1 !pip install ipython-autotime

```
2 !apt-get install openslide-tools
3 !pip install openslide-python
4 %load ext autotime
    Downloading https://files.pythonhosted.org/packages/b4/c9/b413a24f759641bc27ef
  Requirement already satisfied: ipython in /usr/local/lib/python3.7/dist-packages
  Requirement already satisfied: traitlets>=4.2 in /usr/local/lib/python3.7/dist-p
  Requirement already satisfied: setuptools>=18.5 in /usr/local/lib/python3.7/dist
  Requirement already satisfied: pickleshare in /usr/local/lib/python3.7/dist-pack
  Requirement already satisfied: pexpect; sys_platform != "win32" in /usr/local/li
  Requirement already satisfied: prompt-toolkit<2.0.0,>=1.0.4 in /usr/local/lib/py
  Requirement already satisfied: decorator in /usr/local/lib/python3.7/dist-packag
  Requirement already satisfied: pygments in /usr/local/lib/python3.7/dist-package
  Requirement already satisfied: simplegeneric>0.8 in /usr/local/lib/python3.7/dis
  Requirement already satisfied: ipython-genutils in /usr/local/lib/python3.7/dist
  Requirement already satisfied: ptyprocess>=0.5 in /usr/local/lib/python3.7/dist-
  Requirement already satisfied: wcwidth in /usr/local/lib/python3.7/dist-packages
  Requirement already satisfied: six>=1.9.0 in /usr/local/lib/python3.7/dist-packa
   Installing collected packages: ipython-autotime
  Successfully installed ipython-autotime-0.3.1
  Reading package lists... Done
  Building dependency tree
  Reading state information... Done
  The following package was automatically installed and is no longer required:
    libnvidia-common-460
  Use 'apt autoremove' to remove it.
  The following additional packages will be installed:
     libopenslide0
  Suggested packages:
    libtiff-tools
  The following NEW packages will be installed:
    libopenslide0 openslide-tools
   0 upgraded, 2 newly installed, 0 to remove and 34 not upgraded.
  Need to get 92.5 kB of archives.
  After this operation, 268 kB of additional disk space will be used.
  Get:1 http://archive.ubuntu.com/ubuntu bionic/universe amd64 libopenslide0 amd64
  Get: 2 http://archive.ubuntu.com/ubuntu bionic/universe amd64 openslide-tools amd
  Fetched 92.5 kB in 1s (149 kB/s)
  Selecting previously unselected package libopenslide0.
   (Reading database ... 160690 files and directories currently installed.)
  Preparing to unpack .../libopenslide0 3.4.1+dfsg-2 amd64.deb ...
  Unpacking libopenslide0 (3.4.1+dfsg-2) ...
  Selecting previously unselected package openslide-tools.
  Preparing to unpack .../openslide-tools 3.4.1+dfsg-2 amd64.deb ...
  Unpacking openslide-tools (3.4.1+dfsq-2) ...
  Setting up libopenslide0 (3.4.1+dfsg-2) ...
  Setting up openslide-tools (3.4.1+dfsg-2) ...
  Processing triggers for libc-bin (2.27-3ubuntu1.2) ...
   /sbin/ldconfig.real: /usr/local/lib/python3.7/dist-packages/ideep4py/lib/libmkld
  Processing triggers for man-db (2.8.3-2ubuntu0.1) ...
  Collecting openslide-python
    Downloading https://files.pythonhosted.org/packages/03/da/12dc0e7566ace61a5a65
```

