Pneumonia Image Classification

Business Understanding:

In this image classification project, I am utilizing neural networks - primarily convolutional neural networks - to create a model that can identify whether or not a patient has pneumonia by analyzing their lung x-rays.

There are two types of pneumonia - bacterial and viral. The image classification system will have to be able to pick out both, while not necessarily knowing which one is which, as the types of pneumonia are not labeled in the dataset.

Data Understanding:

The dataset is organized into 3 folders (train, test, val) and contains subfolders for each image category (Pneumonia/Normal). There are 5,863 X-Ray images (JPEG) and 2 categories (Pneumonia/Normal).

Chest X-ray images (anterior-posterior) were selected from retrospective cohorts of pediatric patients of one to five years old from Guangzhou Women and Children's Medical Center, Guangzhou. All chest X-ray imaging was performed as part of patients' routine clinical care.

Due to the relatively small amount of validation data (16 images), as well as test data, I will create my own validation data instead of using the provided split.

```
In [1]:
        import pandas as pd
        import numpy as np
        import matplotlib.pyplot as plt
        %matplotlib inline
        import keras
        from keras. models import Sequential
        from keras. Layers import Dense, Conv2D, BatchNormalization, MaxPooling2D, D
        ropout, Flatten
        from sklearn.preprocessing import StandardScaler, LabelBinarizer
        import os
        os. envi ron['KMP_DUPLICATE_LIB_OK']='True'
        import tensorflow as tf
        from keras import models
        from keras import layers
        import pathlib
        import PIL
        import seaborn as sns
        import time
        import scipy
        import numpy as np
        from PIL import Image
        from scipy import ndimage
        from sklearn.model_selection import train_test_split, cross_val_score
        from keras.preprocessing.image import ImageDataGenerator, array_to_img, img
        _to_array, load_img
        from sklearn.model_selection import train_test_split, cross_val_score
        from sklearn.preprocessing import OneHotEncoder
        from sklearn.metrics import accuracy_score, confusion_matrix
        import matplotlib.pyplot as plt
        import matplotlib.image as mpimg
        import numpy as np
        import seaborn as sns
        from tensorflow.keras.models import Sequential
        from tensorflow. keras. regularizers import 12
        from tensorflow.keras.optimizers import SGD
        from tensorflow.keras.wrappers import scikit_learn
        from tensorflow. keras. callbacks import EarlyStopping
        import shutil
        import random
        import glob
        import os
        import sys
        import itertools
        import warnings
        import statistics
```

Define Scoring Metric

For the purposes of this analysis, I think a custom scoring metric is necessary.

My reasoning is this:

- False negatives could cause patients with pneumonia to not recieve further testing. This would be the
 worst possibility, out of the options.
- False positives would cost more due to testing people who did not actually have pneumonia, or could cause people without pneumonia to needlessly worry about their health. This is also costly, but not as costly as mis-diagnosing an individual with pneumonia.

Thus, I want to minimize false negatives while keeping false positives to an appropriate level.

According to https://towardsdatascience.com/performance-metrics-confusion-matrix-precision-recall-and-f1-score-a8fe076a2262), the F1 score gives the same weightage to recall and precision, BUT there is a weighted F1 score in which we can give different weightage to recall and precision. As discussed in the previous section, different problems give different weightage to recall and precision.

```
F_B = (1+B^2) [(Precision Recall) / ((B^2 * Precision) + Recall)]
```

Beta represents how many times recall is more important than precision. If the recall is twice as important as precision, the value of Beta is 2.

Thus, for the purpose of this project, my Weighted F1 score will be:

 $F_B = (1+3^2) [(Precision Recall) / ((3^2 * Precision) + Recall)]$

Check Data

```
In [2]: train_norm_size = len(os.listdir('re-split_data/train/normal'))
        train_pneum_size = len(os.listdir('re-split_data/train/pneumonia'))
        test_norm_size = len(os.listdir('re-split_data/test/normal'))
        test_pneum_size = len(os.listdir('re-split_data/test/pneumonia'))
        valid_norm_size = len(os.listdir('re-split_data/validation/normal'))
        valid_pneum_size = len(os.listdir('re-split_data/validation/pneumonia'))
        train_size = train_norm_size + train_pneum_size -1
        test_size = test_norm_size + test_pneum_size -1
        validation_size = valid_norm_size + valid_pneum_size
        print(f' There are {train_size} images in the training set, {test_size} in
        the test set, and {validation_size} in the validation set')
        print(f' train norm is {train_norm_size}')
        print(f' train pneum is {train_pneum_size}')
        print(f' test norm: {test_norm_size}')
        print(f' test pneum: {test_pneum_size}')
        print(f' valid norm: {valid_norm_size}')
        print(f' valid pneum: {valid_pneum_size}')
```

