

C3M3_Assignment

August 10, 2020

1 Model Interpretation Methods

Welcome to the final assignment of course 3! In this assignment we will focus on the interpretation of machine learning and deep learning models. Using the techniques we've learned this week we'll revisit some of the models we've built throughout the course and try to understand a little more about what they're doing.

In this assignment you'll use various methods to interpret different types of machine learning models. In particular, you'll learn about the following topics:

- Interpreting Deep Learning Models
 - Understanding output using GradCAMs
- Feature Importance in Machine Learning
 - Permutation Method
 - SHAP Values

Let's get started.

1.0.1 This assignment covers the following topics:

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1.1 Packages

We'll first import the necessary packages for this assignment.

- keras: we'll use this framework to interact with our deep learning model
- matplotlib: standard plotting library
- pandas: we'll use this to manipulate data
- numpy: standard python library for numerical operations
- cv2: library that contains convenience functions for image processing
- sklearn: standard machine learning library
- lifelines: we'll use their implementation of the c-index
- shap: library for interpreting and visualizing machine learning models using shapley values

```
In [1]: import keras
        from keras import backend as K
        import matplotlib.pyplot as plt
        import pandas as pd
        import numpy as np
        import cv2
        import sklearn
        import lifelines
        import shap

        from util import *

        # This sets a common size for all the figures we will draw.
        plt.rcParams['figure.figsize'] = [10, 7]
```

Using TensorFlow backend.

1 Interpreting Deep Learning Models

To start, let's try understanding our X-ray diagnostic model from Course 1 Week 1. Run the next cell to load in the model (it should take a few seconds to complete).

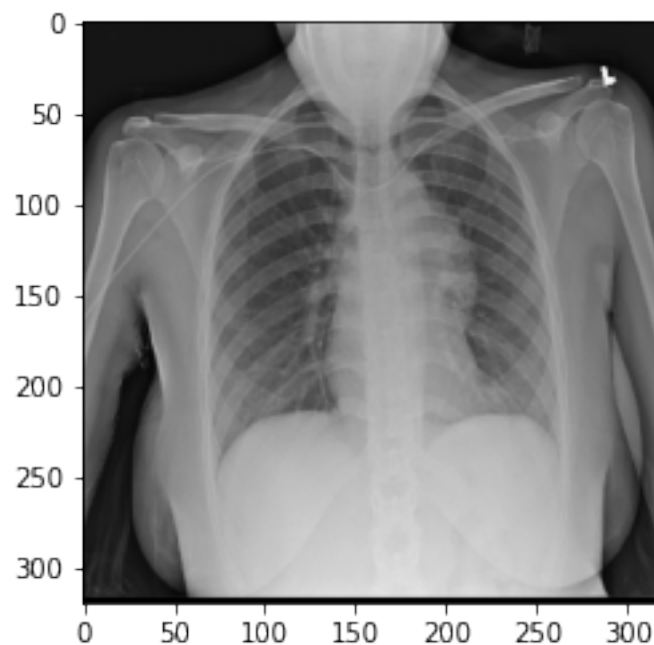
```
In [2]: model = load_C3M3_model()
```

```
Got loss weights
Loaded DenseNet
Added layers
```

Compiled Model
Loaded Weights

Let's load in an X-ray image to develop on. Run the next cell to load and show the image.

```
In [3]: IMAGE_DIR = 'nih_new/images-small/'  
df = pd.read_csv("nih_new/train-small.csv")  
im_path = IMAGE_DIR + '00025288_001.png'  
x = load_image(im_path, df, preprocess=False)  
plt.imshow(x, cmap = 'gray')  
plt.show()
```



```
In [12]: df.head()
```

```
Out[12]:
```

	Image	Atelectasis	Cardiomegaly	Consolidation	Edema	\
0	00008270_015.png	0	0	0	0	
1	00029855_001.png	1	0	0	0	
2	00001297_000.png	0	0	0	0	
3	00012359_002.png	0	0	0	0	
4	00017951_001.png	0	0	0	0	

	Effusion	Emphysema	Fibrosis	Hernia	Infiltration	Mass	Nodule	\
0	0	0	0	0	0	0	0	
1	1	0	0	0	1	0	0	
2	0	0	0	0	0	0	0	

3	0	0	0	0	0	0	0
4	0	0	0	0	1	0	0

	PatientId	Pleural_Thickening	Pneumonia	Pneumothorax
0	8270	0	0	0
1	29855	0	0	0
2	1297	1	0	0
3	12359	0	0	0
4	17951	0	0	0

Next, let's get our predictions. Before we plug the image into our model, we have to normalize it. Run the next cell to compute the mean and standard deviation of the images in our training set.

```
In [4]: mean, std = get_mean_std_per_batch(df)
```

```
In [13]: mean
```

```
Out[13]: 110.8784765625
```

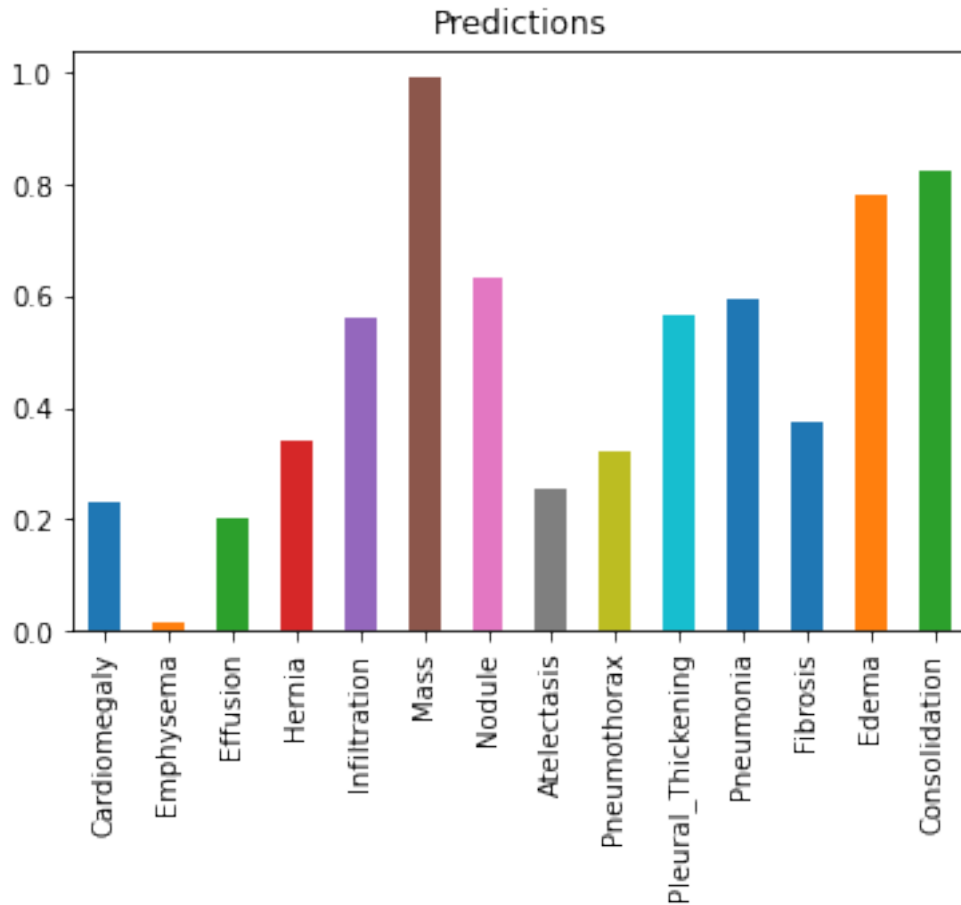
```
In [14]: std
```

```
Out[14]: 71.45956678555459
```

Now we are ready to normalize and run the image through our model to get predictions.

```
In [5]: labels = ['Cardiomegaly', 'Emphysema', 'Effusion', 'Hernia', 'Infiltration', 'Mass', 'Pneumothorax', 'Pleural_Thickening', 'Pneumonia', 'Fibrosis', 'Edema',
```

```
processed_image = load_image_normalize(im_path, mean, std)
preds = model.predict(processed_image)
pred_df = pd.DataFrame(preds, columns = labels)
pred_df.loc[0, :].plot.bar()
plt.title("Predictions")
plt.show()
```



We see, for example, that the model predicts Mass (abnormal spot or area in the lungs that are more than 3 centimeters) with high probability. Indeed, this patient was diagnosed with mass. However, we don't know where the model is looking when it's making its own diagnosis. To gain more insight into what the model is looking at, we can use GradCAMs.

1.1 GradCAM

GradCAM is a technique to visualize the impact of each region of an image on a specific output for a Convolutional Neural Network model. Through GradCAM, we can generate a heatmap by computing gradients of the specific class scores we are interested in visualizing.

1.1.1 Getting Intermediate Layers

Perhaps the most complicated part of computing GradCAM is accessing intermediate activations in our deep learning model and computing gradients with respect to the class output. Now we'll go over one pattern to accomplish this, which you can use when implementing GradCAM.

In order to understand how to access intermediate layers in a computation, first let's see the layers that our model is composed of. This can be done by calling Keras convenience function `model.summary()`. Do this in the cell below.

```
In [6]: model.summary()
```

Layer (type)	Output Shape	Param #	Connected to
--------------	--------------	---------	--------------

input_1 (InputLayer)	(None, None, None, 3 0	
zero_padding2d_1 (ZeroPadding2D)	(None, None, None, 3 0	input_1[0][0]
conv1/conv (Conv2D)	(None, None, None, 6 9408	zero_padding2d_1[0][0]
conv1/bn (BatchNormalization)	(None, None, None, 6 256	conv1/conv[0][0]
conv1/relu (Activation)	(None, None, None, 6 0	conv1/bn[0][0]
zero_padding2d_2 (ZeroPadding2D)	(None, None, None, 6 0	conv1/relu[0][0]
pool1 (MaxPooling2D)	(None, None, None, 6 0	zero_padding2d_2[0][0]
conv2_block1_0_bn (BatchNormali	(None, None, None, 6 256	pool1[0][0]
conv2_block1_0_relu (Activation	(None, None, None, 6 0	conv2_block1_0_bn[0][0]
conv2_block1_1_conv (Conv2D)	(None, None, None, 1 8192	conv2_block1_0_relu[0][0]
conv2_block1_1_bn (BatchNormali	(None, None, None, 1 512	conv2_block1_1_conv[0][0]
conv2_block1_1_relu (Activation	(None, None, None, 1 0	conv2_block1_1_bn[0][0]
conv2_block1_2_conv (Conv2D)	(None, None, None, 3 36864	conv2_block1_1_relu[0][0]
conv2_block1_concat (Concatenat	(None, None, None, 9 0	pool1[0][0] conv2_block1_2_conv[0][0]
conv2_block2_0_bn (BatchNormali	(None, None, None, 9 384	conv2_block1_concat[0][0]
conv2_block2_0_relu (Activation	(None, None, None, 9 0	conv2_block2_0_bn[0][0]
conv2_block2_1_conv (Conv2D)	(None, None, None, 1 12288	conv2_block2_0_relu[0][0]
conv2_block2_1_bn (BatchNormali	(None, None, None, 1 512	conv2_block2_1_conv[0][0]
conv2_block2_1_relu (Activation	(None, None, None, 1 0	conv2_block2_1_bn[0][0]
conv2_block2_2_conv (Conv2D)	(None, None, None, 3 36864	conv2_block2_1_relu[0][0]
conv2_block2_concat (Concatenat	(None, None, None, 1 0	conv2_block1_concat[0][0] conv2_block2_2_conv[0][0]
conv2_block3_0_bn (BatchNormali	(None, None, None, 1 512	conv2_block2_concat[0][0]
conv2_block3_0_relu (Activation	(None, None, None, 1 0	conv2_block3_0_bn[0][0]

conv2_block3_1_conv (Conv2D)	(None, None, None, 1 16384	conv2_block3_0_relu[0][0]
conv2_block3_1_bn (BatchNormali	(None, None, None, 1 512	conv2_block3_1_conv[0][0]
conv2_block3_1_relu (Activation	(None, None, None, 1 0	conv2_block3_1_bn[0][0]
conv2_block3_2_conv (Conv2D)	(None, None, None, 3 36864	conv2_block3_1_relu[0][0]
conv2_block3_concat (Concatenat	(None, None, None, 1 0	conv2_block2_concat[0][0] conv2_block3_2_conv[0][0]
conv2_block4_0_bn (BatchNormali	(None, None, None, 1 640	conv2_block3_concat[0][0]
conv2_block4_0_relu (Activation	(None, None, None, 1 0	conv2_block4_0_bn[0][0]
conv2_block4_1_conv (Conv2D)	(None, None, None, 1 20480	conv2_block4_0_relu[0][0]
conv2_block4_1_bn (BatchNormali	(None, None, None, 1 512	conv2_block4_1_conv[0][0]
conv2_block4_1_relu (Activation	(None, None, None, 1 0	conv2_block4_1_bn[0][0]
conv2_block4_2_conv (Conv2D)	(None, None, None, 3 36864	conv2_block4_1_relu[0][0]
conv2_block4_concat (Concatenat	(None, None, None, 1 0	conv2_block3_concat[0][0] conv2_block4_2_conv[0][0]
conv2_block5_0_bn (BatchNormali	(None, None, None, 1 768	conv2_block4_concat[0][0]
conv2_block5_0_relu (Activation	(None, None, None, 1 0	conv2_block5_0_bn[0][0]
conv2_block5_1_conv (Conv2D)	(None, None, None, 1 24576	conv2_block5_0_relu[0][0]
conv2_block5_1_bn (BatchNormali	(None, None, None, 1 512	conv2_block5_1_conv[0][0]
conv2_block5_1_relu (Activation	(None, None, None, 1 0	conv2_block5_1_bn[0][0]
conv2_block5_2_conv (Conv2D)	(None, None, None, 3 36864	conv2_block5_1_relu[0][0]
conv2_block5_concat (Concatenat	(None, None, None, 2 0	conv2_block4_concat[0][0] conv2_block5_2_conv[0][0]
conv2_block6_0_bn (BatchNormali	(None, None, None, 2 896	conv2_block5_concat[0][0]
conv2_block6_0_relu (Activation	(None, None, None, 2 0	conv2_block6_0_bn[0][0]
conv2_block6_1_conv (Conv2D)	(None, None, None, 1 28672	conv2_block6_0_relu[0][0]