317kB 6.6MB/s

```
Requirement already satisfied: Pillow in /usr/local/lib/python3.7/dist-packages
   Building wheels for collected packages: openslide-python
     Building wheel for openslide-python (setup.py) ... done
     Created wheel for openslide-python: filename=openslide python-1.1.2-cp37-cp37m
     Stored in directory: /root/.cache/pip/wheels/6b/55/74/ba9d3dcc2c5c0f1282e08bae
   Successfully built openslide-python
   Installing collected packages: openslide-python
   Successfully installed openslide-python-1.1.2
   time: 3.5 ms (started: 2021-04-29 14:38:17 +00:00)
 1 %matplotlib inline
 2 import matplotlib.pyplot as plt
 3 import matplotlib.image as mpimg
 4 from openslide import open slide, library version as openslide vers:
 5 import seaborn as sns; sns.set theme()
 6 from sklearn.model selection import train test split
 7 import numpy as np
 8 import pickle
 9 import os
10 from PIL import Image
   time: 866 ms (started: 2021-04-29 14:38:22 +00:00)
 1 import tensorflow as tf
 2 from tensorflow import keras
 3 from tensorflow.keras import layers, models
 4 from tensorflow.keras.layers import Conv2D, MaxPooling2D, Flatten
 5 from tensorflow.keras.layers import Input, LSTM, Embedding, Dense, Glob;
 6 from tensorflow.keras.models import Model, Sequential
 7 tf. version
    '2.4.1' time: 14.6 ms (started: 2021-04-29 14:49:04 +00:00)
 1 # mounting the drive
 2 from google.colab import drive
 3 drive.mount('/content/drive/')
   Mounted at /content/drive/
   time: 15.1 s (started: 2021-04-29 14:38:33 +00:00)
```

# Inception Model with Data Augmentation

```
1 PATCH_SIZE = 75 # size of the patches
2 LEVEL = 4
3 LEVEL_1 = 4 # first zoom level
4 LEVEL 2 = 3 # second zoom level
```

```
5 dataset_path = "/content/drive/MyDrive/Columbia_Assignments/ADL/Project/
6
7 train_file_normal = os.path.join(dataset_path, 'train/camelyon_preproces
8 train_file_zoomed = os.path.join(dataset_path, 'train/camelyon_preproces
9
10 test_file_normal = os.path.join(dataset_path, 'test/camelyon_preprocesse
11 test_file_zoomed = os.path.join(dataset_path, 'test/camelyon_preprocesse
time: 9.71 ms (started: 2021-04-29 14:43:11 +00:00)
```

#### Load train and test dataset

Load the data from the pickle file uploaded to the drive.

```
1 def load dataset(file path):
     with open(file path, 'rb') as f:
          slides, labels = pickle.load(f)
3
     return slides, labels
4
  time: 2.54 ms (started: 2021-04-29 14:40:29 +00:00)
1 # load the training dataset and labels
3 train slides, train labels = load dataset(train file normal)
4 train slides = np.array(train slides)
5 train labels = np.array(train labels)
6 print("Training slides length: {}".format(len(train_slides)))
7 print("Train labels: {}".format(train labels))
8 print("Number of cancerous labels: {}".format(np.sum(train labels)))
  Training slides length: 30065
  Train labels: [0 0 0 ... 0 0 0]
  Number of cancerous labels: 2016
  time: 9.98 s (started: 2021-04-29 17:27:51 +00:00)
1 # load the zoomed training dataset and labels
2
3 train zoomed slides, train labels = load dataset(train file zoomed)
4 train zoomed slides = np.array(train zoomed slides)
5 train labels = np.array(train labels)
6 print("Training slides length: {}".format(len(train zoomed slides)))
7 print("Train labels: {}".format(train labels))
8 print("Number of cancerous labels: {}".format(np.sum(train labels)))
  Training slides length: 30065
  Train labels: [0 0 0 ... 0 0 0]
```

```
Number of cancerous labels: 2016
  time: 3.47 s (started: 2021-04-29 16:36:10 +00:00)
1 # load the test dataset and labels
2
3 test slides, test labels = load dataset(test file normal)
4 test slides = np.array(test slides)
5 test labels = np.array(test labels)
6 print("Test slides length: {}".format(len(test slides)))
7 print("Test labels: {}".format(test labels))
8 print("Number of cancerous labels: {}".format(np.sum(test labels)))
  Test slides length: 11446
  Test labels: [0 0 0 ... 0 0 0]
  Number of cancerous labels: 1149
  time: 4.63 s (started: 2021-04-29 14:43:15 +00:00)
1 # load the zoomed test dataset and labels
3 test zoomed slides, test labels = load dataset(test file zoomed)
4 test zoomed slides = np.array(test zoomed slides)
5 test labels = np.array(test labels)
6 print("Test slides length: {}".format(len(test zoomed slides)))
7 print("Test labels: {}".format(test labels))
8 print("Number of cancerous labels: {}".format(np.sum(test labels)))
  Test slides length: 11446
  Test labels: [0 0 0 ... 0 0 0]
  Number of cancerous labels: 1149
  time: 2.72 s (started: 2021-04-29 16:40:16 +00:00)
```