There are 3272 images in the training set, 1752 in the test set, and 816 in the validation set train norm is 883 train pneum is 2390 test norm: 473 test pneum: 1280

valid norm: 220 valid pneum: 596

Add Functions

```
In [3]:
        # Define Result Saving Initial Function
        dfcols = ['Model', 'RW Score', 'F1', 'Recall', 'Precision', 'Accuracy']
        model_summary = pd. DataFrame(columns=dfcols)
        def save_result(cf, model_name):
                    global model_summary
                    accuracy = np. trace(cf) / float(np. sum(cf))
                    recall = cf[1, 1] / sum(cf[:, 1])
                    precision = cf[1, 1] / sum(cf[1, :])
                    f1_score = 2*precision*recall / (precision + recall)
                    beta = 9 \# 3^2
                    rw_score = (1+beta) * ((precision * recall) / ((beta * precisio
        n) + recall))
                    #cv_std = statistics.stdev([cv1, cv2, cv3, cv4, cv5])
                    \#cv_avg = (cv1 + cv2 + cv3 + cv4 + cv5) / 5
                    results_columns = ['Model', 'RW Score', 'F1', 'Recall', 'Precis
        ion', 'Accuracy']
                    row = [(model_name, rw_score, f1_score, recall, precision, accu
        racy)]
                    res = pd. DataFrame(columns = results_columns, data = row)
                    yeep = [model_summary, res]
                    model_summary = pd.concat(yeep)
                    model_summary = model_summary.sort_values('Accuracy', ascending
        = False)
                    model_summary = model_summary.drop_duplicates()
                    return model_summary.round(3)
```

```
In [4]: # 1. Confusion Matrix
        # SOURCE: The origin of this confusion matrix code was found on medium,
        # from https://medium.com/@dtuk81/confusion-matrix-visualization-fc31e3f30f
        def make_confusion_matrix(cf,
                                   group_names=None,
                                   categori es=' auto',
                                   count=True.
                                   percent=True.
                                   cbar=True,
                                   xyticks=True,
                                   xyplotlabels=True,
                                   sum_stats=True,
                                   fi qsi ze=None,
                                   cmap='Bl ues',
                                   title=None):
             # CODE TO GENERATE TEXT INSIDE EACH SQUARE
            blanks = ['' for i in range(cf. size)]
            if group_names and len(group_names)==cf. size:
                 group_labels = ["{}\n".format(value) for value in group_names]
            el se:
                group_l abel s = bl anks
            if count:
                 group_counts = ["{0:0.0f}\n".format(value) for value in cf.flatten
         ()]
            el se:
                 group_counts = blanks
            if percent:
                 group_percentages = ["{0:.2%}".format(value) for value in cf.flatte
        n()/np.sum(cf)
            el se:
                 group_percentages = blanks
            box_labels = [f''(v1)(v2)(v3)''.strip() for v1, v2, v3 in zip(group_label
        s, group_counts, group_percentages)]
            box_labels = np. asarray(box_labels). reshape(cf. shape[0], cf. shape[1])
             # CODE TO GENERATE SUMMARY STATISTICS & TEXT FOR SUMMARY STATS
            if sum_stats:
                 #Accuracy is sum of diagonal divided by total observations
                 accuracy = np. trace(cf) / float(np. sum(cf))
                 #if it is a binary confusion matrix, show some more stats
                 if len(cf) == 2:
                     #Metrics for Binary Confusion Matrices
                     recall = cf[1, 1] / sum(cf[:, 1])
                     preci si on
                                 = cf[1,1] / sum(cf[1,:])
                     f1_score = 2*precision*recall / (precision + recall)
                     beta = 9
                     rw_score = (1+beta) * ((precision * recall) / ((beta * precisio
        n) + recall))
```

```
stats_text = "\n\nAccuracy={: 0. 3f}\nPrecision={: 0. 3f}\nRecall = "...]
{: 0. 3f}\nF1 Score={: 0. 3f}\nRW Score={: 0. 3f}". format(
                 accuracy, precision, recall, f1_score, rw_score)
        el se:
            stats_text = "\n\nAccuracy={: 0. 3f}". format(accuracy)
    el se:
        stats_text = ""
    # SET FIGURE PARAMETERS ACCORDING TO OTHER ARGUMENTS
    if figsize==None:
        #Get default figure size if not set
        figsize = plt.rcParams.get('figure.figsize')
    if xyticks==False:
        #Do not show categories if xyticks is False
        categori es=Fal se
    # MAKE THE HEATMAP VISUALIZATION
    plt.figure(figsize=figsize)
    sns. heatmap(cf, annot=box_l abels, fmt="", cmap=cmap, cbar=cbar, xti ckl abels=
categories, yticklabels=categories)
    if xyplotlabels:
        plt.ylabel('True label')
        plt.xlabel('Predicted label' + stats_text)
    el se:
        plt.xlabel(stats_text)
    if title:
        plt.title(title)
```

