## **Coursework 2: Image segmentation**

In this coursework you will develop and train a convolutional neural network for brain tumour image segmentation. Please read both the text and the code in this notebook to get an idea what you are expected to implement. Pay attention to the missing code blocks that look like this:

```
### Insert your code ###
...
### End of your code ###
```

## What to do?

- Complete and run the code using jupyter-lab or jupyter-notebook to get the results.
- Export (File | Save and Export Notebook As...) the notebook as a PDF file, which contains your code, results and answers, and upload the PDF file onto <a href="Scientia.doc.ic.ac.uk">Scientia.doc.ic.ac.uk</a>).
- Instead of clicking the Export button, you can also run the following command instead: jupyter nbconvert coursework.ipynb --to pdf
- If Jupyter complains about some problems in exporting, it is likely that pandoc
   (<a href="https://pandoc.org/installing.html">https://pandoc.org/installing.html</a>) or latex is not installed, or their
   paths have not been included. You can install the relevant libraries and retry.
- If Jupyter-lab does not work for you at the end, you can use Google Colab to write the code and export the PDF file.

## **Dependencies**

You need to install Jupyter-Lab (<a href="https://jupyterlab.readthedocs.io/en/stable/getting\_started/installation.html">https://jupyterlab.readthedocs.io/en/stable/getting\_started/installation.html</a>)) and other libraries used in this coursework, such as by running the command: pip3 install [package\_name]

## **GPU** resource

The coursework is developed to be able to run on CPU, as all images have been pre-processed to be 2D and of a smaller size, compared to original 3D volumes.

However, to save training time, you may want to use GPU. In that case, you can run this notebook on Google Colab. On Google Colab, go to the menu, Runtime - Change runtime type, and select **GPU** as the hardware acceleartor. At the end, please still export everything and submit as a PDF file on Scientia.

#### In [1]:

```
# Import libraries
# These libraries should be sufficient for this tutorial.
# However, if any other library is needed, please install by yourself.
import tarfile
import imageio
import torch
import torch.nn as nn
import torch.nn.functional as F
import torch.optim as optim
from torch.utils.data import Dataset
import numpy as np
import time
import os
import random
import matplotlib.pyplot as plt
from matplotlib import colors
```

## 1. Download and visualise the imaging dataset.

The dataset is curated from the brain imaging dataset in <u>Medical Decathlon Challenge</u> (<a href="http://medicaldecathlon.com/">http://medicaldecathlon.com/</a>). To save the storage and reduce the computational cost for this tutorial, we extract 2D image slices from T1-Gd contrast enhanced 3D brain volumes and downsample the images.

The dataset consists of a training set and a test set. Each image is of dimension 120 x 120, with a corresponding label map of the same dimension. There are four number of classes in the label map:

- · 0: background
- 1: edema
- · 2: non-enhancing tumour
- 3: enhancing tumour

### In [2]:

```
# Download the dataset
!wget https://www.dropbox.com/s/zmytk2yu284af6t/Task01_BrainTumour_2D.tar.gz
# Unzip the '.tar.gz' file to the current directory
datafile = tarfile.open('Task01_BrainTumour_2D.tar.gz')
datafile.extractall()
datafile.close()
```

```
--2023-03-03 13:17:14-- https://www.dropbox.com/s/zmytk2yu284af6t/Tas
k01 BrainTumour 2D.tar.gz (https://www.dropbox.com/s/zmytk2yu284af6t/T
ask01 BrainTumour 2D.tar.gz)
Resolving www.dropbox.com (www.dropbox.com)... 162.125.1.18, 2620:100:
6016:18::a27d:112
Connecting to www.dropbox.com (www.dropbox.com) | 162.125.1.18 | :443... c
onnected.
HTTP request sent, awaiting response... 302 Found
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--2023-03-03 13:17:15-- https://www.dropbox.com/s/raw/zmytk2yu284af6
t/Task01 BrainTumour 2D.tar.gz (https://www.dropbox.com/s/raw/zmytk2yu
284af6t/Task01 BrainTumour 2D.tar.gz)
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HTTP request sent, awaiting response... 302 Found
Location: https://uc04907ead56e5d8bd17897f5bed.dl.dropboxusercontent.c
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yAHndhF8PnOUT-aIesswSuSIKabDJy-dDDXoD6vWVwB9v5kFg/file# (https://uc049
07ead56e5d8bd17897f5bed.dl.dropboxusercontent.com/cd/0/inline/B3iC93bW
2LeM7a2IOn30gBpzXkS2HAHiw 6Cler7GYgv-1rDajmK4RXaosliY692eNydDje5L9sYhE
uRWLaPBhwmchRJUPkUkSHhe3P2p2XeMwRKlhaWLur95HML-yAHndhF8PnOUT-aIesswSuS
IKabDJy-dDDXoD6vWVwB9v5kFg/file#) [following]
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oxusercontent.com/cd/0/inline/B3iC93bW2LeM7a2IOn30gBpzXkS2HAHiw 6Cler7
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RKlhaWLur95HML-yAHndhF8PnOUT-aIesswSuSIKabDJy-dDDXoD6vWVwB9v5kFg/file
(https://uc04907ead56e5d8bd17897f5bed.dl.dropboxusercontent.com/cd/0/i
nline/B3iC93bW2LeM7a2IOn30gBpzXkS2HAHiw 6Cler7GYgv-1rDajmK4RXaosliY692
eNydDje5L9sYhEuRWLaPBhwmchRJUPkUkSHhe3P2p2XeMwRKlhaWLur95HML-yAHndhF8P
nOUT-alesswSuSIKabDJy-dDDXoD6vWVwB9v5kFg/file)
Resolving uc04907ead56e5d8bd17897f5bed.dl.dropboxusercontent.com (uc04
907ead56e5d8bd17897f5bed.dl.dropboxusercontent.com)... 162.125.3.15, 2
620:100:6016:15::a27d:10f
Connecting to uc04907ead56e5d8bd17897f5bed.dl.dropboxusercontent.com
(uc04907ead56e5d8bd17897f5bed.dl.dropboxusercontent.com) | 162.125.3.15
:443... connected.
HTTP request sent, awaiting response... 302 Found
Location: /cd/0/inline2/B3hSRiIcx0wADdFSmDYAINdSJ7q0FwA0T3p-sj2XXU8eL6
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ljBocgMqM7-Q3U6RyUrmJnUXYYgeaY973UKpA4OH74euw72fttqxB3LJSmmJ5c4Z2bhcca
m0o8i6h3Wmsu54iersx265yCMKNbeK5Z0sVWxyN6O2RRm0AIfnUSVdaGVkG5ZA1cYWN-tk
6gZnDh3npTtLY3fCu5NVPibFCbpod2XJvRMfpipJaZIrPuR R4Jzp3AXoo819NDfnlIBl0
iUpU0DqXmM9vyj2MRaxCQ-jB040bOAgdZw/file [following]
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oxusercontent.com/cd/0/inline2/B3hSRiIcx0wADdFSmDYAINdSJ7q0FwA0T3p-sj2
XXU8eL6IzXFOpqMlX9hDFcn5EGKfifZFtwiTbUPUtZZ8HAAvHvEETEf-n9VfqmV06zVuuc
vH-M2Gf2LBVxAvtIlvlVgrd7Wei14i-qsJ1WgsUcrouF5BxkSDpgJnLIiLFoSlpLwZbn37
4XANn2S1jBocqMqM7-Q3U6RyUrmJnUXYYqeaY973UKpA4OH74euw72fttqxB3LJSmmJ5c4
Z2bhccam0o8i6h3Wmsu54iersx265yCMKNbeK5Z0sVWxyN6O2RRm0AIfnUSVdaGVkG5ZA1
cYWN-tk6qZnDh3npTtLY3fCu5NVPibFCbpod2XJvRMfpipJaZIrPuR R4Jzp3AXoo819ND
fnlIBl0iUpU0DqXmM9vyj2MRaxCQ-jB040bOAgdZw/file (https://uc04907ead56e5
d8bd17897f5bed.dl.dropboxusercontent.com/cd/0/inline2/B3hSRiIcx0wADdFS
mDYAINdSJ7q0FwA0T3p-sj2XXU8eL6IzXFOpgMlX9hDFcn5EGKfifZFtwiTbUPUtZZ8HAA
vHvEETEf-n9VfgmVO6zVuucvH-M2Gf2LBVxAvtIlvlVgrd7Wei14i-qsJ1WgsUcrouF5Bx
kSDpgJnLIiLFoSlpLwZbn374XANn2SljBocgMqM7-Q3U6RyUrmJnUXYYgeaY973UKpA4OH
74euw72fttqxB3LJSmmJ5c4Z2bhccam0o8i6h3Wmsu54iersx265yCMKNbeK5Z0sVWxyN6
O2RRmOAIfnUSVdaGVkG5ZA1cYWN-tk6qZnDh3npTtLY3fCu5NVPibFCbpod2XJvRMfpipJ
aZIrPuR_R4Jzp3AXoo819NDfnlIBl0iUpU0DqXmM9vyj2MRaxCQ-jB040b0AgdZw/file)
Reusing existing connection to uc04907ead56e5d8bd17897f5bed.dl.dropbox
```

```
usercontent.com:443.
HTTP request sent, awaiting response... 200 OK
Length: 9251149 (8.8M) [application/octet-stream]
Saving to: 'Task01_BrainTumour_2D.tar.gz'

Task01_BrainTumour_ 100%[=============]] 8.82M 21.8MB/s in 0.4s

2023-03-03 13:17:17 (21.8 MB/s) - 'Task01_BrainTumour_2D.tar.gz' saved [9251149/9251149]
```