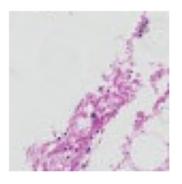
## Split the training dataset into validation

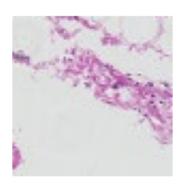
```
1:rain_slides, val_slides, train_zoomed_slides, val_zoomed_slides, train_
2:rint("Validation slides length: {}".format(len(val_slides)))
3:rint("Validation zoomed slides length: {}".format(len(val_zoomed_slides 4:rint("Validation labels: {}".format(val_labels))
5:rint("Number of cancerous labels: {}".format(np.sum(val_labels)))

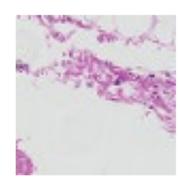
Validation slides length: 7517
   Validation zoomed slides length: 7517
   Validation labels: [1 0 0 ... 0 0 0]
   Number of cancerous labels: 494
   time: 1.02 s (started: 2021-04-29 14:46:23 +00:00)
```

- Define the model
- Data Augmentation

```
1 # adding a data augmentation layer to the model
2 data augmentation = keras.Sequential(
3
             Γ
4
                     layers.experimental.preprocessing.Rescaling(scale=1./255, input share
5
                     layers.experimental.preprocessing.RandomFlip("horizontal and vertical 
                     layers.experimental.preprocessing.RandomRotation(0.2),
6
7
                     # layers.experimental.preprocessing.RandomZoom(0.2)
8
             1
9)
         time: 148 ms (started: 2021-04-29 17:22:52 +00:00)
1 # displaying some patches with data augmentation
2 plt.figure(figsize=(10, 10))
3 image = train slides[19]
4 for i in range(9):
                     image expand = tf.expand dims(image, axis=0)
                     augmented images = data augmentation(image expand)
6
                     ax = plt.subplot(3, 3, i + 1)
7
                     plt.imshow(augmented images[0])
                    plt.axis("off")
9
```







### Two Inception Models

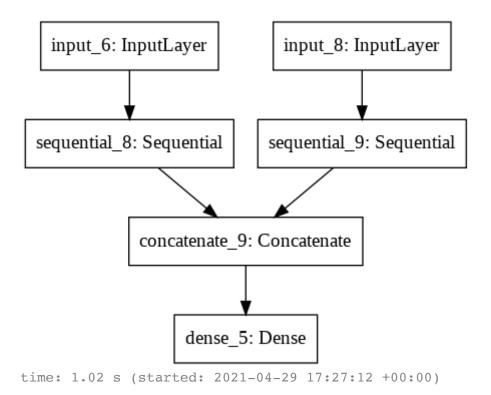
```
1 # inception model 1 for level = LEVEL 1 images
 2 base model 1 = tf.keras.applications.InceptionV3(input shape=[PATCH SIZE
 3 base model 1.trainable = False
 5 # global average layer
 6 global average layer = GlobalAveragePooling2D()
 7 dense = Dense(16, activation='relu')
 9 model 1 = Sequential()
10 model 1.add(data augmentation)
11 model 1.add(base model 1)
12 model 1.add(global_average_layer)
13 model 1.add(dense)
14
15 level 1 input = Input(shape=(PATCH SIZE, PATCH SIZE, 3))
16 level 1 image = model 1(level 1 input)
   time: 6.89 s (started: 2021-04-29 17:26:18 +00:00)
 1 # inception model 1 for level = LEVEL 1 images
 2 base model 2 = tf.keras.applications.InceptionV3(input shape=[PATCH SIZ]
 3 base model 2.trainable = False
 4
 5 # global average layer
 6 global average layer = GlobalAveragePooling2D()
 7 dense = Dense(16, activation='relu')
 8
 9 model 2 = Sequential()
10 model 2.add(data augmentation)
11 model 2.add(base model 2)
12 model 2.add(global average layer)
13 model 2.add(dense)
14
15 level 2 input = Input(shape=(PATCH SIZE, PATCH SIZE, 3))
```

```
coms_4995_project_training_inception_augmentation.ipynb - Colaboratory
16 Level 2 image = model 2(level 2 input)
   time: 5.26 s (started: 2021-04-29 17:26:29 +00:00)
 1 # concatenating the outputs from the two models
 2 concat layer = keras.layers.concatenate([level 1 image, level 2 image])
 3 output = Dense(1, activation='sigmoid')(concat layer)
 5 # final model
 6 model = Model(inputs=[level 1 input, level 2 input], outputs=output)
   time: 34.1 ms (started: 2021-04-29 17:26:36 +00:00)
 1 model.compile(optimizer='adam',
 2
                  loss='binary crossentropy',
 3
                  metrics=['accuracy'])
   time: 28.9 ms (started: 2021-04-29 17:26:39 +00:00)
 1 model.summary()
   Model: "model 1"
```