Re-Splitting (Hide)

Check re-aggregated images

Due to the issues with the given train/test split, i re-aggregated the images. From here, I will train/test split the data myself.

```
In [5]:
        folder = 're-split_data/NORMAL'
        path = folder
        p = os. listdir(path)
        pf = pd. DataFrame(p)
        norm_tot = len(pf)
        folder = 're-split_data/PNEUMONIA'
        path = folder
        p = os. listdir(path)
        pf = pd. DataFrame(p)
        pneum_tot = len(pf)
        pneum_weight = len(pf) / 5863
        norm_weight = 1 - pneum_weight
        print(f' There are {len(pf[0])} images in the pneumonia folder')
        print(f' there are {1576 + len(pf[0])} total images in the dataset')
        print(f' The weight of pneumonia is {round(pneum_weight, 2)}')
        print(f' The weight of normal is {round(norm_weight, 2)}')
        1.1.1
```

Out[5]: "\nfolder = 're-split_data/NORMAL'\npath = folder\np = os.listdir(path)\npf = pd.DataFrame(p)\nnorm_tot = len(pf)\n\nfolder = 're-split_data/PNEUMONIA '\npath = folder\np = os.listdir(path)\npf = pd.DataFrame(p)\npneum_tot = len(pf)\npneum_weight = len(pf) / 5863\nnorm_weight = 1 - pneum_weight\n\npr int(f' There are {len(pf[0])} images in the pneumonia folder')\nprint(f' there are {1576 + len(pf[0])} total images in the dataset')\nprint(f' The weight of pneumonia is {round(pneum_weight, 2)}')\nprint(f' The weight of norma lis {round(norm_weight, 2)}')\n\n"

From here, there is some code which I used to re-split the data.

```
In [7]:
        p1 = os.listdir('re-split_data/train/normal')
        p1 = pd. DataFrame(p1)
        p2 = os.listdir('re-split_data/NORMAL')
        p2 = pd. DataFrame(p2)
        tester_files = pd. concat([p1[0], p2[0]]). drop_duplicates(keep=False)
Out[7]: "\np1 = os.listdir('re-split_data/train/normal')\np1 = pd.DataFrame(p1)\n\n
        p2 = os.listdir('re-split_data/NORMAL')\np2 = pd.DataFrame(p2)\n\ntester_fi
        les = pd. concat([p1[0], p2[0]]). drop_duplicates(keep=False)\n"
        #for file in tester_files:
In [8]:
            #shutil.copy('re-split_data/NORMAL/' + file, 're-split_data/test/normal
         ')
In [9]:
        # The pneumonia files
        pf = os.listdir('re-split_data/PNEUMONIA')
        rand_Pfiles = random.sample(pf, int(pneum_train))
        for file in rand_Pfiles:
            shutil.copy('re-split_data/PNEUMONIA/' + file, 're-split_data/train/pne
        umoni a')
        p3 = os.listdir('re-split_data/train/pneumonia')
        p3 = pd. DataFrame(p3)
        p4 = os. listdir('re-split_data/PNEUMONIA')
        p4 = pd. DataFrame(p4)
        tester_p = pd. concat([p3[0], p4[0]]). drop_duplicates(keep=False)
        for file in tester_p:
            shutil.copy('re-split_data/PNEUMONIA/' + file, 're-split_data/test/pneu
        moni a')
        111
        "\n# The pneumonia files\npf = os.listdir('re-split_data/PNEUMONIA')\n\nran
```