conv2_block6_1_bn (BatchNormali	(None, None, None, 1 512	conv2_block6_1_conv[0][0]
conv2_block6_1_relu (Activation	(None, None, None, 1 0	conv2_block6_1_bn[0][0]
conv2_block6_2_conv (Conv2D)	(None, None, None, 3 36864	conv2_block6_1_relu[0][0]
conv2_block6_concat (Concatenat	(None, None, None, 2 0	conv2_block5_concat[0][0] conv2_block6_2_conv[0][0]
pool2_bn (BatchNormalization)	(None, None, None, 2 1024	conv2_block6_concat[0][0]
pool2_relu (Activation)	(None, None, None, 2 0	pool2_bn[0][0]
pool2_conv (Conv2D)	(None, None, None, 1 32768	pool2_relu[0][0]
pool2_pool (AveragePooling2D)	(None, None, None, 1 0	pool2_conv[0][0]
conv3_block1_0_bn (BatchNormali	(None, None, None, 1 512	pool2_pool[0][0]
conv3_block1_0_relu (Activation	(None, None, None, 1 0	conv3_block1_0_bn[0][0]
conv3_block1_1_conv (Conv2D)	(None, None, None, 1 16384	conv3_block1_0_relu[0][0]
conv3_block1_1_bn (BatchNormali	(None, None, None, 1 512	conv3_block1_1_conv[0][0]
conv3_block1_1_relu (Activation	(None, None, None, 1 0	conv3_block1_1_bn[0][0]
conv3_block1_2_conv (Conv2D)	(None, None, None, 3 36864	conv3_block1_1_relu[0][0]
conv3_block1_concat (Concatenat	(None, None, None, 1 0	pool2_pool[0][0] conv3_block1_2_conv[0][0]
conv3_block2_0_bn (BatchNormali	(None, None, None, 1 640	conv3_block1_concat[0][0]
conv3_block2_0_relu (Activation	(None, None, None, 1 0	conv3_block2_0_bn[0][0]
conv3_block2_1_conv (Conv2D)	(None, None, None, 1 20480	conv3_block2_0_relu[0][0]
conv3_block2_1_bn (BatchNormali	(None, None, None, 1 512	conv3_block2_1_conv[0][0]
conv3_block2_1_relu (Activation	(None, None, None, 1 0	conv3_block2_1_bn[0][0]
conv3_block2_2_conv (Conv2D)	(None, None, None, 3 36864	conv3_block2_1_relu[0][0]
conv3_block2_concat (Concatenat	(None, None, None, 1 0	conv3_block1_concat[0][0] conv3_block2_2_conv[0][0]
conv3_block3_0_bn (BatchNormali	(None, None, None, 1 768	conv3_block2_concat[0][0]

conv3_block3_0_relu	(Activation (None, None, None, 1 0	conv3_block3_0_bn[0][0]
conv3_block3_1_conv	(Conv2D) (None, None, None, 1 24576	conv3_block3_0_relu[0][0]
conv3_block3_1_bn	(BatchNormali (None, None, None, 1 512	conv3_block3_1_conv[0][0]
conv3_block3_1_relu	(Activation (None, None, None, 1 0	conv3_block3_1_bn[0][0]
conv3_block3_2_conv	(Conv2D) (None, None, None, 3 36864	conv3_block3_1_relu[0][0]
conv3_block3_concat	(Concatenat (None, None, None, 2 0	conv3_block2_concat[0][0] conv3_block3_2_conv[0][0]
conv3_block4_0_bn	(BatchNormali (None, None, None, 2 896	conv3_block3_concat[0][0]
conv3_block4_0_relu	(Activation (None, None, None, 2 0	conv3_block4_0_bn[0][0]
conv3_block4_1_conv	(Conv2D) (None, None, None, 1 28672	conv3_block4_0_relu[0][0]
conv3_block4_1_bn	(BatchNormali (None, None, None, 1 512	conv3_block4_1_conv[0][0]
conv3_block4_1_relu	(Activation (None, None, None, 1 0	conv3_block4_1_bn[0][0]
conv3_block4_2_conv	(Conv2D) (None, None, None, 3 36864	conv3_block4_1_relu[0][0]
conv3_block4_concat	(Concatenat (None, None, None, 2 0	conv3_block3_concat[0][0] conv3_block4_2_conv[0][0]
conv3_block5_0_bn	(BatchNormali (None, None, None, 2 1024	conv3_block4_concat[0][0]
conv3_block5_0_relu	(Activation (None, None, None, 2 0	conv3_block5_0_bn[0][0]
conv3_block5_1_conv	(Conv2D) (None, None, None, 1 32768	conv3_block5_0_relu[0][0]
conv3_block5_1_bn	(BatchNormali (None, None, None, 1 512	conv3_block5_1_conv[0][0]
conv3_block5_1_relu	(Activation (None, None, None, 1 0	conv3_block5_1_bn[0][0]
conv3_block5_2_conv	(Conv2D) (None, None, None, 3 36864	conv3_block5_1_relu[0][0]
conv3_block5_concat	(Concatenat (None, None, None, 2 0	conv3_block4_concat[0][0] conv3_block5_2_conv[0][0]
conv3_block6_0_bn	(BatchNormali (None, None, None, 2 1152	conv3_block5_concat[0][0]
conv3_block6_0_relu	(Activation (None, None, None, 2 0	conv3_block6_0_bn[0][0]

conv3_block6_1_conv (Conv2D)	(None, None, None, 1 36864	conv3_block6_0_relu[0][0]
conv3_block6_1_bn (BatchNormali	(None, None, None, 1 512	conv3_block6_1_conv[0][0]
conv3_block6_1_relu (Activation	(None, None, None, 1 0	conv3_block6_1_bn[0][0]
conv3_block6_2_conv (Conv2D)	(None, None, None, 3 36864	conv3_block6_1_relu[0][0]
conv3_block6_concat (Concatenat	(None, None, None, 3 0	conv3_block5_concat[0][0] conv3_block6_2_conv[0][0]
conv3_block7_0_bn (BatchNormali	(None, None, None, 3 1280	conv3_block6_concat[0][0]
conv3_block7_0_relu (Activation	(None, None, None, 3 0	conv3_block7_0_bn[0][0]
conv3_block7_1_conv (Conv2D)	(None, None, None, 1 40960	conv3_block7_0_relu[0][0]
conv3_block7_1_bn (BatchNormali	(None, None, None, 1 512	conv3_block7_1_conv[0][0]
conv3_block7_1_relu (Activation	(None, None, None, 1 0	conv3_block7_1_bn[0][0]
conv3_block7_2_conv (Conv2D)	(None, None, None, 3 36864	conv3_block7_1_relu[0][0]
conv3_block7_concat (Concatenat	(None, None, None, 3 0	conv3_block6_concat[0][0] conv3_block7_2_conv[0][0]
conv3_block8_0_bn (BatchNormali	(None, None, None, 3 1408	conv3_block7_concat[0][0]
conv3_block8_0_relu (Activation	(None, None, None, 3 0	conv3_block8_0_bn[0][0]
conv3_block8_1_conv (Conv2D)	(None, None, None, 1 45056	conv3_block8_0_relu[0][0]
conv3_block8_1_bn (BatchNormali	(None, None, None, 1 512	conv3_block8_1_conv[0][0]
conv3_block8_1_relu (Activation	(None, None, None, 1 0	conv3_block8_1_bn[0][0]
conv3_block8_2_conv (Conv2D)	(None, None, None, 3 36864	conv3_block8_1_relu[0][0]
conv3_block8_concat (Concatenat	(None, None, None, 3 0	conv3_block7_concat[0][0] conv3_block8_2_conv[0][0]
conv3_block9_0_bn (BatchNormali	(None, None, None, 3 1536	conv3_block8_concat[0][0]
conv3_block9_0_relu (Activation	(None, None, None, 3 0	conv3_block9_0_bn[0][0]
conv3_block9_1_conv (Conv2D)	(None, None, None, 1 49152	conv3_block9_0_relu[0][0]
conv3_block9_1_bn (BatchNormali	(None, None, None, 1 512	conv3_block9_1_conv[0][0]

conv3_block9_1_relu (Activation	(None, None, None, 1 0	conv3_block9_1_bn[0][0]
conv3_block9_2_conv (Conv2D)	(None, None, None, 3 36864	conv3_block9_1_relu[0][0]
conv3_block9_concat (Concatenat	(None, None, None, 4 0	conv3_block8_concat[0][0] conv3_block9_2_conv[0][0]
conv3_block10_0_bn (BatchNormal	(None, None, None, 4 1664	conv3_block9_concat[0][0]
conv3_block10_0_relu (Activatio	(None, None, None, 4 0	conv3_block10_0_bn[0][0]
conv3_block10_1_conv (Conv2D)	(None, None, None, 1 53248	conv3_block10_0_relu[0][0]
conv3_block10_1_bn (BatchNormal	(None, None, None, 1 512	conv3_block10_1_conv[0][0]
conv3_block10_1_relu (Activatio	(None, None, None, 1 0	conv3_block10_1_bn[0][0]
conv3_block10_2_conv (Conv2D)	(None, None, None, 3 36864	conv3_block10_1_relu[0][0]
conv3_block10_concat (Concatena	(None, None, None, 4 0	conv3_block9_concat[0][0] conv3_block10_2_conv[0][0]
conv3_block11_0_bn (BatchNormal	(None, None, None, 4 1792	conv3_block10_concat[0][0]
conv3_block11_0_relu (Activatio	(None, None, None, 4 0	conv3_block11_0_bn[0][0]
conv3_block11_1_conv (Conv2D)	(None, None, None, 1 57344	conv3_block11_0_relu[0][0]
conv3_block11_1_bn (BatchNormal	(None, None, None, 1 512	conv3_block11_1_conv[0][0]
conv3_block11_1_relu (Activatio	(None, None, None, 1 0	conv3_block11_1_bn[0][0]
conv3_block11_2_conv (Conv2D)	(None, None, None, 3 36864	conv3_block11_1_relu[0][0]
conv3_block11_concat (Concatena	(None, None, None, 4 0	conv3_block10_concat[0][0] conv3_block11_2_conv[0][0]
conv3_block12_0_bn (BatchNormal	(None, None, None, 4 1920	conv3_block11_concat[0][0]
conv3_block12_0_relu (Activatio	(None, None, None, 4 0	conv3_block12_0_bn[0][0]
conv3_block12_1_conv (Conv2D)	(None, None, None, 1 61440	conv3_block12_0_relu[0][0]
conv3_block12_1_bn (BatchNormal	(None, None, None, 1 512	conv3_block12_1_conv[0][0]
conv3_block12_1_relu (Activatio	(None, None, None, 1 0	conv3_block12_1_bn[0][0]

conv3_block12_2_conv (Conv2D)	(None, None, None, 3 36864	conv3_block12_1_relu[0][0]
conv3_block12_concat (Concatena	(None, None, None, 5 0	conv3_block11_concat[0][0] conv3_block12_2_conv[0][0]
pool3_bn (BatchNormalization)	(None, None, None, 5 2048	conv3_block12_concat[0][0]
pool3_relu (Activation)	(None, None, None, 5 0	pool3_bn[0][0]
pool3_conv (Conv2D)	(None, None, None, 2 131072	pool3_relu[0][0]
pool3_pool (AveragePooling2D)	(None, None, None, 2 0	pool3_conv[0][0]
conv4_block1_0_bn (BatchNormali	(None, None, None, 2 1024	pool3_pool[0][0]
conv4_block1_0_relu (Activation	(None, None, None, 2 0	conv4_block1_0_bn[0][0]
conv4_block1_1_conv (Conv2D)	(None, None, None, 1 32768	conv4_block1_0_relu[0][0]
conv4_block1_1_bn (BatchNormali	(None, None, None, 1 512	conv4_block1_1_conv[0][0]
conv4_block1_1_relu (Activation	(None, None, None, 1 0	conv4_block1_1_bn[0][0]
conv4_block1_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block1_1_relu[0][0]
conv4_block1_concat (Concatenat	(None, None, None, 2 0	pool3_pool[0][0] conv4_block1_2_conv[0][0]
conv4_block2_0_bn (BatchNormali	(None, None, None, 2 1152	conv4_block1_concat[0][0]
conv4_block2_0_relu (Activation	(None, None, None, 2 0	conv4_block2_0_bn[0][0]
conv4_block2_1_conv (Conv2D)	(None, None, None, 1 36864	conv4_block2_0_relu[0][0]
conv4_block2_1_bn (BatchNormali	(None, None, None, 1 512	conv4_block2_1_conv[0][0]
conv4_block2_1_relu (Activation	(None, None, None, 1 0	conv4_block2_1_bn[0][0]
conv4_block2_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block2_1_relu[0][0]
conv4_block2_concat (Concatenat	(None, None, None, 3 0	conv4_block1_concat[0][0] conv4_block2_2_conv[0][0]
conv4_block3_0_bn (BatchNormali	(None, None, None, 3 1280	conv4_block2_concat[0][0]
conv4_block3_0_relu (Activation	(None, None, None, 3 0	conv4_block3_0_bn[0][0]
conv4_block3_1_conv (Conv2D)	(None, None, None, 1 40960	conv4_block3_0_relu[0][0]