# Visualise a random set of 4 training images along with their label maps.

Suggested colour map for brain MR image:

```
cmap = 'gray'
```

Suggested colour map for segmentation map:

```
cmap = colors.ListedColormap(['black', 'green', 'blue', 'red'])
```

#### In [3]:

```
# Define paths for training images and labels
train_image_path = "Task01_BrainTumour_2D/training_images"
train label path = "Task01 BrainTumour 2D/training labels"
image files = os.listdir(train image path)
label files = os.listdir(train label path)
# Define colour maps for image and label
image cmap = "gray"
label cmap = colors.ListedColormap(["black", "green", "blue", "red"])
plt.figure(figsize=(10, 10))
# Loop over 4 random indices
for i in range(4):
    index = random.randint(0, len(image files) - 1)
    image file = image files[index]
    label file = label files[index]
    # Read the images
    image = plt.imread(os.path.join(train image path, image file))
    label = plt.imread(os.path.join(train label path, label file))
    # Plot the images
    plt.subplot(4, 2, i * 2 + 1)
    plt.imshow(image, cmap=image cmap)
    plt.title(f"Image: {image_file}")
    plt.axis("off")
    plt.subplot(4, 2, i * 2 + 2)
    plt.imshow(label, cmap=label cmap)
    plt.title(f"Label: {label file}")
    plt.axis("off")
# Show the figure
plt.show()
```

Image: BRATS\_260\_z124.png

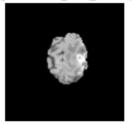


Image: BRATS\_355\_z108.png

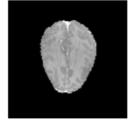


Image: BRATS 279 z93.png

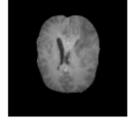
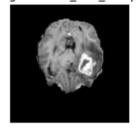
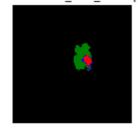


Image: BRATS\_431\_z62.png



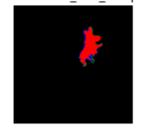
Label: BRATS 260 z124.png



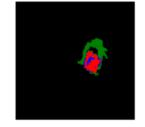
Label: BRATS\_355\_z108.png



Label: BRATS\_279\_z93.png



Label: BRATS\_431\_z62.png



## 2. Implement a dataset class.

It can read the imaging dataset and get items, pairs of images and label maps, as training batches.

In [4]:

```
def normalise intensity(image, thres roi=1.0):
    """ Normalise the image intensity by the mean and standard deviation """
    # ROI defines the image foreground
   val l = np.percentile(image, thres roi)
   roi = (image >= val 1)
   mu, sigma = np.mean(image[roi]), np.std(image[roi])
   eps = 1e-6
   image2 = (image - mu) / (sigma + eps)
   return image2
class BrainImageSet(Dataset):
    """ Brain image set """
   def init (self, image path, label path='', deploy=False):
       self.image path = image path
       self.deploy = deploy
       self.images = []
       self.labels = []
       image names = sorted(os.listdir(image path))
       for image name in image names:
           # Read the image
           image = imageio.imread(os.path.join(image path, image name))
           self.images += [image]
           # Read the label map
           if not self.deploy:
               label name = os.path.join(label path, image name)
               label = imageio.imread(label name)
               self.labels += [label]
   def len (self):
       return len(self.images)
   def getitem (self, idx):
       # Get an image and perform intensity normalisation
       # Dimension: XY
       image = normalise_intensity(self.images[idx])
       # Get its label map
       # Dimension: XY
       label = self.labels[idx]
       return image, label
   def get_random_batch(self, batch_size):
       # Get a batch of paired images and label maps
       # Dimension of images: NCXY
       # Dimension of labels: NXY
       images, labels = [], []
       # randomly select number of images
       indices = random.sample(range(len(self)), batch size)
       for idx in indices:
           # perform intensity normalisation
           # Dimension is normal: XY
           image = normalise intensity(self.images[idx])
```

#### In [5]:

```
Image shape: (120, 120), Label shape: (120, 120)
Images shape: (4, 1, 120, 120), Labels shape: (4, 120, 120)
```

## 3. Build a U-net architecture.

You will implement a U-net architecture. If you are not familiar with U-net, please read this paper:

[1] Olaf Ronneberger et al. <u>U-Net: Convolutional networks for biomedical image segmentation</u> (https://link.springer.com/chapter/10.1007/978-3-319-24574-4\_28). MICCAI, 2015.

For the first convolutional layer, you can start with 16 filters. We have implemented the encoder path. Please complete the decoder path.