Validation Files

Make a validation set from the train set

```
In [10]:
         pf = os.listdir('re-split_data/train/normal')
         norm\_tot = Ien(pf)
         pf1 = os.listdir('re-split_data/train/pneumonia')
         pneum tot = Ien(pf1)
         print(f' There are {norm_tot} images in the normal training folder and {pne
         um_tot} in the pneumonia training folder')
         "\npf = os.listdir('re-split_data/train/normal')\nnorm_tot = len(pf)\npf1 =
Out[10]:
         os. listdir('re-split_data/train/pneumonia')\npneum_tot = len(pf1)\n\nprint
         (f' There are {norm_tot} images in the normal training folder and {pneum_to
         t} in the pneumonia training folder')\n"
In [11]:
         pf2 = os.listdir('re-split_data/validation/normal')
         valid_norm_tot = len(pf2)
         pf3 = os.listdir('re-split_data/validation/pneumonia')
         valid_pneum_tot = len(pf3)
         rand_sample_valid_norm = random.sample(pf2, 110)
         rand sample valid pneum = random.sample(pf3, 298)
         for file in rand_sample_valid_norm:
             shutil.move('re-split_data/validation/normal/' + file, 're-split_data/t
         rai n/normal')
         for file in rand sample valid pneum:
             shutil.move('re-split_data/validation/pneumonia/' + file, 're-split_dat
         a/trai n/pneumoni a')
         . . .
         "\npf2 = os.listdir('re-split_data/validation/normal')\nvalid_norm_tot = le
Out[11]:
         n(pf2)\npf3 = os.listdir('re-split_data/validation/pneumonia')\nvalid_pneum
```

Out[11]: "\npf2 = os.listdir('re-split_data/validation/normal')\nvalid_norm_tot = le n(pf2)\npf3 = os.listdir('re-split_data/validation/pneumonia')\nvalid_pneum _tot = len(pf3)\n\nrand_sample_valid_norm = random.sample(pf2, 110)\nrand_s ample_valid_pneum = random.sample(pf3, 298)\n\nfor file in rand_sample_valid_norm:\n shutil.move('re-split_data/validation/normal/' + file, 're-split_data/train/normal')\n\nfor file in rand_sample_valid_pneum:\n shutil.move('re-split_data/validation/pneumonia/' + file, 're-split_data/train/pneumonia')\n \n"

Generate Test and Train Images

```
In [12]:
          # get all the data in the directory split/test , and reshape them
           train_generator = ImageDataGenerator(rescal e=1. /255). flow_from_directory('r
           e-split_data/train',
                   target_size=(64, 64), batch_size = train_size)
           test_generator = I mageDataGenerator(rescal e=1. /255). flow_from_directory('re-
           split_data/test',
                   target_size=(64, 64), batch_size = test_size, shuffle= False)
           valid_generator = I mageDataGenerator(rescale=1./255). flow_from_directory('re
           -split_data/validation',
                   target_size=(64, 64), batch_size = validation_size)
           Found 3272 images belonging to 2 classes.
           Found 1752 images belonging to 2 classes.
           Found 816 images belonging to 2 classes.
 In [13]: # create the data sets
           train_i mages, train_labels = next(train_generator)
           test_i mages, test_l abel s = next(test_generator)
           valid_images, valid_labels = next(valid_generator)
In [185]:
          train_labels2 = pd. DataFrame(train_labels)
           train_labels2
Out[185]:
                  0
                      1
              0 0.0 1.0
              1 0.0 1.0
              2 0.0 1.0
              3 0.0 1.0
              4 0.0 1.0
           3267 0.0 1.0
           3268 0.0 1.0
           3269 0.0 1.0
           3270 0.0 1.0
           3271 0.0 1.0
           3272 rows x 2 columns
In [186]: train_labels2[0].sum()
Out[186]: 882.0
In [187]: train_labels2[1].sum()
Out[187]: 2390.0
```

This shows us that if column '0' equals 1, it is a NORMAL label.

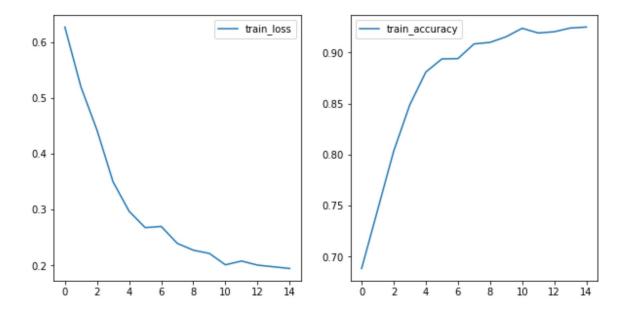
```
In [14]: def show_i mages(i mages):
              fig, axes = plt. subplots (1, 10, figsize=(12, 12))
              axes = axes.flatten()
              for img, ax in zip(images, axes):
                  ax.imshow(img)
                  ax. axi s('off')
              plt.tight_layout()
              pl t. show()
In [15]:
          show_i mages(trai n_i mages)
         train_img = train_images.reshape(train_images.shape[0], -1)
In [16]:
          test_i mg = test_i mages. reshape(test_i mages. shape[0], -1)
          valid_img = valid_images.reshape(valid_images.shape[0], -1)
          print(train_img. shape)
          print(test_i mg. shape)
          print(valid_img.shape)
          (3272, 12288)
          (1752, 12288)
          (816, 12288)
In [17]: train_y = np. reshape(train_labels[:,0], (train_size, 1))
          test_y = np. reshape(test_l abel s[:, 0], (test_si ze, 1))
          valid_y = np.reshape(valid_labels[:,0], (validation_size,1))
          pri nt(trai n_y. shape)
          print(test_y. shape)
          pri nt (val i d_y. shape)
          (3272, 1)
          (1752, 1)
          (816, 1)
```