Layer (type)	Output Shape	Param #	Connected to
input_6 (InputLayer)	[(None, 75, 75, 3)]	0	
input_8 (InputLayer)	[(None, 75, 75, 3)]	0	
sequential_8 (Sequential)	(None, 16)	21835568	input_6[0][0]
sequential_9 (Sequential)	(None, 16)	21835568	input_8[0][0]
concatenate_9 (Concatenate)	(None, 32)	0	sequential_8[0] sequential_9[0]
dense_5 (Dense)	(None, 1)	33	concatenate_9[0
Total params: 43,671,169 Trainable params: 65,601 Non-trainable params: 43,605,5	568		
time: 66.4 ms (started: 2021-0	04-29 17:26:40 +00:00)		

## Plot the model for visualization

```
1 from tensorflow.keras.utils import plot model
2
3 plot model(model)
```



## Training the model

```
1 \text{ EPOCHS} = 10
2 callbacks = [
  keras.callbacks.ModelCheckpoint("save at {epoch}.h5"),
4 ]
5 history = model.fit(
   [train slides, train zoomed slides],
7
   train labels,
  validation data=([val slides, val zoomed slides], val labels),
8
9
  callbacks=callbacks,
  epochs=EPOCHS
10
11)
 ========= | - 659s 702ms/step - loss: 0.1363 - accuracy: 0.9563
```

## Plot the graph for accuracy and loss

```
1 # Your code here
 2 acc = history.history['accuracy']
 3 val acc = history.history['val accuracy']
 4 loss = history.history['loss']
 5 val loss = history.history['val loss']
 7 # Get the number of epochs
 8 epochs = range(len(acc))
10 plt.title('Training and validation accuracy')
11 plt.plot(epochs, acc, color='blue', label='Train')
12 plt.plot(epochs, val acc, color='orange', label='Val')
13 plt.xlabel('Epoch')
14 plt.ylabel('Accuracy')
15 plt.legend()
16
17 = plt.figure()
18 plt.title('Training and validation loss')
19 plt.plot(epochs, loss, color='blue', label='Train')
20 plt.plot(epochs, val loss, color='orange', label='Val')
21 plt.xlabel('Epoch')
22 plt.ylabel('Loss')
23 plt.legend()
```