#### In [6]:

```
""" U-net """
class UNet(nn.Module):
    def init (self, input channel=1, output channel=1, num filter=16):
        super(UNet, self). init ()
        # BatchNorm: by default during training this layer keeps running estimates
        # of its computed mean and variance, which are then used for normalization
        # during evaluation.
        # Encoder path
        n = num filter # 16
        self.conv1 = nn.Sequential(
            nn.Conv2d(input channel, n, kernel size=3, padding=1),
            nn.BatchNorm2d(n),
            nn.ReLU(),
            nn.Conv2d(n, n, kernel_size=3, padding=1),
            nn.BatchNorm2d(n),
            nn.ReLU()
        )
        n *= 2 # 32
        self.conv2 = nn.Sequential(
            nn.Conv2d(int(n / 2), n, kernel_size=3, stride=2, padding=1),
            nn.BatchNorm2d(n),
            nn.ReLU(),
            nn.Conv2d(n, n, kernel size=3, padding=1),
            nn.BatchNorm2d(n),
            nn.ReLU()
        )
        n *= 2 # 64
        self.conv3 = nn.Sequential(
            nn.Conv2d(int(n / 2), n, kernel_size=3, stride=2, padding=1),
            nn.BatchNorm2d(n),
            nn.ReLU(),
            nn.Conv2d(n, n, kernel size=3, padding=1),
            nn.BatchNorm2d(n),
            nn.ReLU()
        )
        n *= 2 # 128
        self.conv4 = nn.Sequential(
            nn.Conv2d(int(n / 2), n, kernel_size=3, stride=2, padding=1),
            nn.BatchNorm2d(n),
            nn.ReLU(),
            nn.Conv2d(n, n, kernel size=3, padding=1),
            nn.BatchNorm2d(n),
            nn.ReLU()
        )
        # Decoder path
        ### Insert your code
        # Bottom layer
        self.inter = nn.Sequential(
            nn.Conv2d(n, 2*n, kernel size=3, stride=2, padding=1),
            nn.BatchNorm2d(2*n),
            nn.ReLU(),
```

```
nn.ConvTranspose2d(2*n, n, kernel size=3, dilation = 3, output padding=1
        nn.BatchNorm2d(n),
        nn.ReLU()
    )
    self.conv up4 = nn.Sequential(
        nn.Conv2d(2*n, n, kernel size=3, padding=1),
        nn.BatchNorm2d(n),
        nn.ReLU(),
        nn.ConvTranspose2d(n, int(n/2), kernel size=3, stride=2, padding=1, outp
        nn.BatchNorm2d(int(n/2)),
        nn.ReLU()
    )
    n //= 2 # 64
    self.conv up3 = nn.Sequential(
        nn.Conv2d(2*n, n, kernel size=3, padding=1),
        nn.BatchNorm2d(n),
        nn.ReLU(),
        nn.ConvTranspose2d(n, int(n/2), kernel size=3, stride=2, padding=1, outp
        nn.BatchNorm2d(int(n/2)),
        nn.ReLU()
    )
    n //= 2 # 32
    self.conv up2 = nn.Sequential(
        nn.Conv2d(2*n, n, kernel size=3, padding=1),
        nn.BatchNorm2d(n),
        nn.ReLU(),
        nn.ConvTranspose2d(n, int(n/2), kernel size=3, stride=2, padding=1, out
        nn.BatchNorm2d(int(n/2)),
        nn.ReLU()
    )
    n //= 2 # 16
    self.conv up1 = nn.Sequential(
        nn.Conv2d(n*2, n, kernel size=3, padding=1),
        nn.BatchNorm2d(n),
        nn.ReLU(),
        nn.Conv2d(n, output channel, kernel size=1)
    )
             # 64
    # self.conv up4 = nn.Sequential(
         nn.ConvTranspose2d(n, int(n / 2), kernel size=3, stride=2, padding=1,
    #
    #
          nn.BatchNorm2d(int(n / 2)),
    #
          nn.ReLU(),
    #
          nn.Conv2d(int(n / 2), int(n / 2), kernel size=3, padding=1),
    #
          nn.BatchNorm2d(int(n / 2)),
    #
          nn.ReLU()
    # )
    ### End of your code
def forward(self, x):
    # Use the convolutional operators defined above to build the U-net
    # The encoder part is already done for you.
    # You need to complete the decoder part.
    # Encoder
    x = self.conv1(x)
```

```
conv1_skip = x
x = self.conv2(x)
conv2\_skip = x
x = self.conv3(x)
conv3\_skip = x
x = self.conv4(x)
conv4\_skip = x
x = self.inter(x)
# Decoder
### Insert your code
x = torch.cat([x, conv4_skip], 1)
x = self.conv_up4(x)
x = torch.cat([x, conv3_skip], 1)
x = self.conv up3(x)
x = torch.cat([x, conv2_skip], 1)
x = self.conv_up2(x)
x = torch.cat([x, conv1 skip], 1)
x = self.conv_up1(x)
### End of your code
return x
```

