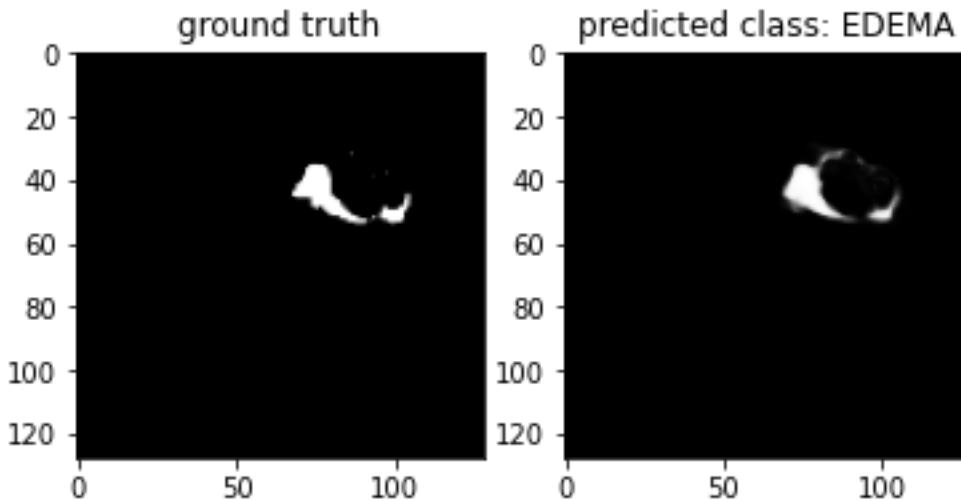
```
gt[gt != eval_class] = 1
```

```
resized_gt = cv2.resize(gt[:, :, i+VOLUME_START_AT], (IMG_SIZE, IMG_SIZE))
```

```
plt.figure()
f, axarr = plt.subplots(1,2)
axarr[0].imshow(resized_gt, cmap="gray")
axarr[0].title.set_text('ground truth')
axarr[1].imshow(p[i, :, :, eval_class], cmap="gray")
axarr[1].title.set_text(f'predicted class: {SEGMENT_CLASSES[eval_class]}')
plt.show()
```

4/4 [=====] - 0s 36ms/step

<Figure size 432x288 with 0 Axes>



```
improved_model.compile(loss="categorical_crossentropy",
optimizer=keras.optimizers.Adam(learning_rate=0.001), metrics =
['accuracy',tf.keras.metrics.MeanIOU(num_classes=4), dice_coef, precision,
sensitivity, specificity, dice_coef_necrotic, dice_coef_edema,
dice_coef_enhancing] )
print("Evaluate on test data")
results = improved_model.evaluate(test_generator, batch_size=100, callbacks=
callbacks)
print("test loss, test acc:", results)
```

Evaluate on test data

```
45/45 [=====] - 25s 538ms/step - loss: 1.3613 -
accuracy: 0.9620 - mean_io_u_1: 0.3755 - dice_coef: 0.3311 - precision:
0.9663 - sensitivity: 0.9603 - specificity: 0.9890 - dice_coef_necrotic:
0.1275 - dice_coef_edema: 0.1491 - dice_coef_enhancing: 0.2706
test loss, test acc: [1.4527249336242676, 0.9519450664520264,
0.37558794021606445, 0.33874499797821045, 0.9571685791015625,
0.9493675231933594, 0.9859769940376282, 0.122862309217453,
0.18301647901535034, 0.2746582329273224]
```