Baseline Model

```
In [18]: # Build a baseline fully connected model
    model = models. Sequential()
    model.add(layers.Dense(20, activation='relu', input_shape=(12288,))) # 2 hi
    dden layers
    model.add(layers.Dense(7, activation='relu'))
    model.add(layers.Dense(5, activation='relu'))
    model.add(layers.Dense(1, activation='sigmoid'))
```

Epoch 1/15	•	75 / .			0 (0(0		
103/103 [======] -	- 89	5 /5ms/step	-	l oss:	0. 6268	-	ac
curacy: 0.6886							
Epoch 2/15							
103/103 [========] -	- 99	86ms/step	-	loss:	0. 5197	-	ac
curacy: 0.7460							
Epoch 3/15							
103/103 [========] -	- 89	80ms/step	-	loss:	0. 4422	-	ac
curacy: 0.8035							
Epoch 4/15							
103/103 [====================================	- 89	77ms/step	_	loss:	0.3498	_	ac
curacy: 0.8487		·					
Epoch 5/15							
103/103 [====================================	- 89	77ms/step	_	loss:	0. 2971	_	ac
curacy: 0.8808							
Epoch 6/15							
103/103 [====================================	_ 89	78ms/sten	_	l nss·	0 2677	_	ac
curacy: 0.8936	0.	, , oms, stop		1055.	0. 2011		uo
Epoch 7/15							
103/103 [====================================	_ &	78ms/sten	_	l nee.	0 2699	_	ac
curacy: 0.8939	U.	5 7011137 3 CCP		1033.	0.2077		ac
Epoch 8/15							
103/103 [====================================	0.	74mc/cton		Locci	0 2205		00
	- 03	5 /ollis/step	-	1055.	0. 2393	-	ac
curacy: 0.9083							
Epoch 9/15	0-	70/			0 0074		
103/103 [======] -	- 89	s /9ms/step	-	TOSS:	0. 2274	-	ac
curacy: 0. 9098							
Epoch 10/15	0	70/			0 0015		
103/103 [======] -	- 89	s /9ms/step	-	TOSS:	0. 2215	-	ac
curacy: 0. 9153							
Epoch 11/15	_			_			
103/103 [======] -	- 89	79ms/step	-	loss:	0. 2012	-	ac
curacy: 0. 9236							
Epoch 12/15							
103/103 [=======] -	- 89	82ms/step	-	loss:	0. 2080	-	ac
curacy: 0.9190							
Epoch 13/15							
103/103 [========] -	- 89	76ms/step	-	loss:	0. 2007	-	ac
curacy: 0.9202							
Epoch 14/15							
103/103 [=========] -	- 89	78ms/step	-	loss:	0. 1976	_	ac
curacy: 0.9239		·					
Epoch 15/15							
103/103 [====================================	- 89	80ms/step	-	loss:	0. 1946	_	ac
curacy: 0. 9248		•					
AvacSubplates							

Out[19]: <AxesSubpl ot: >



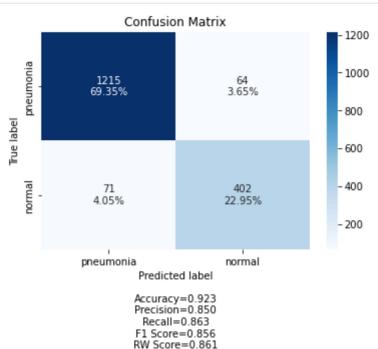
SkLearn Version for Cross-Validation

```
In [20]:
         # Build function that builds the model so we can evaluate in sklearn
         def build_model():
             model.add(layers.Dense(20, activation='relu', input_shape=(12288,))) #
         2 hidden layers
             model.add(layers.Dense(7, activation='relu'))
             model.add(layers.Dense(5, activation='relu'))
             model.add(layers.Dense(1, activation='sigmoid'))
             model.compile(optimizer='sgd',
                        loss='binary_crossentropy',
                        metri cs=['accuracy'])
              return model
         keras_model = scikit_learn. KerasClassifier(build_model,
In [21]:
                                                     epochs=15,
                                                     batch_si ze=32,
                                                     verbose=2)
```