<matplotlib.legend.Legend at 0x7fe755b6d450>





### Test the model on the test data

```
1 # loading the test tif files to be used for plotting
 2 test tumor path = os.path.join(dataset path, 'test/tumor')
 3 test tumor mask path = os.path.join(dataset path, 'test/tumor mask')
 4
 5 test tumors tifs = []
 6 test tumors mask tifs = []
 7
 8 for filename in os.listdir(test tumor path):
      test tumors tifs.append(os.path.join(test tumor path, filename))
10 for filename in os.listdir(test tumor mask path):
      test tumors mask tifs.append(os.path.join(test tumor mask path, file
11
12
13 test tumors tifs.sort()
14 test tumors mask tifs.sort()
15 print("Length of test tumor tiffs: {}".format(len(test tumors tifs)))
16 print("Length of test tumor mask tiffs: {}".format(len(test tumors mask
   Length of test tumor tiffs: 2
   Length of test tumor mask tiffs: 2
   time: 23 ms (started: 2021-04-29 19:38:20 +00:00)
```

```
1 # create separate test slides and test labels for each of the two test +
 2
 3 # this corresponds to the number of patches in the row and column of each
 4 # tif file.
 5 stride width array = [116, 78]
 6 stride height array = [59, 59]
 7
8 first index = stride height array[0]*stride width array[0]
9 test slides normal 1 = test slides[:first index]
10 test labels normal 1 = test labels[:first index]
11
12 test slides normal 2 = test slides[first index:]
13 test labels normal 2 = test labels[first index:]
14
15 print("first tiff index: ", first index)
16 print("length: ", test labels normal 1)
17 print(np.sum(test labels normal 1))
18
19 print("length: ", test_labels_normal_2)
20 print(np.sum(test labels normal 2))
   first tiff index: 6844
   length: [0 0 0 ... 0 0 0]
   222
   length: [0 0 0 ... 0 0 0]
   927
   time: 19.5 ms (started: 2021-04-29 19:38:26 +00:00)
 1 # create separate test slides and test labels for each of the two test t
 2 # stride width array zoom = [232, 157]
 3 # stride height array zoom = [119, 119]
 4
 5 # first tiff index = PATCH SIZE * stride width array[0] + PATCH SIZE * s
 6 # first index = stride height array zoom[0]*stride width array zoom[0]
 7 test slides zoomed 1 = test zoomed slides[:first index]
 8 test labels zoomed 1 = test labels[:first index]
 9
10 test slides zoomed 2 = test zoomed slides[first index:]
11 test labels zoomed 2 = test labels[first index:]
12
13 print("first tiff index: ", first index)
14 print("length: ", test labels zoomed 1)
15 print(np.sum(test labels zoomed 1))
16
17 print("length: ", test labels zoomed 2)
18 print(np.sum(test_labels_zoomed_2))
```

```
first_tiff_index: 6844
length: [0 0 0 ... 0 0 0]
222
length: [0 0 0 ... 0 0 0]
927
time: 16.5 ms (started: 2021-04-29 19:38:30 +00:00)
```

## Predicting the first test tiff file.

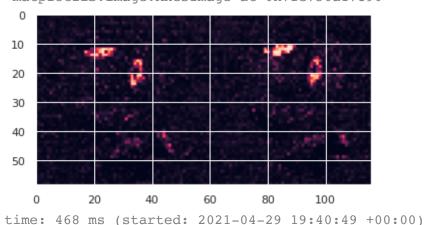
```
1 predictions = model.predict([test_slides_normal_1, test_slides_zoomed_1
    time: lmin 54s (started: 2021-04-29 19:38:33 +00:00)

1 print("length of predictions: {}".format(len(predictions)))
2 print("predictions: {}".format(predictions))

length of predictions: 6844
    predictions: [[0.02210265]
    [0.01815641]
    [0.04221612]
    ...
    [0.01344201]
    [0.01127237]
    [0.00645652]]
    time: 15.6 ms (started: 2021-04-29 19:40:46 +00:00)
```

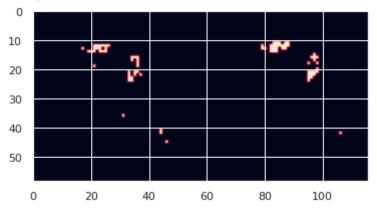
#### 1 plt.imshow(np.reshape(predictions, (116, 59)).T)

<matplotlib.image.AxesImage at 0x7fe756a17c90>



```
1 # plotting the thresholded
2 THRESHOLD = 0.5
3 threshold_predictions = np.where(predictions > THRESHOLD, 1, 0)
4 plt.imshow(np.reshape(threshold predictions, (116, 59)).T)
```





time: 438 ms (started: 2021-04-29 19:41:03 +00:00)

### ▼ F1-Score

```
1 from sklearn.metrics import classification_report
2
3 classification_metrics = classification_report(threshold_predictions, to 4 print(classification metrics)
```

	precision	recall	f1-score	support
0	1.00 0.34	0.98 0.94	0.99	6763 81
accuracy macro avg	0.67	0.96	0.98 0.75	6844 6844
weighted avg	0.99	0.98	0.98	6844

time: 25.3 ms (started: 2021-04-29 20:02:20 +00:00)