### In [7]:

```
# Test part 3
from torchsummary import summary

# CUDA device
device = torch.device('cuda' if torch.cuda.is_available() else 'cpu')
print('Device: {0}'.format(device))

# Build the model
num_class = 4
model = UNet(input_channel=1, output_channel=num_class, num_filter=16)
model = model.to(device)
params = list(model.parameters())

print(summary(model, (1, 120, 120)))
```

Device: cuda

Layer (type)	Output Shape	Param #
 Conv2d-1	[-1, 16, 120, 120]	160
BatchNorm2d-2	[-1, 16, 120, 120]	32
ReLU-3	[-1, 16, 120, 120]	0
Conv2d-4	[-1, 16, 120, 120]	2,320
BatchNorm2d-5	[-1, 16, 120, 120]	32
ReLU-6	[-1, 16, 120, 120]	0
Conv2d-7	[-1, 32, 60, 60]	4,640
BatchNorm2d-8	[-1, 32, 60, 60]	64
ReLU-9	[-1, 32, 60, 60]	0
Conv2d-10	[-1, 32, 60, 60]	9,248
BatchNorm2d-11	[-1, 32, 60, 60]	64
ReLU-12	[-1, 32, 60, 60]	0
Conv2d-13	[-1, 64, 30, 30]	18,496
BatchNorm2d-14	[-1, 64, 30, 30]	128
ReLU-15	[-1, 64, 30, 30]	0
Conv2d-16	[-1, 64, 30, 30]	36,928
BatchNorm2d-17	[-1, 64, 30, 30]	128
ReLU-18	[-1, 64, 30, 30]	0
Conv2d-19	[-1, 128, 15, 15]	73 <b>,</b> 856
BatchNorm2d-20	[-1, 128, 15, 15]	256
ReLU-21	[-1, 128, 15, 15]	0
Conv2d-22	[-1, 128, 15, 15]	147,584
BatchNorm2d-23	[-1, 128, 15, 15]	256
ReLU-24	[-1, 128, 15, 15]	0
Conv2d-25	[-1, 256, 8, 8]	295,168
BatchNorm2d-26	[-1, 256, 8, 8]	512
ReLU-27	[-1, 256, 8, 8]	0
ConvTranspose2d-28	[-1, 128, 15, 15]	295,040
BatchNorm2d-29	[-1, 128, 15, 15]	256
ReLU-30	[-1, 128, 15, 15]	0
Conv2d-31	[-1, 128, 15, 15]	295,040
BatchNorm2d-32	[-1, 128, 15, 15]	256
ReLU-33	[-1, 128, 15, 15]	0
ConvTranspose2d-34	[-1, 64, 30, 30]	73,792
BatchNorm2d-35	[-1, 64, 30, 30]	128
ReLU-36	[-1, 64, 30, 30]	0
Conv2d-37	[-1, 64, 30, 30]	73,792
BatchNorm2d-38	[-1, 64, 30, 30]	128
ReLU-39	[-1, 64, 30, 30]	0
ConvTranspose2d-40	[-1, 32, 60, 60]	18,464
BatchNorm2d-41	[-1, 32, 60, 60]	64
ReLU-42	[-1, 32, 60, 60]	0
Conv2d-43	[-1, 32, 60, 60]	18,464
BatchNorm2d-44	[-1, 32, 60, 60]	64
ReLU-45	[-1, 32, 60, 60]	0
ConvTranspose2d-46	[-1, 16, 120, 120]	4,624
BatchNorm2d-47	[-1, 16, 120, 120]	32
ReLU-48	[-1, 16, 120, 120]	0
Conv2d-49	[-1, 16, 120, 120]	4,624
BatchNorm2d-50	[-1, 16, 120, 120]	32
ReLU-51	[-1, 16, 120, 120]	0
Conv2d-52	[-1, 4, 120, 120]	68