Results

```
In [23]: results_test = model.evaluate(test_img, test_y)
         55/55 [============== ] - 4s 74ms/step - loss: 0.1946 - accu
         racy: 0.9229
In [24]:
         predictions = model.predict(x = test_img, verbose=0)
In [25]:
         pred_check = np. round(predictions)
In [26]:
         pred_check = pred_check[:]
         pred_check = pred_check.flatten()
         pred_check
Out[26]:
         array([0., 1., 1., ..., 0., 0., 0.], dtype=float32)
         test_check = test_labels[:,0]
In [27]:
         test_check
Out[27]:
         array([1., 1., 1., ..., 0., 0., 0.], dtype=float32)
In [28]:
         cm = confusi on_matri x(y_true=test_check, y_pred=pred_check)
In [29]:
         save_result(cm, 'baseline_model')
Out[29]:
                                    F1 Recall Precision Accuracy
                   Model RW Score
          0 baseline_model
                             0.861 0.856
                                        0.863
                                                  0.85
                                                          0.923
```

In [30]: cm_plot_labels = ['pneumonia', 'normal']
 make_confusion_matrix(cm, categories = cm_plot_labels, title='Confusion Mat
 rix')



Convolutional Neural Network (CNN) #1

In [32]: model.summary()

Model: "sequential_1"

Layer (type)	Output	Shape	Param #
conv2d (Conv2D)	(None,	62, 62, 32)	896
max_pooling2d (MaxPooling2D)	(None,	31, 31, 32)	0
conv2d_1 (Conv2D)	(None,	28, 28, 32)	16416
max_pooling2d_1 (MaxPooling2	(None,	14, 14, 32)	0
conv2d_2 (Conv2D)	(None,	12, 12, 64)	18496
max_pooling2d_2 (MaxPooling2	(None,	6, 6, 64)	0
flatten (Flatten)	(None,	2304)	0
dense_4 (Dense)	(None,	64)	147520
dense_5 (Dense)	(None,	1)	65

Total params: 183,393 Trainable params: 183,393 Non-trainable params: 0

Train Initial Simple CNN

```
Epoch 1/25
103/103 [=============== ] - 16s 154ms/step - Loss: 0.5929 -
accuracy: 0.7304 - val_loss: 0.5820 - val_accuracy: 0.7304
Epoch 2/25
accuracy: 0.7304 - val_loss: 0.5572 - val_accuracy: 0.7304
accuracy: 0.7304 - val_loss: 0.4996 - val_accuracy: 0.7304
Epoch 4/25
103/103 [================= ] - 15s 145ms/step - Loss: 0.4832 -
accuracy: 0.7781 - val_loss: 0.3968 - val_accuracy: 0.8542
Epoch 5/25
accuracy: 0.8188 - val_loss: 0.3631 - val_accuracy: 0.8076
Epoch 6/25
accuracy: 0.8579 - val_loss: 0.4914 - val_accuracy: 0.7647
Epoch 7/25
accuracy: 0.8854 - val_loss: 0.2383 - val_accuracy: 0.8971
Epoch 8/25
accuracy: 0.8973 - val_loss: 0.5562 - val_accuracy: 0.7843
accuracy: 0.8985 - val_loss: 0.2822 - val_accuracy: 0.8664
Epoch 10/25
accuracy: 0.9181 - val_loss: 0.2163 - val_accuracy: 0.9093
Epoch 11/25
accuracy: 0.9169 - val_loss: 0.2437 - val_accuracy: 0.8958
Epoch 12/25
103/103 [============ ] - 15s 142ms/step - Loss: 0.1978 -
accuracy: 0.9233 - val_loss: 0.3334 - val_accuracy: 0.8468
Epoch 13/25
accuracy: 0.9224 - val_loss: 0.1677 - val_accuracy: 0.9314
Epoch 14/25
accuracy: 0.9328 - val_loss: 0.1644 - val_accuracy: 0.9449
Epoch 15/25
accuracy: 0.9273 - val_loss: 0.1661 - val_accuracy: 0.9326
Epoch 16/25
103/103 [=========== ] - 13s 131ms/step - Loss: 0.1773 -
accuracy: 0.9318 - val_loss: 0.2322 - val_accuracy: 0.9069
Epoch 17/25
accuracy: 0.9288 - val_loss: 0.1497 - val_accuracy: 0.9461
Epoch 18/25
accuracy: 0.9373 - val_loss: 0.1440 - val_accuracy: 0.9461
Epoch 19/25
103/103 [=============== ] - 17s 162ms/step - Loss: 0.1703 -
```

```
accuracy: 0.9346 - val_loss: 0.4063 - val_accuracy: 0.8554
       Epoch 20/25
       103/103 [============== ] - 16s 160ms/step - Loss: 0.1666 -
       accuracy: 0.9383 - val_loss: 0.1440 - val_accuracy: 0.9498
       Epoch 21/25
       accuracy: 0.9370 - val_loss: 0.1507 - val_accuracy: 0.9510
       Epoch 22/25
       103/103 [=============== ] - 16s 156ms/step - Loss: 0.1582 -
       accuracy: 0.9389 - val_loss: 0.1352 - val_accuracy: 0.9571
       Epoch 23/25
       accuracy: 0.9419 - val_loss: 0.1313 - val_accuracy: 0.9510
       Epoch 24/25
       103/103 [=============== ] - 16s 156ms/step - Loss: 0.1573 -
       accuracy: 0.9407 - val_loss: 0.1347 - val_accuracy: 0.9510
       Epoch 25/25
       103/103 [============ ] - 15s 144ms/step - Loss: 0.1509 -
       accuracy: 0.9425 - val_loss: 0.1316 - val_accuracy: 0.9608
In [35]: results_train = model.evaluate(train_images, train_y)
       103/103 [=============== ] - 4s 40ms/step - loss: 0.1364 - ac
       curacy: 0.9474
In [36]: results_test = model.evaluate(test_images, test_y)
       55/55 [============== ] - 2s 35ms/step - loss: 0.1803 - accu
       racy: 0.9309
```