#### Intersection over Union Score

```
1 y_pred_image = np.reshape(threshold_predictions, (116, 59))
2 y_true_image = np.reshape(test_labels_normal_1, (116, 59))
3
4 inter = np.logical_and(y_pred_image, y_true_image)
5 union = np.logical_or(y_pred_image, y_true_image)
6
7 print(np.sum(inter)/float(np.sum(union)))
0.33480176211453744
time: 6.04 ms (started: 2021-04-29 20:08:58 +00:00)
```

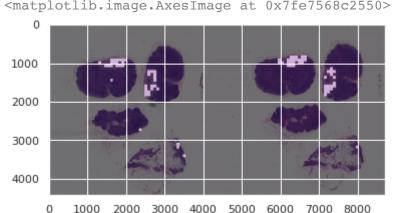
## Heatmap For The test tiff

```
1 # read slide and return an image
 2 def read slide(slide, x, y, level, width, height, as float=False):
      im = slide.read region((x,y), level, (width, height))
 3
      im = im.convert('RGB') # drop the alpha channel
 4
 5
      if as float:
 6
           im = np.asarray(im, dtype=np.float32)
 7
      else:
 8
           im = np.asarray(im)
 9
      assert im.shape == (height, width, 3)
10
      return im
   time: 7.23 ms (started: 2021-04-29 19:41:10 +00:00)
 1 # show a slide
 2 tumor image = open slide(test tumors tifs[0])
 3 mask image = open slide(test tumors mask tifs[0])
 4
 5 width = tumor image.level dimensions[LEVEL 1][0]
 6 height = tumor image.level dimensions[LEVEL 1][1]
 8 tumor slide = read slide(tumor_image, 0, 0, LEVEL_1, width=width, height
9 mask_slide = read_slide(mask image, 0, 0, LEVEL 1, width=width, height=1
10
11 plt.imshow(tumor slide)
12 plt.imshow(mask slide[:, :, 0], cmap='Reds', alpha=0.7)
13 plt.show()
      0
    1000
    2000
```

```
3000
4000
        1000 2000 3000 4000 5000 6000 7000 8000
```

time: 9.54 s (started: 2021-04-29 19:41:12 +00:00)

```
1 # plot the patches to see if patch extraction is working properly
     2 step width = width // PATCH SIZE
     3 step height = height // PATCH SIZE
     4 canvas_slide = Image.new('RGB', (PATCH_SIZE * step_width, PATCH_SIZE * s
     5 canvas mask = Image.new('RGB', (PATCH SIZE * step width, PATCH SIZE * st
     6 mask blank = np.zeros((PATCH SIZE, PATCH SIZE))
     7 mask tumor = 255 * np.ones((PATCH SIZE, PATCH SIZE))
https://colab.research.google.com/drive/1LUO9zbJ02u19esZJBjNocJTYa1INOnHq#scrollTo=lfeMvVqEijie&printMode=true
                                                                                      14/19
```



time: 30.9 s (started: 2021-04-29 19:41:45 +00:00)

# Predicting the second tiff file

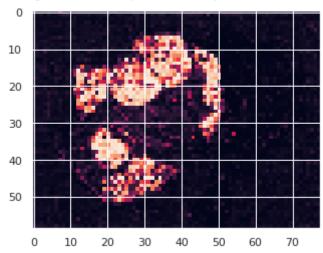
```
1 predictions_2 = model.predict([test_slides_normal_2, test_slides_zoomed_
    time: lmin 15s (started: 2021-04-29 19:42:45 +00:00)

1 print("length of predictions: {}".format(len(predictions_2)))
2 print("predictions: {}".format(predictions_2))
    length of predictions: 4602
    predictions: [[0.04947379]]
```

```
[0.03780037]
[0.02027133]
...
[0.02750137]
[0.04804933]
[0.02972835]]
time: 15.6 ms (started: 2021-04-29 19:44:13 +00:00)
```