Total params: 1,374,740

Trainable params: 1,374,740 Non-trainable params: 0

\_\_\_\_\_

Input size (MB): 0.05

Forward/backward pass size (MB): 40.37

Params size (MB): 5.24

Estimated Total Size (MB): 45.66

None

## 4. Train the segmentation model.

In [8]:

```
# CUDA device
device = torch.device('cuda' if torch.cuda.is available() else 'cpu')
print('Device: {0}'.format(device))
# Build the model
num class = 4
model = UNet(input channel=1, output channel=num class, num filter=16)
model = model.to(device)
params = list(model.parameters())
model dir = 'saved models'
if not os.path.exists(model dir):
        os.makedirs(model dir)
# Optimizer
optimizer = optim.Adam(params, lr=1e-3)
# Segmentation loss
criterion = nn.CrossEntropyLoss()
# Datasets
train set = BrainImageSet('Task01 BrainTumour 2D/training images', 'Task01 BrainTumour
test set = BrainImageSet('Task01 BrainTumour 2D/test images', 'Task01 BrainTumour 2D/
# Train the model
# Note: when you debug the model, you may reduce the number of iterations or batch s
# CHANGE THIS BACK TO BIGGGG
num iter = 10000
train batch size = 16
eval batch size = 16
start = time.time()
for it in range(1, 1 + num iter):
         # Set the modules in training mode, which will have effects on certain modules,
         start iter = time.time()
        model.train()
         # Get a batch of images and labels
         images, labels = train_set.get_random_batch(train_batch_size)
         images, labels = torch.from numpy(images), torch.from numpy(labels)
         images, labels = images.to(device, dtype=torch.float32), labels.to(device, dtype
         logits = model(images)
         # Perform optimisation and print out the training loss
         ### Insert your code ###
        loss = criterion(logits, labels)
        optimizer.zero_grad()
        loss.backward()
        optimizer.step()
        ### End of your code ###
         # Evaluate
         if it % 100 == 0:
                 model.eval()
                 # Disabling gradient calculation during reference to reduce memory consumpti
                 with torch.no grad():
                          # Evaluate on a batch of test images and print out the test loss
                          ### Insert your code ###
```

```
total_loss = 0
    images, labels = test_set.get_random_batch(eval_batch_size)
    images, labels = torch.from_numpy(images), torch.from_numpy(labels)
    images, labels = images.to(device, dtype=torch.float32), labels.to(device)
    logits = model(images)
    loss = criterion(logits, labels)
    total_loss += loss.item()
    avg_loss = total_loss / (len(test_set) / eval_batch_size)
    print('Iteration [{0}/{1}], test loss: {2:.4f}, loss: {3:.4f}'\
        .format(it, num_iter, avg_loss, loss))
    ### End of your code ###

# Save the model
if it % 5000 == 0:
    torch.save(model.state_dict(), os.path.join(model_dir, 'model_{0}.pt'.formatellost)
print('Training took {:.3f}s in total.'.format(time.time() - start))
```

```
Device: cuda
Iteration [100/10000], test loss: 0.0068, loss: 0.3119
Iteration [200/10000], test loss: 0.0027, loss: 0.1213
Iteration [300/10000], test loss: 0.0023, loss: 0.1061
Iteration [400/10000], test loss: 0.0016, loss: 0.0723
Iteration [500/10000], test loss: 0.0018, loss: 0.0831
Iteration [600/10000], test loss: 0.0014, loss: 0.0629
Iteration [700/10000], test loss: 0.0015, loss: 0.0692
Iteration [800/10000], test loss: 0.0015, loss: 0.0677
Iteration [900/10000], test loss: 0.0014, loss: 0.0646
Iteration [1000/10000], test loss: 0.0012, loss: 0.0567
Iteration [1100/10000], test loss: 0.0012, loss: 0.0558
Iteration [1200/10000], test loss: 0.0008, loss: 0.0359
Iteration [1300/10000], test loss: 0.0011, loss: 0.0524
Iteration [1400/10000], test loss: 0.0008, loss: 0.0351
Iteration [1500/10000], test loss: 0.0010, loss: 0.0463
Iteration [1600/10000], test loss: 0.0006, loss: 0.0256
Iteration [1700/10000], test loss: 0.0008, loss: 0.0349
Iteration [1800/10000], test loss: 0.0009, loss: 0.0422
Iteration [1900/10000], test loss: 0.0007, loss: 0.0327
Iteration [2000/10000], test loss: 0.0006, loss: 0.0290
Iteration [2100/10000], test loss: 0.0010, loss: 0.0457
Iteration [2200/10000], test loss: 0.0007, loss: 0.0340
Iteration [2300/10000], test loss: 0.0008, loss: 0.0362
Iteration [2400/10000], test loss: 0.0010, loss: 0.0444
Iteration [2500/10000], test loss: 0.0006, loss: 0.0292
Iteration [2600/10000], test loss: 0.0006, loss: 0.0260
Iteration [2700/10000], test loss: 0.0007, loss: 0.0335
Iteration [2800/10000], test loss: 0.0009, loss: 0.0408
Iteration [2900/10000], test loss: 0.0005, loss: 0.0234
Iteration [3000/10000], test loss: 0.0009, loss: 0.0412
Iteration [3100/10000], test loss: 0.0006, loss: 0.0253
Iteration [3200/10000], test loss: 0.0008, loss: 0.0355
Iteration [3300/10000], test loss: 0.0008, loss: 0.0377
Iteration [3400/10000], test loss: 0.0006, loss: 0.0266
Iteration [3500/10000], test loss: 0.0011, loss: 0.0487
Iteration [3600/10000], test loss: 0.0008, loss: 0.0384
Iteration [3700/10000], test loss: 0.0007, loss: 0.0304
Iteration [3800/10000], test loss: 0.0007, loss: 0.0323
Iteration [3900/10000], test loss: 0.0007, loss: 0.0315
Iteration [4000/10000], test loss: 0.0005, loss: 0.0206
Iteration [4100/10000], test loss: 0.0006, loss: 0.0296
Iteration [4200/10000], test loss: 0.0005, loss: 0.0224
Iteration [4300/10000], test loss: 0.0012, loss: 0.0565
Iteration [4400/10000], test loss: 0.0006, loss: 0.0268
Iteration [4500/10000], test loss: 0.0008, loss: 0.0358
Iteration [4600/10000], test loss: 0.0007, loss: 0.0319
Iteration [4700/10000], test loss: 0.0007, loss: 0.0303
Iteration [4800/10000], test loss: 0.0007, loss: 0.0322
Iteration [4900/10000], test loss: 0.0009, loss: 0.0395
Iteration [5000/10000], test loss: 0.0012, loss: 0.0544
Iteration [5100/10000], test loss: 0.0007, loss: 0.0302
Iteration [5200/10000], test loss: 0.0005, loss: 0.0215
Iteration [5300/10000], test loss: 0.0009, loss: 0.0422
Iteration [5400/10000], test loss: 0.0009, loss: 0.0393
Iteration [5500/10000], test loss: 0.0011, loss: 0.0510
Iteration [5600/10000], test loss: 0.0006, loss: 0.0292
Iteration [5700/10000], test loss: 0.0008, loss: 0.0377
Iteration [5800/10000], test loss: 0.0005, loss: 0.0213
Iteration [5900/10000], test loss: 0.0005, loss: 0.0210
Iteration [6000/10000], test loss: 0.0007, loss: 0.0341
```