Prediction for Confusion Matrix

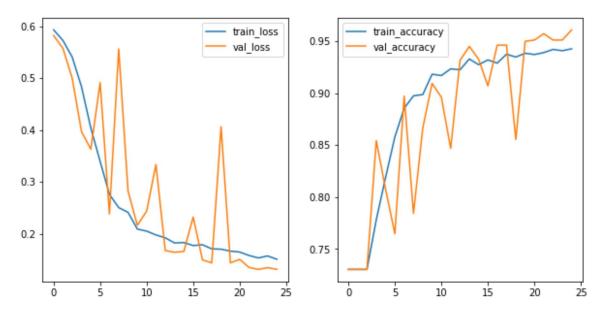
```
In [37]: predictions = model.predict(x = test_images, steps = 10, verbose=0)
In [38]: pred_check = np.round(predictions)
In [39]: pred_check = pred_check[:]
    pred_check = pred_check.flatten()
    pred_check
Out[39]: array([0., 1., 1., ..., 0., 0., 0.], dtype=float32)
In [40]: test_check = test_labels[:,0]
    test_check
Out[40]: array([1., 1., 1., ..., 0., 0., 0.], dtype=float32)
In [41]: cm = confusion_matrix(y_true=test_check, y_pred=pred_check)
```

CNN #1 Results

```
In [42]: train_loss = cnn_1.history['loss']
    train_acc = cnn_1.history['accuracy']
    val_loss = cnn_1.history['val_loss']
    val_acc = cnn_1.history['val_accuracy']

fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(10, 5))
    sns.lineplot(x=cnn_1.epoch, y=train_loss, ax=ax1, label='train_loss')
    sns.lineplot(x=cnn_1.epoch, y=train_acc, ax=ax2, label='train_accuracy')
    sns.lineplot(x=cnn_1.epoch, y=val_loss, ax=ax1, label='val_loss')
    sns.lineplot(x=cnn_1.epoch, y=val_acc, ax=ax2, label='val_accuracy')
```

Out[42]: <AxesSubpl ot: >



```
In [43]: def build_cnn():
             model = models.Sequential()
             model.add(layers.Conv2D(32, (3, 3), activation='relu', input_shape=(64
         , 64,
               3)))
             model.add(layers.MaxPooling2D((2, 2)))
             model.add(layers.Conv2D(32, (4, 4), activation='relu'))
             model.add(layers.MaxPooling2D((2, 2)))
             model.add(layers.Conv2D(64, (3, 3), activation='relu'))
             model.add(layers.MaxPooling2D((2, 2)))
             model.add(layers.Flatten())
             model.add(layers.Dense(64, activation='relu'))
             model.add(layers.Dense(1, activation='sigmoid'))
                                                                    #Last layer mus
         t be 1 for binary classification
             model.compile(loss='binary_crossentropy',
                       optimizer="sgd",
                       metri cs=['accuracy'])
             return model
         keras_model 2 = scikit_learn. KerasClassifier(build_cnn,
In [44]:
                                                      epochs=25,
                                                      validation_data=(valid_images,
         valid_y),
```