conv4_block3_1_bn (BatchNormali	(None, None, None, 1 512	conv4_block3_1_conv[0][0]
conv4_block3_1_relu (Activation	(None, None, None, 1 0	conv4_block3_1_bn[0][0]
conv4_block3_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block3_1_relu[0][0]
conv4_block3_concat (Concatenat	(None, None, None, 3 0	conv4_block2_concat[0][0] conv4_block3_2_conv[0][0]
conv4_block4_0_bn (BatchNormali	(None, None, None, 3 1408	conv4_block3_concat[0][0]
conv4_block4_0_relu (Activation	(None, None, None, 3 0	conv4_block4_0_bn[0][0]
conv4_block4_1_conv (Conv2D)	(None, None, None, 1 45056	conv4_block4_0_relu[0][0]
conv4_block4_1_bn (BatchNormali	(None, None, None, 1 512	conv4_block4_1_conv[0][0]
conv4_block4_1_relu (Activation	(None, None, None, 1 0	conv4_block4_1_bn[0][0]
conv4_block4_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block4_1_relu[0][0]
conv4_block4_concat (Concatenat	(None, None, None, 3 0	conv4_block3_concat[0][0] conv4_block4_2_conv[0][0]
conv4_block5_0_bn (BatchNormali	(None, None, None, 3 1536	conv4_block4_concat[0][0]
conv4_block5_0_relu (Activation	(None, None, None, 3 0	conv4_block5_0_bn[0][0]
conv4_block5_1_conv (Conv2D)	(None, None, None, 1 49152	conv4_block5_0_relu[0][0]
conv4_block5_1_bn (BatchNormali	(None, None, None, 1 512	conv4_block5_1_conv[0][0]
conv4_block5_1_relu (Activation	(None, None, None, 1 0	conv4_block5_1_bn[0][0]
conv4_block5_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block5_1_relu[0][0]
conv4_block5_concat (Concatenat	(None, None, None, 4 0	conv4_block4_concat[0][0] conv4_block5_2_conv[0][0]
conv4_block6_0_bn (BatchNormali	(None, None, None, 4 1664	conv4_block5_concat[0][0]
conv4_block6_0_relu (Activation	(None, None, None, 4 0	conv4_block6_0_bn[0][0]
conv4_block6_1_conv (Conv2D)	(None, None, None, 1 53248	conv4_block6_0_relu[0][0]
conv4_block6_1_bn (BatchNormali	(None, None, None, 1 512	conv4_block6_1_conv[0][0]

conv4_block6_1_relu (Activation (None, None, None, 1 0	conv4_block6_1_bn[0] [0]
conv4_block6_2_conv (Conv2D) (None, None, None, 3 36864	conv4_block6_1_relu[0] [0]
conv4_block6_concat (Concatenat (None, None, None, 4 0	conv4_block5_concat[0] [0] conv4_block6_2_conv[0] [0]
conv4_block7_0_bn (BatchNormali (None, None, None, 4 1792	conv4_block6_concat[0] [0]
conv4_block7_0_relu (Activation (None, None, None, 4 0	conv4_block7_0_bn[0] [0]
conv4_block7_1_conv (Conv2D) (None, None, None, 1 57344	conv4_block7_0_relu[0] [0]
conv4_block7_1_bn (BatchNormali (None, None, None, 1 512	conv4_block7_1_conv[0] [0]
conv4_block7_1_relu (Activation (None, None, None, 1 0	conv4_block7_1_bn[0] [0]
conv4_block7_2_conv (Conv2D) (None, None, None, 3 36864	conv4_block7_1_relu[0] [0]
conv4_block7_concat (Concatenat (None, None, None, 4 0	conv4_block6_concat[0] [0] conv4_block7_2_conv[0] [0]
conv4_block8_0_bn (BatchNormali (None, None, None, 4 1920	conv4_block7_concat[0] [0]
conv4_block8_0_relu (Activation (None, None, None, 4 0	conv4_block8_0_bn[0] [0]
conv4_block8_1_conv (Conv2D) (None, None, None, 1 61440	conv4_block8_0_relu[0] [0]
conv4_block8_1_bn (BatchNormali (None, None, None, 1 512	conv4_block8_1_conv[0] [0]
conv4_block8_1_relu (Activation (None, None, None, 1 0	conv4_block8_1_bn[0] [0]
conv4_block8_2_conv (Conv2D) (None, None, None, 3 36864	conv4_block8_1_relu[0] [0]
conv4_block8_concat (Concatenat (None, None, None, 5 0	conv4_block7_concat[0] [0] conv4_block8_2_conv[0] [0]
conv4_block9_0_bn (BatchNormali (None, None, None, 5 2048	conv4_block8_concat[0] [0]
conv4_block9_0_relu (Activation (None, None, None, 5 0	conv4_block9_0_bn[0] [0]
conv4_block9_1_conv (Conv2D) (None, None, None, 1 65536	conv4_block9_0_relu[0] [0]
conv4_block9_1_bn (BatchNormali (None, None, None, 1 512	conv4_block9_1_conv[0] [0]
conv4_block9_1_relu (Activation (None, None, None, 1 0	conv4_block9_1_bn[0] [0]
conv4_block9_2_conv (Conv2D) (None, None, None, 3 36864	conv4_block9_1_relu[0] [0]

conv4_block9_concat (Concatenat	(None, None, None, 5 0	conv4_block8_concat[0][0] conv4_block9_2_conv[0][0]
conv4_block10_0_bn (BatchNormal	(None, None, None, 5 2176	conv4_block9_concat[0][0]
conv4_block10_0_relu (Activatio	(None, None, None, 5 0	conv4_block10_0_bn[0][0]
conv4_block10_1_conv (Conv2D)	(None, None, None, 1 69632	conv4_block10_0_relu[0][0]
conv4_block10_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block10_1_conv[0][0]
conv4_block10_1_relu (Activatio	(None, None, None, 1 0	conv4_block10_1_bn[0][0]
conv4_block10_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block10_1_relu[0][0]
conv4_block10_concat (Concatena	(None, None, None, 5 0	conv4_block9_concat[0][0] conv4_block10_2_conv[0][0]
conv4_block11_0_bn (BatchNormal	(None, None, None, 5 2304	conv4_block10_concat[0][0]
conv4_block11_0_relu (Activatio	(None, None, None, 5 0	conv4_block11_0_bn[0][0]
conv4_block11_1_conv (Conv2D)	(None, None, None, 1 73728	conv4_block11_0_relu[0][0]
conv4_block11_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block11_1_conv[0][0]
conv4_block11_1_relu (Activatio	(None, None, None, 1 0	conv4_block11_1_bn[0][0]
conv4_block11_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block11_1_relu[0][0]
conv4_block11_concat (Concatena	(None, None, None, 6 0	conv4_block10_concat[0][0] conv4_block11_2_conv[0][0]
conv4_block12_0_bn (BatchNormal	(None, None, None, 6 2432	conv4_block11_concat[0][0]
conv4_block12_0_relu (Activatio	(None, None, None, 6 0	conv4_block12_0_bn[0][0]
conv4_block12_1_conv (Conv2D)	(None, None, None, 1 77824	conv4_block12_0_relu[0][0]
conv4_block12_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block12_1_conv[0][0]
conv4_block12_1_relu (Activatio	(None, None, None, 1 0	conv4_block12_1_bn[0][0]
conv4_block12_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block12_1_relu[0][0]
conv4_block12_concat (Concatena	(None, None, None, 6 0	conv4_block11_concat[0][0] conv4_block12_2_conv[0][0]

conv4_block13_0_bn (BatchNormal	(None, None, None, 6 2560	conv4_block12_concat[0][0]
conv4_block13_0_relu (Activatio	(None, None, None, 6 0	conv4_block13_0_bn[0][0]
conv4_block13_1_conv (Conv2D)	(None, None, None, 1 81920	conv4_block13_0_relu[0][0]
conv4_block13_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block13_1_conv[0][0]
conv4_block13_1_relu (Activatio	(None, None, None, 1 0	conv4_block13_1_bn[0][0]
conv4_block13_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block13_1_relu[0][0]
conv4_block13_concat (Concatena	(None, None, None, 6 0	conv4_block12_concat[0][0] conv4_block13_2_conv[0][0]
conv4_block14_0_bn (BatchNormal	(None, None, None, 6 2688	conv4_block13_concat[0][0]
conv4_block14_0_relu (Activatio	(None, None, None, 6 0	conv4_block14_0_bn[0][0]
conv4_block14_1_conv (Conv2D)	(None, None, None, 1 86016	conv4_block14_0_relu[0][0]
conv4_block14_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block14_1_conv[0][0]
conv4_block14_1_relu (Activatio	(None, None, None, 1 0	conv4_block14_1_bn[0][0]
conv4_block14_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block14_1_relu[0][0]
conv4_block14_concat (Concatena	(None, None, None, 7 0	conv4_block13_concat[0][0] conv4_block14_2_conv[0][0]
conv4_block15_0_bn (BatchNormal	(None, None, None, 7 2816	conv4_block14_concat[0][0]
conv4_block15_0_relu (Activatio	(None, None, None, 7 0	conv4_block15_0_bn[0][0]
conv4_block15_1_conv (Conv2D)	(None, None, None, 1 90112	conv4_block15_0_relu[0][0]
conv4_block15_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block15_1_conv[0][0]
conv4_block15_1_relu (Activatio	(None, None, None, 1 0	conv4_block15_1_bn[0][0]
conv4_block15_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block15_1_relu[0][0]
conv4_block15_concat (Concatena	(None, None, None, 7 0	conv4_block14_concat[0][0] conv4_block15_2_conv[0][0]
conv4_block16_0_bn (BatchNormal	(None, None, None, 7 2944	conv4_block15_concat[0][0]

conv4_block16_0_relu (Activatio	(None, None, None, 7 0	conv4_block16_0_bn[0][0]
conv4_block16_1_conv (Conv2D)	(None, None, None, 1 94208	conv4_block16_0_relu[0][0]
conv4_block16_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block16_1_conv[0][0]
conv4_block16_1_relu (Activatio	(None, None, None, 1 0	conv4_block16_1_bn[0][0]
conv4_block16_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block16_1_relu[0][0]
conv4_block16_concat (Concatena	(None, None, None, 7 0	conv4_block15_concat[0][0] conv4_block16_2_conv[0][0]
conv4_block17_0_bn (BatchNormal	(None, None, None, 7 3072	conv4_block16_concat[0][0]
conv4_block17_0_relu (Activatio	(None, None, None, 7 0	conv4_block17_0_bn[0][0]
conv4_block17_1_conv (Conv2D)	(None, None, None, 1 98304	conv4_block17_0_relu[0][0]
conv4_block17_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block17_1_conv[0][0]
conv4_block17_1_relu (Activatio	(None, None, None, 1 0	conv4_block17_1_bn[0][0]
conv4_block17_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block17_1_relu[0][0]
conv4_block17_concat (Concatena	(None, None, None, 8 0	conv4_block16_concat[0][0] conv4_block17_2_conv[0][0]
conv4_block18_0_bn (BatchNormal	(None, None, None, 8 3200	conv4_block17_concat[0][0]
conv4_block18_0_relu (Activatio	(None, None, None, 8 0	conv4_block18_0_bn[0][0]
conv4_block18_1_conv (Conv2D)	(None, None, None, 1 102400	conv4_block18_0_relu[0][0]
conv4_block18_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block18_1_conv[0][0]
conv4_block18_1_relu (Activatio	(None, None, None, 1 0	conv4_block18_1_bn[0][0]
conv4_block18_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block18_1_relu[0][0]
conv4_block18_concat (Concatena	(None, None, None, 8 0	conv4_block17_concat[0][0] conv4_block18_2_conv[0][0]
conv4_block19_0_bn (BatchNormal	(None, None, None, 8 3328	conv4_block18_concat[0][0]
conv4_block19_0_relu (Activatio	(None, None, None, 8 0	conv4_block19_0_bn[0][0]
conv4_block19_1_conv (Conv2D)	(None, None, None, 1 106496	conv4_block19_0_relu[0][0]