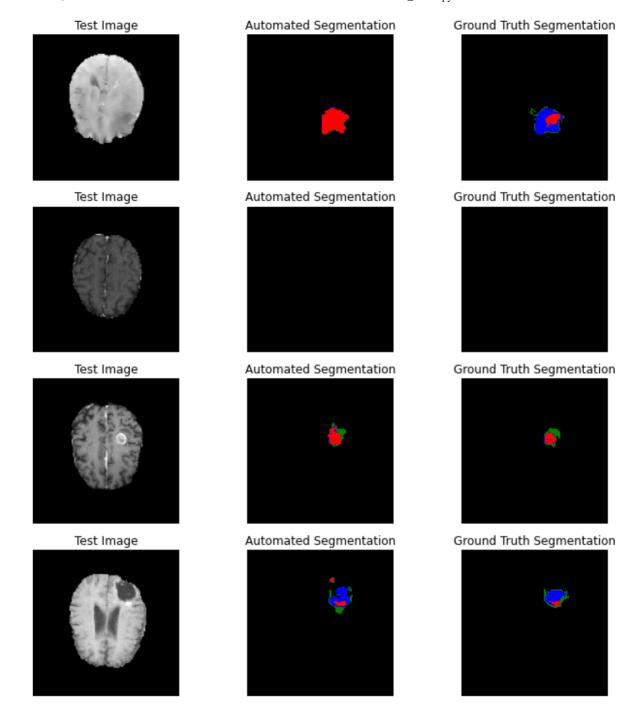
```
Iteration [6100/10000], test loss: 0.0007, loss: 0.0336
Iteration [6200/10000], test loss: 0.0008, loss: 0.0353
Iteration [6300/10000], test loss: 0.0011, loss: 0.0519
Iteration [6400/10000], test loss: 0.0009, loss: 0.0425
Iteration [6500/10000], test loss: 0.0013, loss: 0.0571
Iteration [6600/10000], test loss: 0.0008, loss: 0.0352
Iteration [6700/10000], test loss: 0.0005, loss: 0.0231
Iteration [6800/10000], test loss: 0.0011, loss: 0.0502
Iteration [6900/10000], test loss: 0.0013, loss: 0.0596
Iteration [7000/10000], test loss: 0.0010, loss: 0.0440
Iteration [7100/10000], test loss: 0.0008, loss: 0.0342
Iteration [7200/10000], test loss: 0.0009, loss: 0.0418
Iteration [7300/10000], test loss: 0.0013, loss: 0.0612
Iteration [7400/10000], test loss: 0.0004, loss: 0.0177
Iteration [7500/10000], test loss: 0.0009, loss: 0.0395
Iteration [7600/10000], test loss: 0.0010, loss: 0.0454
Iteration [7700/10000], test loss: 0.0011, loss: 0.0487
Iteration [7800/10000], test loss: 0.0014, loss: 0.0661
Iteration [7900/10000], test loss: 0.0009, loss: 0.0430
Iteration [8000/10000], test loss: 0.0010, loss: 0.0468
Iteration [8100/10000], test loss: 0.0013, loss: 0.0578
Iteration [8200/10000], test loss: 0.0010, loss: 0.0454
Iteration [8300/10000], test loss: 0.0009, loss: 0.0425
Iteration [8400/10000], test loss: 0.0008, loss: 0.0380
Iteration [8500/10000], test loss: 0.0009, loss: 0.0426
Iteration [8600/10000], test loss: 0.0006, loss: 0.0273
Iteration [8700/10000], test loss: 0.0010, loss: 0.0435
Iteration [8800/10000], test loss: 0.0009, loss: 0.0405
Iteration [8900/10000], test loss: 0.0008, loss: 0.0377
Iteration [9000/10000], test loss: 0.0005, loss: 0.0232
Iteration [9100/10000], test loss: 0.0013, loss: 0.0586
Iteration [9200/10000], test loss: 0.0020, loss: 0.0903
Iteration [9300/10000], test loss: 0.0009, loss: 0.0391
Iteration [9400/10000], test loss: 0.0010, loss: 0.0464
Iteration [9500/10000], test loss: 0.0008, loss: 0.0387
Iteration [9600/10000], test loss: 0.0005, loss: 0.0242
Iteration [9700/10000], test loss: 0.0009, loss: 0.0394
Iteration [9800/10000], test loss: 0.0005, loss: 0.0223
Iteration [9900/10000], test loss: 0.0016, loss: 0.0712
Iteration [10000/10000], test loss: 0.0019, loss: 0.0889
Training took 359.858s in total.
```