#### 1 plt.imshow(np.reshape(predictions 2, (78, 59)).T)

<matplotlib.image.AxesImage at 0x7fe7567d1190>

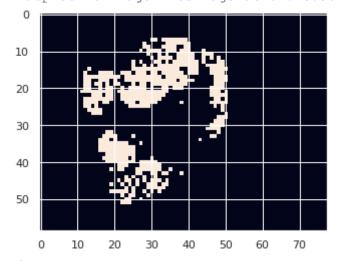


time: 436 ms (started: 2021-04-29 19:44:15 +00:00)

```
1 \text{ THRESHOLD} = 0.5
```

- 2 threshold predictions = np.where(predictions 2 > THRESHOLD, 1, 0)
- 3 plt.imshow(np.reshape(threshold\_predictions, (78, 59)).T)

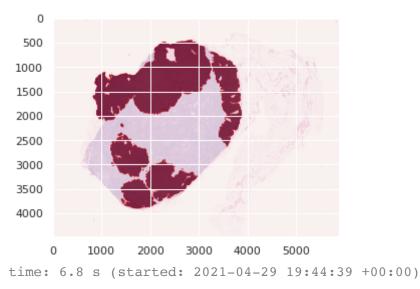
<matplotlib.image.AxesImage at 0x7fe7567ff690>



time: 526 ms (started: 2021-04-29 19:44:29 +00:00)

# Heatmap For The test tiff

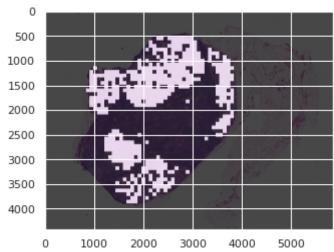
```
1 # read slide and return an image
 2 def read slide(slide, x, y, level, width, height, as float=False):
      im = slide.read_region((x,y), level, (width, height))
 3
      im = im.convert('RGB') # drop the alpha channel
 4
      if as float:
 5
 6
           im = np.asarray(im, dtype=np.float32)
 7
      else:
 8
           im = np.asarray(im)
 9
      assert im.shape == (height, width, 3)
      return im
10
   time: 15.3 ms (started: 2021-04-29 19:44:37 +00:00)
 1 # show a slide
 2 tumor image = open slide(test tumors tifs[1])
 3 mask image = open slide(test tumors mask tifs[1])
 5 width = tumor image.level dimensions[LEVEL 1][0]
 6 height = tumor image.level dimensions[LEVEL 1][1]
 7
 8 tumor slide = read slide(tumor image, 0, 0, LEVEL 1, width=width, height
 9 mask slide = read slide(mask image, 0, 0, LEVEL 1, width=width, height=1
10
11 plt.imshow(tumor slide)
12 plt.imshow(mask slide[:, :, 0], cmap='Reds', alpha=0.7)
13 plt.show()
```



```
1 plot the patches to see if patch extraction is working properly
2 tep width = width // PATCH SIZE
3 tep height = height // PATCH SIZE
4 anvas_slide = Image.new('RGB', (PATCH_SIZE * step_width, PATCH_SIZE * st
5 anvas mask = Image.new('RGB', (PATCH SIZE * step width, PATCH SIZE * ste
```

```
4/30/2021
                             coms_4995_project_training_inception_augmentation.ipynb - Colaboratory
     6 ask blank = np.zeros((PATCH SIZE, PATCH SIZE))
     7 ask tumor = 255 * np.ones((PATCH SIZE, PATCH SIZE))
     8 \text{ ndex} = 0
     9
    10 or i in range(step width):
    11
          for j in range(step height):
    12
              canvas slide.paste(Image.fromarray(test slides normal 2[index],
    13
              if temp 2[index] == 0:
    14
                   canvas mask.paste(Image.fromarray(mask blank), (i*PATCH SIZE
    15
              else:
                   canvas mask.paste(Image.fromarray(mask tumor), (i*PATCH SIZE
    16
    17
              index += 1
    18
    19 lide name = 'patch to slide.png'
    20 ask name = 'patch to mask.png'
    21 anvas slide.save(slide name)
    22 anvas mask.save(mask name)
    23 mage slide = plt.imread(slide name)
    24 mage mask = plt.imread(mask name)
    25
    26 lt.imshow(image slide)
    27 lt.imshow(image mask, cmap='Reds', alpha=0.7)
```





time: 20 s (started: 2021-04-29 19:44:59 +00:00)