conv4_block19_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block19_1_conv[0][0]
conv4_block19_1_relu (Activatio	(None, None, None, 1 0	conv4_block19_1_bn[0][0]
conv4_block19_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block19_1_relu[0][0]
conv4_block19_concat (Concatena	(None, None, None, 8 0	conv4_block18_concat[0][0] conv4_block19_2_conv[0][0]
conv4_block20_0_bn (BatchNormal	(None, None, None, 8 3456	conv4_block19_concat[0][0]
conv4_block20_0_relu (Activatio	(None, None, None, 8 0	conv4_block20_0_bn[0][0]
conv4_block20_1_conv (Conv2D)	(None, None, None, 1 110592	conv4_block20_0_relu[0][0]
conv4_block20_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block20_1_conv[0][0]
conv4_block20_1_relu (Activatio	(None, None, None, 1 0	conv4_block20_1_bn[0][0]
conv4_block20_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block20_1_relu[0][0]
conv4_block20_concat (Concatena	(None, None, None, 8 0	conv4_block19_concat[0][0] conv4_block20_2_conv[0][0]
conv4_block21_0_bn (BatchNormal	(None, None, None, 8 3584	conv4_block20_concat[0][0]
conv4_block21_0_relu (Activatio	(None, None, None, 8 0	conv4_block21_0_bn[0][0]
conv4_block21_1_conv (Conv2D)	(None, None, None, 1 114688	conv4_block21_0_relu[0][0]
conv4_block21_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block21_1_conv[0][0]
conv4_block21_1_relu (Activatio	(None, None, None, 1 0	conv4_block21_1_bn[0][0]
conv4_block21_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block21_1_relu[0][0]
conv4_block21_concat (Concatena	(None, None, None, 9 0	conv4_block20_concat[0][0] conv4_block21_2_conv[0][0]
conv4_block22_0_bn (BatchNormal	(None, None, None, 9 3712	conv4_block21_concat[0][0]
conv4_block22_0_relu (Activatio	(None, None, None, 9 0	conv4_block22_0_bn[0][0]
conv4_block22_1_conv (Conv2D)	(None, None, None, 1 118784	conv4_block22_0_relu[0][0]
conv4_block22_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block22_1_conv[0][0]

conv4_block22_1_relu (Activation)	(None, None, None, 1 0	conv4_block22_1_bn[0][0]
conv4_block22_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block22_1_relu[0][0]
conv4_block22_concat (Concatenation)	(None, None, None, 9 0	conv4_block21_concat[0][0] conv4_block22_2_conv[0][0]
conv4_block23_0_bn (BatchNormal	(None, None, None, 9 3840	conv4_block22_concat[0][0]
conv4_block23_0_relu (Activation)	(None, None, None, 9 0	conv4_block23_0_bn[0][0]
conv4_block23_1_conv (Conv2D)	(None, None, None, 1 122880	conv4_block23_0_relu[0][0]
conv4_block23_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block23_1_conv[0][0]
conv4_block23_1_relu (Activation)	(None, None, None, 1 0	conv4_block23_1_bn[0][0]
conv4_block23_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block23_1_relu[0][0]
conv4_block23_concat (Concatenation)	(None, None, None, 9 0	conv4_block22_concat[0][0] conv4_block23_2_conv[0][0]
conv4_block24_0_bn (BatchNormal	(None, None, None, 9 3968	conv4_block23_concat[0][0]
conv4_block24_0_relu (Activation)	(None, None, None, 9 0	conv4_block24_0_bn[0][0]
conv4_block24_1_conv (Conv2D)	(None, None, None, 1 126976	conv4_block24_0_relu[0][0]
conv4_block24_1_bn (BatchNormal	(None, None, None, 1 512	conv4_block24_1_conv[0][0]
conv4_block24_1_relu (Activation)	(None, None, None, 1 0	conv4_block24_1_bn[0][0]
conv4_block24_2_conv (Conv2D)	(None, None, None, 3 36864	conv4_block24_1_relu[0][0]
conv4_block24_concat (Concatenation)	(None, None, None, 1 0	conv4_block23_concat[0][0] conv4_block24_2_conv[0][0]
pool4_bn (BatchNormalization)	(None, None, None, 1 4096	conv4_block24_concat[0][0]
pool4_relu (Activation)	(None, None, None, 1 0	pool4_bn[0][0]
pool4_conv (Conv2D)	(None, None, None, 5 524288	pool4_relu[0][0]
pool4_pool (AveragePooling2D)	(None, None, None, 5 0	pool4_conv[0][0]
conv5_block1_0_bn (BatchNormali	(None, None, None, 5 2048	pool4_pool[0][0]
conv5_block1_0_relu (Activation)	(None, None, None, 5 0	conv5_block1_0_bn[0][0]

conv5_block1_1_conv (Conv2D)	(None, None, None, 1 65536	conv5_block1_0_relu[0][0]
conv5_block1_1_bn (BatchNormali	(None, None, None, 1 512	conv5_block1_1_conv[0][0]
conv5_block1_1_relu (Activation	(None, None, None, 1 0	conv5_block1_1_bn[0][0]
conv5_block1_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block1_1_relu[0][0]
conv5_block1_concat (Concatenat	(None, None, None, 5 0	pool4_pool[0][0] conv5_block1_2_conv[0][0]
conv5_block2_0_bn (BatchNormali	(None, None, None, 5 2176	conv5_block1_concat[0][0]
conv5_block2_0_relu (Activation	(None, None, None, 5 0	conv5_block2_0_bn[0][0]
conv5_block2_1_conv (Conv2D)	(None, None, None, 1 69632	conv5_block2_0_relu[0][0]
conv5_block2_1_bn (BatchNormali	(None, None, None, 1 512	conv5_block2_1_conv[0][0]
conv5_block2_1_relu (Activation	(None, None, None, 1 0	conv5_block2_1_bn[0][0]
conv5_block2_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block2_1_relu[0][0]
conv5_block2_concat (Concatenat	(None, None, None, 5 0	conv5_block1_concat[0][0] conv5_block2_2_conv[0][0]
conv5_block3_0_bn (BatchNormali	(None, None, None, 5 2304	conv5_block2_concat[0][0]
conv5_block3_0_relu (Activation	(None, None, None, 5 0	conv5_block3_0_bn[0][0]
conv5_block3_1_conv (Conv2D)	(None, None, None, 1 73728	conv5_block3_0_relu[0][0]
conv5_block3_1_bn (BatchNormali	(None, None, None, 1 512	conv5_block3_1_conv[0][0]
conv5_block3_1_relu (Activation	(None, None, None, 1 0	conv5_block3_1_bn[0][0]
conv5_block3_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block3_1_relu[0][0]
conv5_block3_concat (Concatenat	(None, None, None, 6 0	conv5_block2_concat[0][0] conv5_block3_2_conv[0][0]
conv5_block4_0_bn (BatchNormali	(None, None, None, 6 2432	conv5_block3_concat[0][0]
conv5_block4_0_relu (Activation	(None, None, None, 6 0	conv5_block4_0_bn[0][0]
conv5_block4_1_conv (Conv2D)	(None, None, None, 1 77824	conv5_block4_0_relu[0][0]

conv5_block4_1_bn (BatchNormali	(None, None, None, 1 512	conv5_block4_1_conv[0][0]
conv5_block4_1_relu (Activation	(None, None, None, 1 0	conv5_block4_1_bn[0][0]
conv5_block4_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block4_1_relu[0][0]
conv5_block4_concat (Concatenat	(None, None, None, 6 0	conv5_block3_concat[0][0] conv5_block4_2_conv[0][0]
conv5_block5_0_bn (BatchNormali	(None, None, None, 6 2560	conv5_block4_concat[0][0]
conv5_block5_0_relu (Activation	(None, None, None, 6 0	conv5_block5_0_bn[0][0]
conv5_block5_1_conv (Conv2D)	(None, None, None, 1 81920	conv5_block5_0_relu[0][0]
conv5_block5_1_bn (BatchNormali	(None, None, None, 1 512	conv5_block5_1_conv[0][0]
conv5_block5_1_relu (Activation	(None, None, None, 1 0	conv5_block5_1_bn[0][0]
conv5_block5_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block5_1_relu[0][0]
conv5_block5_concat (Concatenat	(None, None, None, 6 0	conv5_block4_concat[0][0] conv5_block5_2_conv[0][0]
conv5_block6_0_bn (BatchNormali	(None, None, None, 6 2688	conv5_block5_concat[0][0]
conv5_block6_0_relu (Activation	(None, None, None, 6 0	conv5_block6_0_bn[0][0]
conv5_block6_1_conv (Conv2D)	(None, None, None, 1 86016	conv5_block6_0_relu[0][0]
conv5_block6_1_bn (BatchNormali	(None, None, None, 1 512	conv5_block6_1_conv[0][0]
conv5_block6_1_relu (Activation	(None, None, None, 1 0	conv5_block6_1_bn[0][0]
conv5_block6_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block6_1_relu[0][0]
conv5_block6_concat (Concatenat	(None, None, None, 7 0	conv5_block5_concat[0][0] conv5_block6_2_conv[0][0]
conv5_block7_0_bn (BatchNormali	(None, None, None, 7 2816	conv5_block6_concat[0][0]
conv5_block7_0_relu (Activation	(None, None, None, 7 0	conv5_block7_0_bn[0][0]
conv5_block7_1_conv (Conv2D)	(None, None, None, 1 90112	conv5_block7_0_relu[0][0]
conv5_block7_1_bn (BatchNormali	(None, None, None, 1 512	conv5_block7_1_conv[0][0]
conv5_block7_1_relu (Activation	(None, None, None, 1 0	conv5_block7_1_bn[0][0]

conv5_block7_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block7_1_relu[0][0]
conv5_block7_concat (Concatenat	(None, None, None, 7 0	conv5_block6_concat[0][0] conv5_block7_2_conv[0][0]
conv5_block8_0_bn (BatchNormali	(None, None, None, 7 2944	conv5_block7_concat[0][0]
conv5_block8_0_relu (Activation	(None, None, None, 7 0	conv5_block8_0_bn[0][0]
conv5_block8_1_conv (Conv2D)	(None, None, None, 1 94208	conv5_block8_0_relu[0][0]
conv5_block8_1_bn (BatchNormali	(None, None, None, 1 512	conv5_block8_1_conv[0][0]
conv5_block8_1_relu (Activation	(None, None, None, 1 0	conv5_block8_1_bn[0][0]
conv5_block8_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block8_1_relu[0][0]
conv5_block8_concat (Concatenat	(None, None, None, 7 0	conv5_block7_concat[0][0] conv5_block8_2_conv[0][0]
conv5_block9_0_bn (BatchNormali	(None, None, None, 7 3072	conv5_block8_concat[0][0]
conv5_block9_0_relu (Activation	(None, None, None, 7 0	conv5_block9_0_bn[0][0]
conv5_block9_1_conv (Conv2D)	(None, None, None, 1 98304	conv5_block9_0_relu[0][0]
conv5_block9_1_bn (BatchNormali	(None, None, None, 1 512	conv5_block9_1_conv[0][0]
conv5_block9_1_relu (Activation	(None, None, None, 1 0	conv5_block9_1_bn[0][0]
conv5_block9_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block9_1_relu[0][0]
conv5_block9_concat (Concatenat	(None, None, None, 8 0	conv5_block8_concat[0][0] conv5_block9_2_conv[0][0]
conv5_block10_0_bn (BatchNormal	(None, None, None, 8 3200	conv5_block9_concat[0][0]
conv5_block10_0_relu (Activatio	(None, None, None, 8 0	conv5_block10_0_bn[0][0]
conv5_block10_1_conv (Conv2D)	(None, None, None, 1 102400	conv5_block10_0_relu[0][0]
conv5_block10_1_bn (BatchNormal	(None, None, None, 1 512	conv5_block10_1_conv[0][0]
conv5_block10_1_relu (Activatio	(None, None, None, 1 0	conv5_block10_1_bn[0][0]
conv5_block10_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block10_1_relu[0][0]

conv5_block10_concat (Concatena	(None, None, None, 8 0	conv5_block9_concat[0][0] conv5_block10_2_conv[0][0]
conv5_block11_0_bn (BatchNormal	(None, None, None, 8 3328	conv5_block10_concat[0][0]
conv5_block11_0_relu (Activatio	(None, None, None, 8 0	conv5_block11_0_bn[0][0]
conv5_block11_1_conv (Conv2D)	(None, None, None, 1 106496	conv5_block11_0_relu[0][0]
conv5_block11_1_bn (BatchNormal	(None, None, None, 1 512	conv5_block11_1_conv[0][0]
conv5_block11_1_relu (Activatio	(None, None, None, 1 0	conv5_block11_1_bn[0][0]
conv5_block11_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block11_1_relu[0][0]
conv5_block11_concat (Concatena	(None, None, None, 8 0	conv5_block10_concat[0][0] conv5_block11_2_conv[0][0]
conv5_block12_0_bn (BatchNormal	(None, None, None, 8 3456	conv5_block11_concat[0][0]
conv5_block12_0_relu (Activatio	(None, None, None, 8 0	conv5_block12_0_bn[0][0]
conv5_block12_1_conv (Conv2D)	(None, None, None, 1 110592	conv5_block12_0_relu[0][0]
conv5_block12_1_bn (BatchNormal	(None, None, None, 1 512	conv5_block12_1_conv[0][0]
conv5_block12_1_relu (Activatio	(None, None, None, 1 0	conv5_block12_1_bn[0][0]
conv5_block12_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block12_1_relu[0][0]
conv5_block12_concat (Concatena	(None, None, None, 8 0	conv5_block11_concat[0][0] conv5_block12_2_conv[0][0]
conv5_block13_0_bn (BatchNormal	(None, None, None, 8 3584	conv5_block12_concat[0][0]
conv5_block13_0_relu (Activatio	(None, None, None, 8 0	conv5_block13_0_bn[0][0]
conv5_block13_1_conv (Conv2D)	(None, None, None, 1 114688	conv5_block13_0_relu[0][0]
conv5_block13_1_bn (BatchNormal	(None, None, None, 1 512	conv5_block13_1_conv[0][0]
conv5_block13_1_relu (Activatio	(None, None, None, 1 0	conv5_block13_1_bn[0][0]
conv5_block13_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block13_1_relu[0][0]
conv5_block13_concat (Concatena	(None, None, None, 9 0	conv5_block12_concat[0][0] conv5_block13_2_conv[0][0]