# 5. Deploy the trained model to a random set of 4 test images and visualise the automated segmentation.

You can show the images as a 4 x 3 panel. Each row shows one example, with the 3 columns being the test image, automated segmentation and ground truth segmentation.

#### In [23]:

```
### Insert your code ###
label cmap = colors.ListedColormap(["black", "green", "blue", "red"])
# get test images and labels
test images, test labels = test set.get random batch(4)
# use evaluation mode
model.eval()
# get the automated segmentation for each test image using model
test images = torch.from numpy(test images).to(device, dtype=torch.float32)
with torch.no grad():
    test logits = model(test images)
test segmentations = torch.argmax(test logits, dim=1).cpu().numpy()
num test images = 4
fig, axs = plt.subplots(nrows=num test images, ncols=3, figsize=(10, 10))
for i in range(num test images):
    # test image
    axs[i, 0].imshow(test images[i, 0, :, :].cpu(), cmap='gray')
    axs[i, 0].axis('off')
    axs[i, 0].set_title('Test Image')
    # automated segmentation
    axs[i, 1].imshow(test segmentations[i, :, :], cmap=label cmap)
    axs[i, 1].axis('off')
    axs[i, 1].set title('Automated Segmentation')
    # ground truth segmentation
    axs[i, 2].imshow(test labels[i, :, :], cmap=label cmap)
    axs[i, 2].axis('off')
    axs[i, 2].set title('Ground Truth Segmentation')
plt.tight layout()
plt.show()
### End of your code ###
```



# 6. Discussion. Does your trained model work well? How would you improve this model so it can be deployed to the real clinic?

To evaluate the performance of the trained model, various evaluation metrics can be computed, such as sensitivity, specificity, precision, and others. These metrics are crucial in determining the model's efficacy, as they directly impact patient outcomes. A model with high sensitivity can accurately detect a larger proportion of true positive cases, such as tumors, ensuring timely and appropriate treatment. Similarly, a model with high specificity can minimize false positive cases, avoiding unnecessary interventions or misdiagnosis.

One important technique that can be employed to improve the model's performance is early stopping. This technique can help prevent overfitting and ensure that the model is generalizing well to new data. By implementing early stopping, the model can be trained for the optimal number of epochs and achieve the best performance.

To further improve the model's performance, various techniques can be employed. Data augmentation techniques, such as flipping, rotating, and scaling, can increase the size of the training dataset and improve the model's generalization ability. Hyperparameters, such as learning rate, batch size, and number of filters, can be optimized through tuning. Transfer learning can be used to initialize the model with pre-trained weights from a similar dataset or task, improving convergence and performance. Ensemble methods can combine the predictions of multiple models trained on different subsets of data to enhance the model's robustness and performance.

Ultimately, to ensure the model's effectiveness in clinical settings, it should be validated on a diverse set of clinical data. Clinical validation is essential to verify that the model is reliable and robust in different clinical contexts, leading to better patient outcomes.