conv5_block14_0_bn (BatchNormal	(None, None, None, 9 3712	conv5_block13_concat[0][0]
conv5_block14_0_relu (Activatio	(None, None, None, 9 0	conv5_block14_0_bn[0][0]
conv5_block14_1_conv (Conv2D)	(None, None, None, 1 118784	conv5_block14_0_relu[0][0]
conv5_block14_1_bn (BatchNormal	(None, None, None, 1 512	conv5_block14_1_conv[0][0]
conv5_block14_1_relu (Activatio	(None, None, None, 1 0	conv5_block14_1_bn[0][0]
conv5_block14_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block14_1_relu[0][0]
conv5_block14_concat (Concatena	(None, None, None, 9 0	conv5_block13_concat[0][0] conv5_block14_2_conv[0][0]
conv5_block15_0_bn (BatchNormal	(None, None, None, 9 3840	conv5_block14_concat[0][0]
conv5_block15_0_relu (Activatio	(None, None, None, 9 0	conv5_block15_0_bn[0][0]
conv5_block15_1_conv (Conv2D)	(None, None, None, 1 122880	conv5_block15_0_relu[0][0]
conv5_block15_1_bn (BatchNormal	(None, None, None, 1 512	conv5_block15_1_conv[0][0]
conv5_block15_1_relu (Activatio	(None, None, None, 1 0	conv5_block15_1_bn[0][0]
conv5_block15_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block15_1_relu[0][0]
conv5_block15_concat (Concatena	(None, None, None, 9 0	conv5_block14_concat[0][0] conv5_block15_2_conv[0][0]
conv5_block16_0_bn (BatchNormal	(None, None, None, 9 3968	conv5_block15_concat[0][0]
conv5_block16_0_relu (Activatio	(None, None, None, 9 0	conv5_block16_0_bn[0][0]
conv5_block16_1_conv (Conv2D)	(None, None, None, 1 126976	conv5_block16_0_relu[0][0]
conv5_block16_1_bn (BatchNormal	(None, None, None, 1 512	conv5_block16_1_conv[0][0]
conv5_block16_1_relu (Activatio	(None, None, None, 1 0	conv5_block16_1_bn[0][0]
conv5_block16_2_conv (Conv2D)	(None, None, None, 3 36864	conv5_block16_1_relu[0][0]
conv5_block16_concat (Concatena	(None, None, None, 1 0	conv5_block15_concat[0][0] conv5_block16_2_conv[0][0]
bn (BatchNormalization)	(None, None, None, 1 4096	conv5_block16_concat[0][0]
global_average_pooling2d_1 (Glo	(None, 1024) 0	bn[0][0]


```
-----
dense_1 (Dense)                (None, 14)                14350                global_average_pooling2d_1[0]
=====
```

Total params: 7,051,854

Trainable params: 6,968,206

Non-trainable params: 83,648

```
-----
```

There are a lot of layers, but typically we'll only be extracting one of the last few. Remember that the last few layers usually have more abstract information. To access a layer, we can use `model.get_layer(layer).output`, which takes in the name of the layer in question. Let's try getting the `conv5_block16_concat` layer, the raw output of the last convolutional layer.

```
In [7]: spatial_maps = model.get_layer('conv5_block16_concat').output
        print(spatial_maps)
```

```
Tensor("conv5_block16_concat/concat:0", shape=(?, ?, ?, 1024), dtype=float32)
```

Now, this tensor is just a placeholder, it doesn't contain the actual activations for a particular image. To get this we will use [Keras.backend.function](#) to return intermediate computations while the model is processing a particular input. This method takes in an input and output placeholders and returns a function. This function will compute the intermediate output (until it reaches the given placeholder) evaluated given the input. For example, if you want the layer that you just retrieved (`conv5_block16_concat`), you could write the following:

```
In [15]: model.input
```

```
Out[15]: <tf.Tensor 'input_1:0' shape=(?, ?, ?, 3) dtype=float32>
```

```
In [16]: get_spatial_maps = K.function([model.input], [spatial_maps])
        print(get_spatial_maps)
```

```
<keras.backend.tensorflow_backend.Function object at 0x7fee5915f320>
```

We see that we now have a Function object. Now, to get the actual intermediate output evaluated with a particular input, we just plug in an image to this function:

```
In [17]: # get an image
        x = load_image_normalize(im_path, mean, std)
        print(f"x is of type {type(x)}")
        print(f"x is of shape {x.shape}")
```

```
x is of type <class 'numpy.ndarray'>
```

```
x is of shape (1, 320, 320, 3)
```

```
In [18]: # get the spatial maps layer activations (a list of numpy arrays)
         spatial_maps_x_l = get_spatial_maps([x])
```

```
print(f"spatial_maps_x_l is of type {type(spatial_maps_x_l)}")
print(f"spatial_maps_x_l is has length {len(spatial_maps_x_l)}")
```

```
spatial_maps_x_l is of type <class 'list'>
spatial_maps_x_l is has length 1
```

```
In [19]: # get the 0th item in the list
```

```
spatial_maps_x = spatial_maps_x_l[0]
print(f"spatial_maps_x is of type {type(spatial_maps_x)}")
print(f"spatial_maps_x is of shape {spatial_maps_x.shape}")
```

```
spatial_maps_x is of type <class 'numpy.ndarray'>
spatial_maps_x is of shape (1, 10, 10, 1024)
```

Notice that the shape is (1, 10, 10, 1024). The 0th dimension of size 1 is the batch dimension. Remove the batch dimension for later calculations by taking the 0th index of `spatial_maps_x`.

```
In [20]: # Get rid of the batch dimension
```

```
spatial_maps_x = spatial_maps_x[0] # equivalent to spatial_maps_x[0,:]
print(f"spatial_maps_x without the batch dimension has shape {spatial_maps_x.shape}")
print("Output some of the content:")
print(spatial_maps_x[0])
```

```
spatial_maps_x without the batch dimension has shape (10, 10, 1024)
```

```
Output some of the content:
```

```
[[-0.40058142  0.2602039 -0.6559637 ...  0.14879027 -0.07433726
   0.1822857 ]
 [-0.7217703 -0.21797124 -0.9300643 ...  0.24638593 -0.11097135
   0.30256408]
 [-0.41521266 -0.05806366 -0.6248067 ...  0.30840862 -0.13081692
   0.35475287]
 ...
 [-0.2644133 -0.41372856 -1.2155564 ...  0.18217763 -0.08790095
   0.25714245]
 [-0.29200602 -0.28028086 -0.7485931 ...  0.23563018 -0.12183944
   0.33535504]
 [-0.17051847  0.53123987 -0.2356185 ...  0.10086929 -0.05561769
   0.19629744]]
```

We now have the activations for that particular image, and we can use it for interpretation. The function that is returned by calling `K.function([model.input], [spatial_maps])` (saved here in the variable `get_spatial_maps`) is sometimes referred to as a “hook”, letting you peek into the intermediate computations in the model.

1.1.2 Getting Gradients

The other major step in computing GradCAMs is getting gradients with respect to the output for a particular class. Luckily, Keras makes getting gradients simple. We can use the [Keras.backend.gradients](#) function. The first parameter is the value you are taking the gradient of, and the second is the parameter you are taking that gradient with respect to. We illustrate below:

```
In [21]: # get the output of the model
        output_with_batch_dim = model.output
        print(f"Model output includes batch dimension, has shape {output_with_batch_dim.shape}")
```

Model output includes batch dimension, has shape (?, 14)

To get the output without the batch dimension, you can take the 0th index of the tensor. Note that because the batch dimension is 'None', you could actually enter any integer index, but let's just use 0.

```
In [22]: # Get the output without the batch dimension
        output_all_categories = output_with_batch_dim[0]
        print(f"The output for all 14 categories of disease has shape {output_all_categories.shape}")
```

The output for all 14 categories of disease has shape (14,)

The output has 14 categories, one for each disease category, indexed from 0 to 13. Cardiomegaly is the disease category at index 0.

```
In [23]: # Get the first category's output (Cardiomegaly) at index 0
        y_category_0 = output_all_categories[0]
        print(f"The Cardiomegaly output is at index 0, and has shape {y_category_0.shape}")
```

The Cardiomegaly output is at index 0, and has shape ()

```
In [24]: # Get gradient of y_category_0 with respect to spatial_maps
```

```
        gradient_l = K.gradients(y_category_0, spatial_maps)
        print(f"gradient_l is of type {type(gradient_l)} and has length {len(gradient_l)}")

        # gradient_l is a list of size 1. Get the gradient at index 0
        gradient = gradient_l[0]
        print(gradient)
```

gradient_l is of type <class 'list'> and has length 1
Tensor("gradients/AddN:0", shape=(?, ?, ?, 1024), dtype=float32)

Again, this is just a placeholder. Just like for intermediate layers, we can use `K.function` to compute the value of the gradient for a particular input.

The `K.function()` takes in - a list of inputs: in this case, one input, 'model.input' - a list of tensors: in this case, one output tensor 'gradient'

It returns a function that calculates the activations of the list of tensors. - This returned function returns a list of the activations, one for each tensor that was passed into `K.function()`.

```
In [25]: # Create the function that gets the gradient
get_gradient = K.function([model.input], [gradient])
type(get_gradient)
```

```
Out[25]: keras.backend.tensorflow_backend.Function
```

```
In [26]: # get an input x-ray image
x = load_image_normalize(im_path, mean, std)
print(f"X-ray image has shape {x.shape}")
```

```
X-ray image has shape (1, 320, 320, 3)
```

The `get_gradient` function takes in a list of inputs, and returns a list of the gradients, one for each image.

```
In [27]: # use the get_gradient function to get the gradient (pass in the input image inside a
grad_x_l = get_gradient([x])
print(f"grad_x_l is of type {type(grad_x_l)} and length {len(grad_x_l)}")

# get the gradient at index 0 of the list.
grad_x_with_batch_dim = grad_x_l[0]
print(f"grad_x_with_batch_dim is type {type(grad_x_with_batch_dim)} and shape {grad_x_

# To remove the batch dimension, take the value at index 0 of the batch dimension
grad_x = grad_x_with_batch_dim[0]
print(f"grad_x is type {type(grad_x)} and shape {grad_x.shape}")

print("Gradient grad_x (show some of its content:")
print(grad_x[0])
```

```
grad_x_l is of type <class 'list'> and length 1
grad_x_with_batch_dim is type <class 'numpy.ndarray'> and shape (1, 10, 10, 1024)
grad_x is type <class 'numpy.ndarray'> and shape (10, 10, 1024)
Gradient grad_x (show some of its content:
[[-1.3060314e-09  2.6313329e-09  3.0835795e-07 ...  8.6101973e-05
  -5.7629448e-05  6.0046797e-05]
 [-1.3060314e-09  2.6313329e-09  3.0835795e-07 ...  8.6101973e-05
  -5.7629448e-05  6.0046797e-05]
 [-1.3060314e-09  2.6313329e-09  3.0835795e-07 ...  8.6101973e-05
  -5.7629448e-05  6.0046797e-05]
 ...
```

```
[-1.3060314e-09  2.6313329e-09  3.0835795e-07 ...  8.6101973e-05
 -5.7629448e-05  6.0046797e-05]
[-1.3060314e-09  2.6313329e-09  3.0835795e-07 ...  8.6101973e-05
 -5.7629448e-05  6.0046797e-05]
[-1.3060314e-09  2.6313329e-09  3.0835795e-07 ...  8.6101973e-05
 -5.7629448e-05  6.0046797e-05]]
```

Just like we had a hook into the penultimate layer, we now have a hook into the gradient! This allows us to easily compute pretty much anything relevant to our model output.

We can also combine the two to have one function call which gives us both the gradient and the last layer (this might come in handy when implementing GradCAM in the next section).

```
In [28]: # Use K.function to generate a single function
        # Notice that a list of two tensors, is passed in as the second argument of K.function
        get_spatial_maps_and_gradient = K.function([model.input], [spatial_maps, gradient])
        print(type(get_spatial_maps_and_gradient))
```

```
<class 'keras.backend.tensorflow_backend.Function'>
```

```
In [29]: # The returned function returns a list of the evaluated tensors
        tensor_eval_l = get_spatial_maps_and_gradient([x])
        print(f"tensor_eval_l is type {type(tensor_eval_l)} and length {len(tensor_eval_l)}")
```

```
tensor_eval_l is type <class 'list'> and length 2
```

```
In [30]: # store the two numpy arrays from index 0 and 1 into their own variables
        spatial_maps_x_with_batch_dim, grad_x_with_batch_dim = tensor_eval_l
        print(f"spatial_maps_x_with_batch_dim has shape {spatial_maps_x_with_batch_dim.shape}")
        print(f"grad_x_with_batch_dim has shape {grad_x_with_batch_dim.shape}")
```

```
spatial_maps_x_with_batch_dim has shape (1, 10, 10, 1024)
```

```
grad_x_with_batch_dim has shape (1, 10, 10, 1024)
```

```
In [31]: # Note: you could also do this directly from the function call:
        spatial_maps_x_with_batch_dim, grad_x_with_batch_dim = get_spatial_maps_and_gradient([x])
        print(f"spatial_maps_x_with_batch_dim has shape {spatial_maps_x_with_batch_dim.shape}")
        print(f"grad_x_with_batch_dim has shape {grad_x_with_batch_dim.shape}")
```

```
spatial_maps_x_with_batch_dim has shape (1, 10, 10, 1024)
```

```
grad_x_with_batch_dim has shape (1, 10, 10, 1024)
```

```
In [32]: # Remove the batch dimension by taking the 0th index at the batch dimension
        spatial_maps_x = spatial_maps_x_with_batch_dim[0]
        grad_x = grad_x_with_batch_dim[0]
```

```

print(f"spatial_maps_x shape {spatial_maps_x.shape}")
print(f"grad_x shape {grad_x.shape}")

print("\nSpatial maps (print some content):")
print(spatial_maps_x[0])
print("\nGradient (print some content:")
print(grad_x[0])

```

```

spatial_maps_x shape (10, 10, 1024)
grad_x shape (10, 10, 1024)

```

```

Spatial maps (print some content):
[[-0.40058142  0.2602039 -0.6559637 ...  0.14879027 -0.07433726
  0.1822857 ]
 [-0.7217703 -0.21797124 -0.9300643 ...  0.24638593 -0.11097135
  0.30256408]
 [-0.41521266 -0.05806366 -0.6248067 ...  0.30840862 -0.13081692
  0.35475287]
 ...
 [-0.2644133 -0.41372856 -1.2155564 ...  0.18217763 -0.08790095
  0.25714245]
 [-0.29200602 -0.28028086 -0.7485931 ...  0.23563018 -0.12183944
  0.33535504]
 [-0.17051847  0.53123987 -0.2356185 ...  0.10086929 -0.05561769
  0.19629744]]

```

```

Gradient (print some content:
[[-1.3060314e-09  2.6313329e-09  3.0835795e-07 ...  8.6101973e-05
 -5.7629448e-05  6.0046797e-05]
 [-1.3060314e-09  2.6313329e-09  3.0835795e-07 ...  8.6101973e-05
 -5.7629448e-05  6.0046797e-05]
 [-1.3060314e-09  2.6313329e-09  3.0835795e-07 ...  8.6101973e-05
 -5.7629448e-05  6.0046797e-05]
 ...
 [-1.3060314e-09  2.6313329e-09  3.0835795e-07 ...  8.6101973e-05
 -5.7629448e-05  6.0046797e-05]
 [-1.3060314e-09  2.6313329e-09  3.0835795e-07 ...  8.6101973e-05
 -5.7629448e-05  6.0046797e-05]
 [-1.3060314e-09  2.6313329e-09  3.0835795e-07 ...  8.6101973e-05
 -5.7629448e-05  6.0046797e-05]]

```

1.1.3 Implementing GradCAM

Exercise 1

In the next cell, fill in the `grad_cam` method to produce GradCAM visualizations for an input model and image. This is fairly complicated, so it might help to break it down into these steps:

1. Hook into model output and last layer activations.

2. Get gradients of last layer activations with respect to output.
3. Compute value of last layer and gradients for input image.
4. Compute weights from gradients by global average pooling.
5. Compute the dot product between the last layer and weights to get the score for each pixel.
6. Resize, take ReLU, and return cam.

Hints

The following hints follow the order of the sections described above. 1. Remember that the output shape of our model will be [1, class_amount]. 1. The input in this case will always have batch_size = 1 2. See [K.gradients](#) 3. Follow the procedure we used in the previous two sections. 4. Check the axis; make sure weights have shape (C)! 5. See [np.dot](#)

To test, you will compare your output on an image to the output from a correct implementation of GradCAM. You will receive full credit if the pixel-wise mean squared error is less than 0.05.

```
In [35]: # UNQ_C1 (UNIQUE CELL IDENTIFIER, DO NOT EDIT)
def grad_cam(input_model, image, category_index, layer_name):
    """
    GradCAM method for visualizing input saliency.

    Args:
        input_model (Keras.model): model to compute cam for
        image (tensor): input to model, shape (1, H, W, 3)
        cls (int): class to compute cam with respect to
        layer_name (str): relevant layer in model
        H (int): input height
        W (int): input width
    Return:
        cam ()
    """
    cam = None

    ### START CODE HERE (REPLACE INSTANCES OF 'None' with your code) ###

    # 1. Get placeholders for class output and last layer
    # Get the model's output
    output_with_batch_dim = input_model.output

    # Remove the batch dimension
    output_all_categories = output_with_batch_dim[0]

    # Retrieve only the disease category at the given category index
    y_c = output_all_categories[category_index]

    # Get the input model's layer specified by layer_name, and retrieve the layer's output
    spatial_map_layer = model.get_layer(layer_name).output

    # 2. Get gradients of last layer with respect to output
```

```

# get the gradients of y_c with respect to the spatial map layer (it's a list of
grads_l = K.gradients(y_c, spatial_map_layer)

# Get the gradient at index 0 of the list
grads = grads_l[0]

# 3. Get hook for the selected layer and its gradient, based on given model's inp
# Hint: Use the variables produced by the previous two lines of code
spatial_map_and_gradient_function = K.function([input_model.input], [spatial_maps

# Put in the image to calculate the values of the spatial_maps (selected layer) a
spatial_map_all_dims, grads_val_all_dims = spatial_map_and_gradient_function([ima

# Reshape activations and gradient to remove the batch dimension
# Shape goes from (B, H, W, C) to (H, W, C)
# B: Batch. H: Height. W: Width. C: Channel
# Reshape spatial map output to remove the batch dimension
spatial_map_val = spatial_map_all_dims[0]

# Reshape gradients to remove the batch dimension
grads_val = grads_val_all_dims[0]

# 4. Compute weights using global average pooling on gradient
# grads_val has shape (Height, Width, Channels) (H,W,C)
# Take the mean across the height and also width, for each channel
# Make sure weights have shape (C)
weights = np.mean(grads_val,axis=(0,1))

# 5. Compute dot product of spatial map values with the weights
cam = np.dot(spatial_map_val,weights)

### END CODE HERE ###

# We'll take care of the postprocessing.
H, W = image.shape[1], image.shape[2]
cam = np.maximum(cam, 0) # ReLU so we only get positive importance
cam = cv2.resize(cam, (W, H), cv2.INTER_NEAREST)
cam = cam / cam.max()

return cam

```

Below we generate the CAM for the image and compute the error (pixel-wise mean squared difference) from the expected values according to our reference.

```

In [36]: im = load_image_normalize(im_path, mean, std)
cam = grad_cam(model, im, 5, 'conv5_block16_concat') # Mass is class 5

# Loads reference CAM to compare our implementation with.

```



```
reference = np.load("reference_cam.npy")
error = np.mean((cam-reference)**2)

print(f"Error from reference: {error:.4f}, should be less than 0.05")
```

Error from reference: 0.0606, should be less than 0.05

Run the next cell to visualize the CAM and the original image.

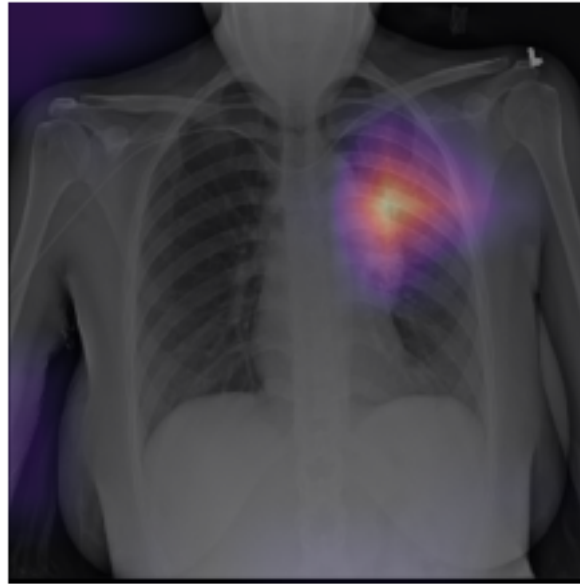
```
In [37]: plt.imshow(load_image(im_path, df, preprocess=False), cmap='gray')
plt.title("Original")
plt.axis('off')

plt.show()

plt.imshow(load_image(im_path, df, preprocess=False), cmap='gray')
plt.imshow(cam, cmap='magma', alpha=0.5)
plt.title("GradCAM")
plt.axis('off')
plt.show()
```



GradCAM



We can see that it focuses on the large (white) empty area on the right lung. Indeed this is a clear case of Mass.

1.1.4 Using GradCAM to Visualize Multiple Labels

Exercise 2

We can use GradCAMs for multiple labels on the same image. Let's do it for the labels with best AUC for our model, Cardiomegaly, Mass, and Edema.

```
In [50]: # UNQ_C2 (UNIQUE CELL IDENTIFIER, DO NOT EDIT)
def compute_gradcam(model, img, mean, std, data_dir, df,
                    labels, selected_labels, layer_name='conv5_block16_concat'):
    """
    Compute GradCAM for many specified labels for an image.
    This method will use the `grad_cam` function.

    Args:
        model (Keras.model): Model to compute GradCAM for
        img (string): Image name we want to compute GradCAM for.
        mean (float): Mean to normalize to image.
        std (float): Standard deviation to normalize the image.
        data_dir (str): Path of the directory to load the images from.
        df(pd.DataFrame): Dataframe with the image features.
        labels ([str]): All output labels for the model.
        selected_labels ([str]): All output labels we want to compute the GradCAM for
        layer_name: Intermediate layer from the model we want to compute the GradCAM for.
    """
    img_path = data_dir + img
    preprocessed_input = load_image_normalize(img_path, mean, std)
```

```

predictions = model.predict(preprocessed_input)
print("Ground Truth: ", " ", ".join(np.take(labels, np.nonzero(df[df["Image"] == im

plt.figure(figsize=(15, 10))
plt.subplot(151)
plt.title("Original")
plt.axis('off')
plt.imshow(load_image(img_path, df, preprocess=False), cmap='gray')

j = 1

### START CODE HERE (REPLACE INSTANCES OF 'None' with your code) ###
# Loop through all labels
for i in range(len(labels)): # complete this line
    # Compute CAM and show plots for each selected label.

    # Check if the label is one of the selected labels
    if labels[i] in selected_labels: # complete this line

        # Use the grad_cam function to calculate gradcam
        gradcam = grad_cam(model, preprocessed_input, i, layer_name)

    ### END CODE HERE ###

    print("Generating gradcam for class %s (p=%2.2f)" % (labels[i], round(pre
plt.subplot(151 + j)
plt.title(labels[i] + ": " + str(round(predictions[0][i], 3)))
plt.axis('off')
plt.imshow(load_image(img_path, df, preprocess=False), cmap='gray')
plt.imshow(gradcam, cmap='magma', alpha=min(0.5, predictions[0][i]))
j +=1

```

Run the following cells to print the ground truth diagnosis for a given case and show the original x-ray as well as GradCAMs for Cardiomegaly, Mass, and Edema.

```

In [51]: df = pd.read_csv("nih_new/train-small.csv")

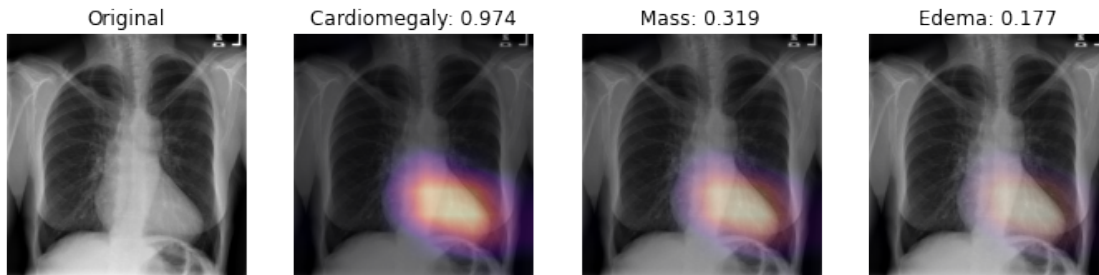
image_filename = '00016650_000.png'
labels_to_show = ['Cardiomegaly', 'Mass', 'Edema']
compute_gradcam(model, image_filename, mean, std, IMAGE_DIR, df, labels, labels_to_sho

```

```

Ground Truth: Cardiomegaly
Generating gradcam for class Cardiomegaly (p=0.97)
Generating gradcam for class Mass (p=0.32)
Generating gradcam for class Edema (p=0.18)

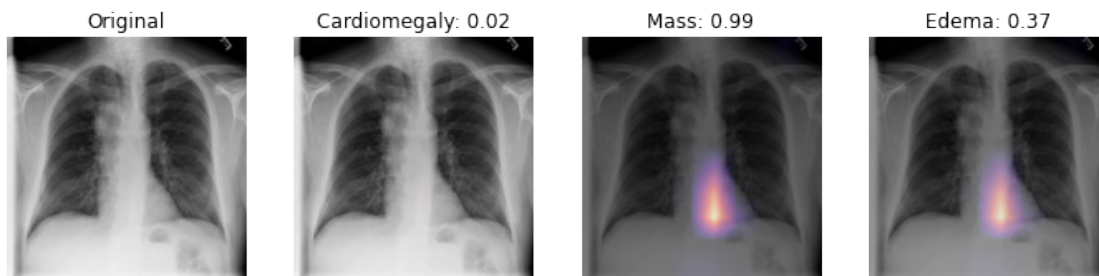
```



The model correctly predicts absence of mass or edema. The probability for mass is higher, and we can see that it may be influenced by the shapes in the middle of the chest cavity, as well as around the shoulder. We'll run it for two more images.

```
In [52]: image_filename = '00005410_000.png'
         compute_gradcam(model, image_filename, mean, std, IMAGE_DIR, df, labels, labels_to_show)
```

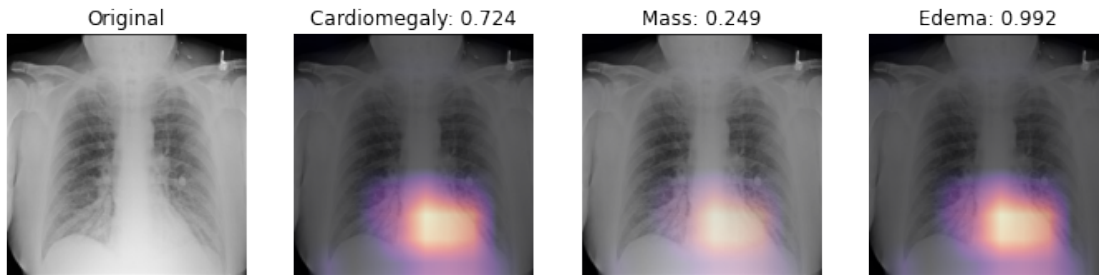
```
Ground Truth:  Mass
Generating gradcam for class Cardiomegaly (p=0.02)
Generating gradcam for class Mass (p=0.99)
Generating gradcam for class Edema (p=0.37)
```



In the example above, the model correctly focuses on the mass near the center of the chest cavity.

```
In [53]: image_name = '00004090_002.png'
         compute_gradcam(model, image_name, mean, std, IMAGE_DIR, df, labels, labels_to_show)
```

```
Ground Truth:  Edema
Generating gradcam for class Cardiomegaly (p=0.72)
Generating gradcam for class Mass (p=0.25)
Generating gradcam for class Edema (p=0.99)
```



Here the model correctly picks up the signs of edema near the bottom of the chest cavity. We can also notice that Cardiomegaly has a high score for this image, though the ground truth doesn't include it. This visualization might be helpful for error analysis; for example, we can notice that the model is indeed looking at the expected area to make the prediction.

This concludes the section on GradCAMs. We hope you've gained an appreciation for the importance of interpretation when it comes to deep learning models in medicine. Interpretation tools like this one can be helpful for discovery of markers, error analysis, and even in deployment.

2 Feature Importance in Machine Learning

When developing predictive models and risk measures, it's often helpful to know which features are making the most difference. This is easy to determine in simpler models such as linear models and decision trees. However as we move to more complex models to achieve high performance, we usually sacrifice some interpretability. In this assignment we'll try to regain some of that interpretability using Shapley values, a technique which has gained popularity in recent years, but which is based on classic results in cooperative game theory.

We'll revisit our random forest model from course 2 module 2 and try to analyze it more closely using Shapley values. Run the next cell to load in the data and model from that assignment and recalculate the test set c-index.

```
In [54]: rf = pickle.load(open('nhanes_rf.sav', 'rb')) # Loading the model
test_df = pd.read_csv('nhanestest.csv')
test_df = test_df.drop(test_df.columns[0], axis=1)
X_test = test_df.drop('y', axis=1)
y_test = test_df.loc[:, 'y']
cindex_test = cindex(y_test, rf.predict_proba(X_test)[:, 1])

print("Model C-index on test: {}".format(cindex_test))
```

Model C-index on test: 0.7776169781865744

Run the next cell to print out the riskiest individuals according to our model.

```
In [55]: X_test_risky = X_test.copy(deep=True)
X_test_risky.loc[:, 'risk'] = rf.predict_proba(X_test)[:, 1] # Predicting our risk.
X_test_risky = X_test_risky.sort_values(by='risk', ascending=False) # Sorting by risk
X_test_risky.head()
```

```

Out [55]:      Age  Diastolic BP  Poverty index  Race  Red blood cells  \
572   70.0         80.0         312.0   1.0         54.8
190   69.0        100.0         316.0   1.0         77.7
1300  73.0         80.0         999.0   1.0         52.6
634   66.0        100.0          69.0   2.0         42.9
1221  74.0         80.0          67.0   1.0         40.3

      Sedimentation rate  Serum Albumin  Serum Cholesterol  Serum Iron  \
572                   7.0           4.4           222.0        52.0
190                   26.0           4.2           197.0        65.0
1300                   35.0           3.9           258.0        61.0
634                   47.0           3.8           233.0       170.0
1221                   24.0           3.7           139.0        28.0

      Serum Magnesium  Serum Protein  Sex  Systolic BP  TIBC  TS  \
572                1.57           7.2  1.0        180.0  417.0  12.5
190                1.49           7.5  1.0        165.0  298.0  21.8
1300                1.66           6.8  1.0        150.0  314.0  19.4
634                1.42           8.6  1.0        180.0  411.0  41.4
1221                1.91           6.4  2.0        140.0  495.0   5.7

      White blood cells      BMI  Pulse pressure  risk
572                7.5  45.770473        100.0  0.77
190                8.8  22.129018         65.0  0.69
1300                9.4  26.466850         70.0  0.69
634                7.2  22.129498         80.0  0.68
1221                4.1  22.066389         60.0  0.68

```

2.1 Permutation Method for Feature Importance

First we'll try to determine feature importance using the permutation method. In the permutation method, the importance of feature i would be the regular performance of the model minus the performance with the values for feature i permuted in the dataset. This way we can assess how well a model without that feature would do without having to train a new model for each feature.

2.1.1 Implementing Permutation

Exercise 3

Complete the implementation of the function below, which given a feature name returns a dataset with those feature values randomly permuted.

Hints

See `np.random.permutation`

```

In [62]: # UNQ_C3 (UNIQUE CELL IDENTIFIER, DO NOT EDIT)
def permute_feature(df, feature):
    """
    Given dataset, returns version with the values of
    the given feature randomly permuted.

    Args:

```

```

    df (dataframe): The dataset, shape (num subjects, num features)
    feature (string): Name of feature to permute
Returns:
    permuted_df (dataframe): Exactly the same as df except the values
                             of the given feature are randomly permuted.
"""
permuted_df = df.copy(deep=True) # Make copy so we don't change original df

### START CODE HERE (REPLACE INSTANCES OF 'None' with your code) ###

# Permute the values of the column 'feature'
permuted_features = np.random.permutation(permuted_df[feature])

# Set the column 'feature' to its permuted values.
permuted_df[feature] = permuted_features

### END CODE HERE ###

return permuted_df

```

```
In [63]: print("Test Case")
```

```

example_df = pd.DataFrame({'col1': [0, 1, 2], 'col2': ['A', 'B', 'C']})
print("Original dataframe:")
print(example_df)
print("\n")

print("col1 permuted:")
print(permute_feature(example_df, 'col1'))

print("\n")
print("Compute average values over 1000 runs to get expected values:")
col1_values = np.zeros((3, 1000))
np.random.seed(0) # Adding a constant seed so we can always expect the same values and
for i in range(1000):
    col1_values[:, i] = permute_feature(example_df, 'col1')['col1'].values

print("Average of col1: {}, expected value: [0.976, 1.03, 0.994]".format(np.mean(col1_values),

```

Test Case

Original dataframe:

	col1	col2
0	0	A
1	1	B
2	2	C

col1 permuted:

	col1	col2	
0	1	A	
1	0	B	
2	2	C	

Compute average values over 1000 runs to get expected values:

Average of col1: [0.976 1.03 0.994], expected value: [0.976, 1.03, 0.994]

2.1.2 Implementing Importance

Exercise 4

Now we will use the function we just created to compute feature importances (according to the permutation method) in the function below.

Hints

$$I_x = |perf - perf_x| \quad (1)$$

where I_x is the importance of feature x and

$$perf_x = \frac{1}{n} \cdot \sum_{i=1}^n perf_i^{sx} \quad (2)$$

where $perf_i^{sx}$ is the performance with the feature x shuffled in the i th permutation.

In [70]: # UNQ_C4 (UNIQUE CELL IDENTIFIER, DO NOT EDIT)

```
def permutation_importance(X, y, model, metric, num_samples = 100):
    """
```

Compute permutation importance for each feature.

Args:

X (dataframe): Dataframe for test data, shape (num subject, num features)

y (np.array): Labels for each row of X, shape (num subjects,)

model (object): Model to compute importances for, guaranteed to have a 'predict_proba' method to compute probabilistic predictions given input

metric (function): Metric to be used for feature importance. Takes in ground truth and predictions as the only two arguments

num_samples (int): Number of samples to average over when computing change in performance for each feature

Returns:

importances (dataframe): Dataframe containing feature importance for each column of df with shape (1, num_features)

```
    """
```

```
importances = pd.DataFrame(index = ['importance'], columns = X.columns)
```

```
# Get baseline performance (note, you'll use this metric function again later)
```

```
baseline_performance = metric(y, model.predict_proba(X)[: , 1])
```



```

### START CODE HERE (REPLACE INSTANCES OF 'None' with your code) ###

# Iterate over features (the columns in the importances dataframe)
for feature in importances.columns: # complete this line

    # Compute 'num_sample' performances by permutating that feature

    # You'll see how the model performs when the feature is permuted
    # You'll do this num_samples number of times, and save the performance each time
    # To store the feature performance,
    # create a numpy array of size num_samples, initialized to all zeros
    feature_performance_arr = np.zeros(num_samples)

    # Loop through each sample
    for i in range(num_samples): # complete this line

        # permute the column of dataframe X
        perm_X = permute_feature(X, feature)

        # calculate the performance with the permuted data
        # Use the same metric function that was used earlier
        feature_performance_arr[i] = metric(y, model.predict_proba(perm_X)[:, 1])

    # Compute importance: absolute difference between
    # the baseline performance and the average across the feature performance
    importances[feature]['importance'] = baseline_performance - np.mean(feature_performance_arr)

### END CODE HERE ###

return importances

```

Test Case

```

In [71]: print("Test Case")
         print("\n")
         print("We check our answers on a Logistic Regression on a dataset")
         print("where y is given by a sigmoid applied to the important feature.")
         print("The unimportant feature is random noise.")
         print("\n")
         example_df = pd.DataFrame({'important': np.random.normal(size=(1000)), 'unimportant': np.random.normal(size=(1000))})
         example_y = np.round(1 / (1 + np.exp(-example_df.important)))
         example_model = sklearn.linear_model.LogisticRegression(fit_intercept=False).fit(example_df, example_y)

         example_importances = permutation_importance(example_df, example_y, example_model, ci=0.05)
         print("Computed importances:")
         print(example_importances)

```

```

print("\n")
print("Expected importances (approximate values):")
print(pd.DataFrame({"important": 0.50, "unimportant": 0.00}, index=['importance']))
print("If you round the actual values, they will be similar to the expected values")

```

Test Case

We check our answers on a Logistic Regression on a dataset where y is given by a sigmoid applied to the important feature. The unimportant feature is random noise.

Computed importances:

```

            important  unimportant
importance  0.500501  8.40165e-05

```

Expected importances (approximate values):

```

            important  unimportant
importance          0.5          0.0

```

If you round the actual values, they will be similar to the expected values

2.1.3 Computing our Feature Importance

Next, we compute importances on our dataset. Since we are computing the permutation importance for all the features, it might take a few minutes to run.

```

In [72]: importances = permutation_importance(X_test, y_test, rf, cindex, num_samples=100)
importances

```

```

Out[72]:
            Age Diastolic BP Poverty index          Race Red blood cells \
importance  0.147385    0.0114287    0.0109559  1.87334e-05    0.00107613

            Sedimentation rate Serum Albumin Serum Cholesterol  Serum Iron \
importance          0.00549156    0.00476165    -0.000836536  0.000241709

            Serum Magnesium Serum Protein          Sex Systolic BP          TIBC \
importance          0.00264893   -0.00167158  0.0268127  0.00600149  0.0019791

            TS White blood cells          BMI Pulse pressure
importance  0.000174254          0.00264314  0.00269314    0.00401883

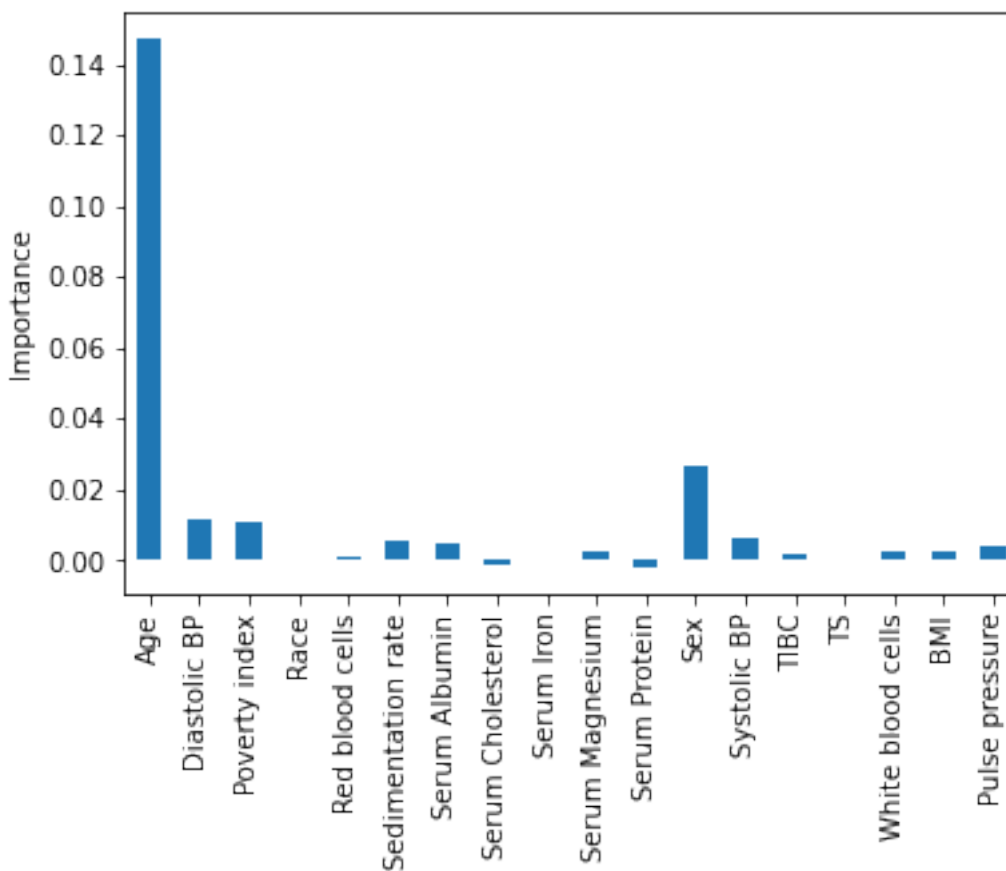
```

Let's plot these in a bar chart for easier comparison.

```

In [73]: importances.T.plot.bar()
plt.ylabel("Importance")
l = plt.legend()
l.remove()
plt.show()

```



You should see age as by far the best prediction of near term mortality, as one might expect. Next is sex, followed by diastolic blood pressure. Interestingly, the poverty index also has a large impact, despite the fact that it is not directly related to an individual's health. This alludes to the importance of social determinants of health in our model.

2.2 Shapley Values for Random Forests

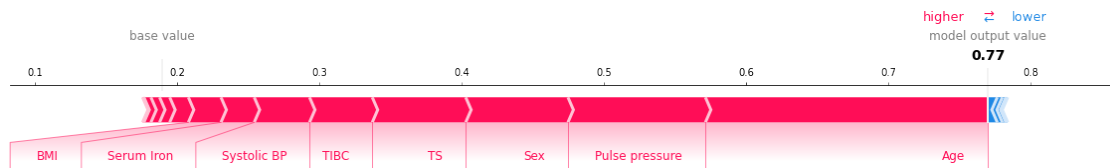
We'll contrast the permutation method with a more recent technique known as Shapley values (actually, Shapley values date back to the mid 20th century, but have only been applied to machine learning very recently).

2.2.1 Visualizing Feature Importance on Specific Individuals

We can use Shapley values to try and understand the model output on specific individuals. In general Shapley values take exponential time to compute, but luckily there are faster approximations for forests in particular that run in polynomial time. Run the next cell to display a 'force plot' showing how each feature influences the output for the first person in our dataset. If you want more information about 'force plots' and other decision plots, please take a look at [this notebook](#) by the shap library creators.

```
In [74]: explainer = shap.TreeExplainer(rf)
         i = 0 # Picking an individual
         shap_value = explainer.shap_values(X_test.loc[X_test_risky.index[i], :])[1]
         shap.force_plot(explainer.expected_value[1], shap_value, feature_names=X_test.columns)
```

Setting `feature_perturbation = "tree_path_dependent"` because no background data was given.



For this individual, their age, pulse pressure, and sex were the biggest contributors to their high risk prediction. Note how shapley values give us greater granularity in our interpretations.

Feel free to change the `i` value above to explore the feature influences for different individuals.

2.2.2 Visualizing Feature Importance on Aggregate

Just like with the permutation method, we might also want to understand model output in aggregate. Shapley values allow us to do this as well. Run the next cell to initialize the shapley values for each example in the test set (this may also take a few minutes).

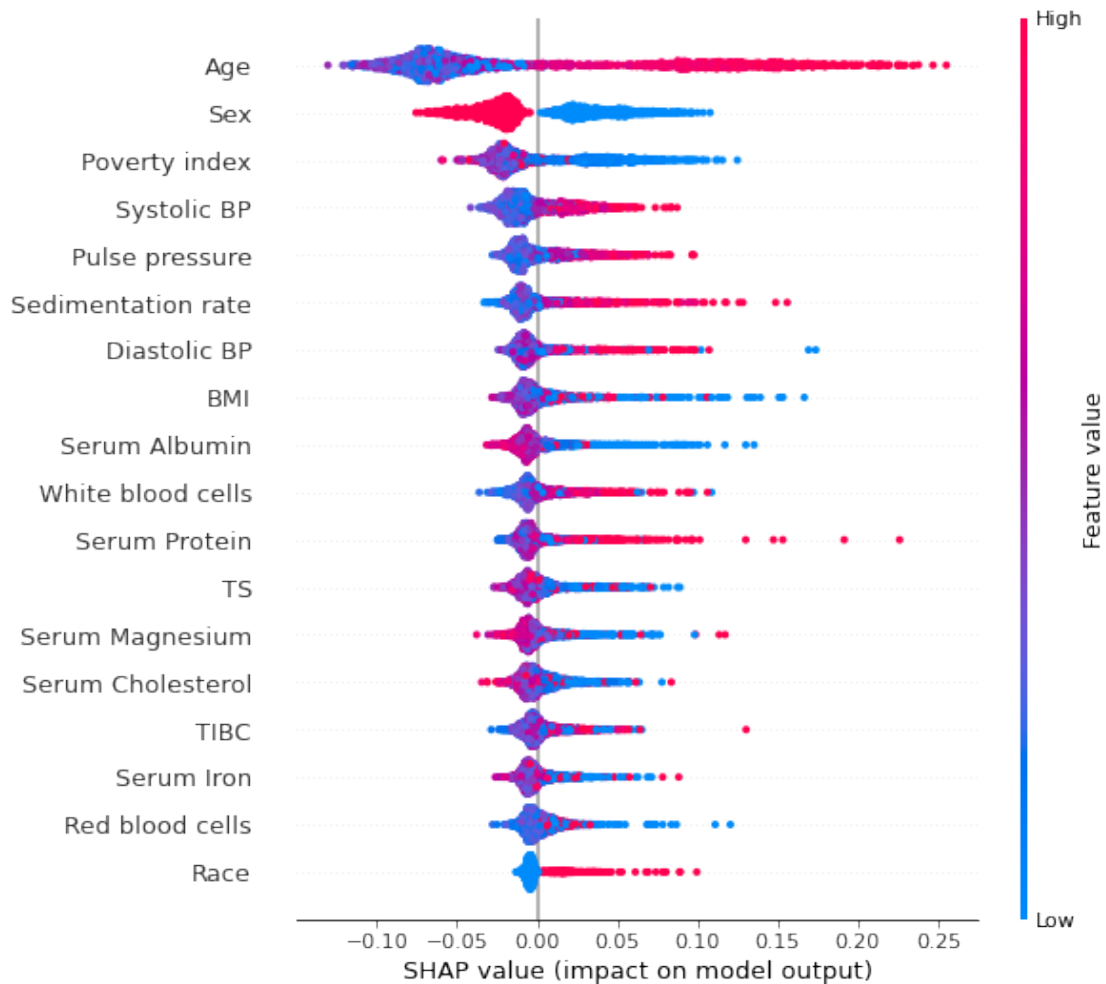
```
In [75]: shap_values = shap.TreeExplainer(rf).shap_values(X_test)[1]
```

Setting `feature_perturbation = "tree_path_dependent"` because no background data was given.

You can ignore the setting `feature_perturbation` message.

Run the next cell to see a summary plot of the shapley values for each feature on each of the test examples. The colors indicate the value of the feature. The features are listed in terms of decreasing absolute average shapley value over all the individuals in the dataset.

```
In [76]: shap.summary_plot(shap_values, X_test)
```



In the above plot, you might be able to notice a high concentration of points on specific SHAP value ranges. This means that a high proportion of our test set lies on those ranges.

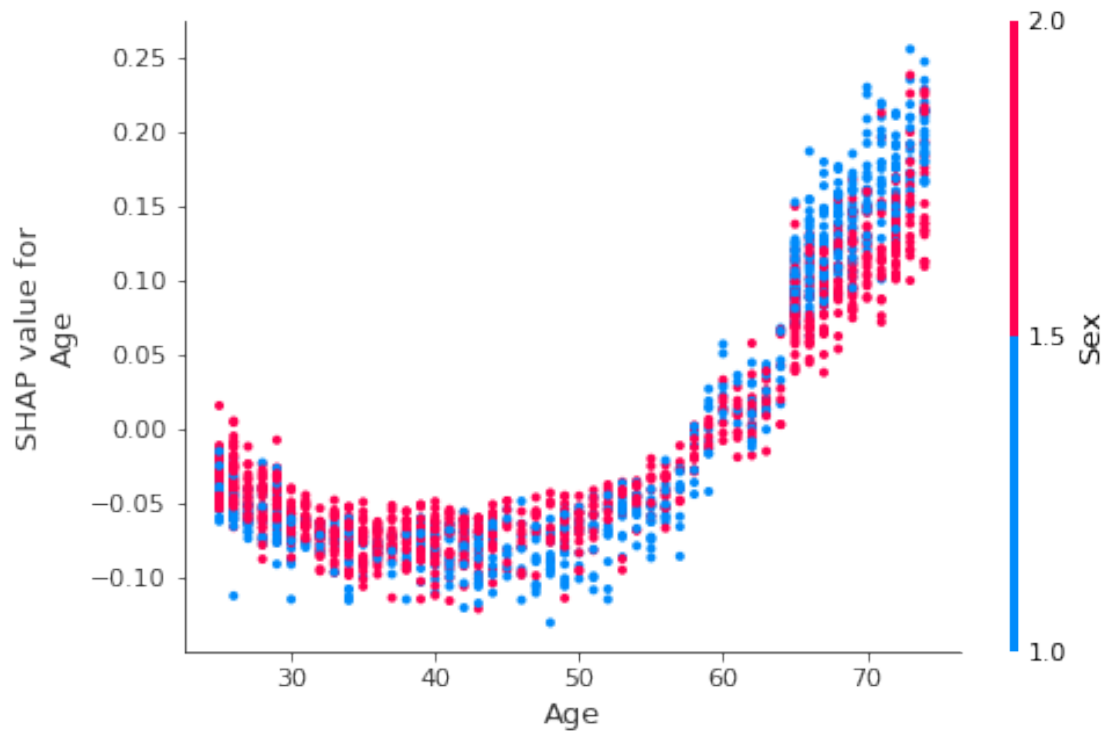
As with the permutation method, age, sex, poverty index, and diastolic BP seem to be the most important features. Being older has a negative impact on mortality, and being a woman (sex=2.0) has a positive effect.

2.2.3 Visualizing Interactions between Features

The shap library also lets you visualize interactions between features using dependence plots. These plot the Shapley value for a given feature for each data point, and color the points in using the value for another feature. This lets us begin to explain the variation in shapley value for a single value of the main feature.

Run the next cell to see the interaction between Age and Sex.

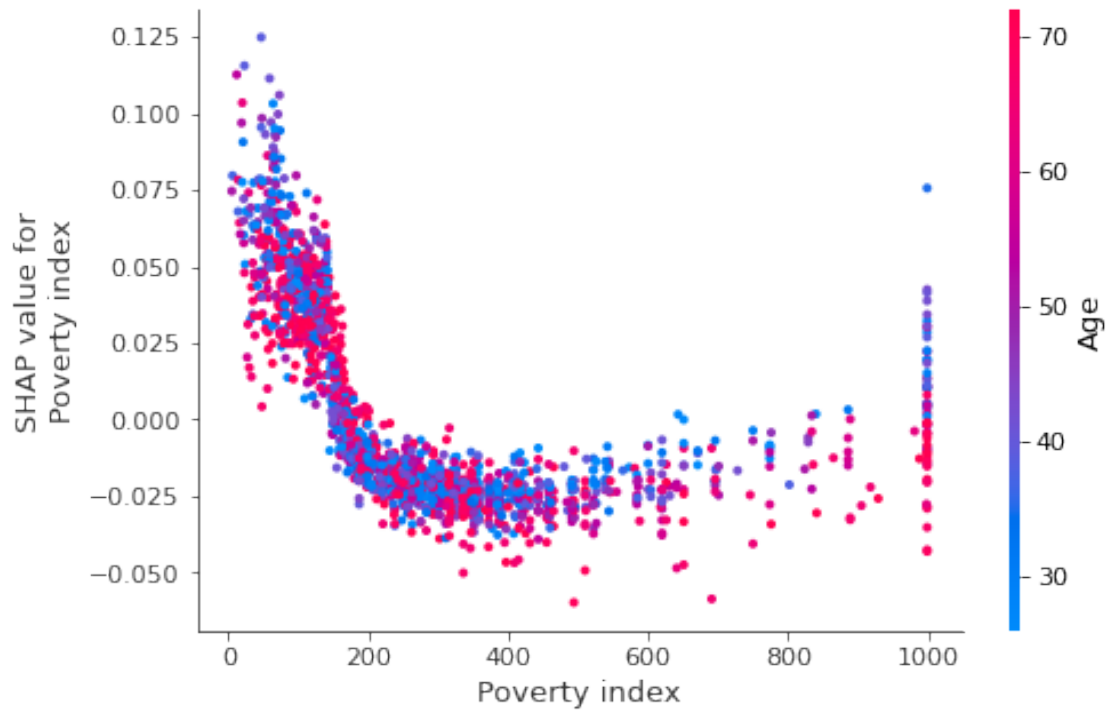
```
In [77]: shap.dependence_plot('Age', shap_values, X_test, interaction_index = 'Sex')
```



We see that while Age > 50 is generally bad (positive Shapley value), being a woman (red points) generally reduces the impact of age. This makes sense since we know that women generally live longer than men.

Run the next cell to see the interaction between Poverty index and Age

```
In [78]: shap.dependence_plot('Poverty index', shap_values, X_test, interaction_index='Age')
```



We see that the impact of poverty index drops off quickly, and for higher income individuals age begins to explain much of variation in the impact of poverty index. We encourage you to try some other pairs and see what other interesting relationships you can find!

Congratulations! You've completed the final assignment of course 3, well done!