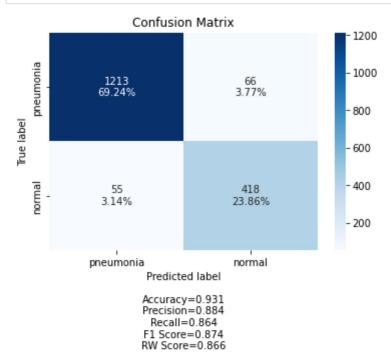
```
validation_steps = validation_s
ize)
```

In [45]: save_result(cm, 'CNN #1')

Out[45]:

	Model	RW Score	F1	Recall	Precision	Accuracy
0	CNN #1	0.866	0.874	0.864	0.884	0.931
0	baseline_model	0.861	0.856	0.863	0.850	0.923

In [46]: cm_plot_labels = ['pneumonia', 'normal']
make_confusion_matrix(cm, categories = cm_plot_labels, title='Confusion Mat
rix')



CNN Model 2

For this model, I added another round of Conv2D and MaxPooling layers, and changed the optimizer to "adam"

```
In [47]: model 2 = model s. Sequential()
          model 2. add(layers. Conv2D(32, (3, 3), activation='relu',
                                    input_shape=(64,64, 3)))
          model 2. add(layers. MaxPooling2D((2, 2)))
          model 2. add(layers. Conv2D(32, (4, 4), activation='relu'))
          model 2. add(layers. MaxPooling2D((2, 2)))
          model 2. add(l ayers. Conv2D(64, (3, 3), activation='relu'))
          model 2. add(layers. MaxPooling2D((2, 2)))
          model 2. add(layers. Conv2D(96, (3, 3), activation='relu'))
          model 2. add(layers. MaxPooling2D((2, 2)))
          model 2. add(l ayers. Fl atten())
          model 2. add(layers. Dense(64, activation='relu'))
          model 2. add(l ayers. Dense(1, activation='sigmoid'))
          model 2. compile(loss='binary_crossentropy',
                         opti mi zer="adam",
                         metri cs=['accuracy'])
```

```
Epoch 1/50
103/103 [=============== ] - 14s 128ms/step - Loss: 0.5128 -
accuracy: 0.7549 - val_loss: 0.4897 - val_accuracy: 0.7402
Epoch 2/50
103/103 [============ ] - 13s 127ms/step - Loss: 0.2891 -
accuracy: 0.8811 - val_loss: 0.1954 - val_accuracy: 0.9167
accuracy: 0.9141 - val_loss: 0.1449 - val_accuracy: 0.9436
Epoch 4/50
103/103 [================ ] - 11s 112ms/step - Loss: 0.1590 -
accuracy: 0.9377 - val_loss: 0.1154 - val_accuracy: 0.9596
Epoch 5/50
accuracy: 0.9542 - val_loss: 0.1255 - val_accuracy: 0.9632
Epoch 6/50
accuracy: 0.9514 - val_loss: 0.1058 - val_accuracy: 0.9620
Epoch 7/50
accuracy: 0.9603 - val_loss: 0.0941 - val_accuracy: 0.9657
Epoch 8/50
accuracy: 0.9633 - val_loss: 0.0930 - val_accuracy: 0.9669
accuracy: 0.9649 - val_loss: 0.1270 - val_accuracy: 0.9559
Epoch 10/50
accuracy: 0.9710 - val_loss: 0.1081 - val_accuracy: 0.9694
Epoch 11/50
103/103 [=============== ] - 13s 127ms/step - Loss: 0.0830 -
accuracy: 0.9676 - val_loss: 0.1210 - val_accuracy: 0.9596
Epoch 12/50
accuracy: 0.9759 - val_loss: 0.1048 - val_accuracy: 0.9608
Epoch 13/50
accuracy: 0.9780 - val_loss: 0.0923 - val_accuracy: 0.9706
Epoch 14/50
accuracy: 0.9780 - val_loss: 0.0915 - val_accuracy: 0.9706
Epoch 15/50
accuracy: 0.9850 - val_loss: 0.1108 - val_accuracy: 0.9669
Epoch 16/50
103/103 [=========== ] - 12s 120ms/step - Loss: 0.0328 -
accuracy: 0.9887 - val_loss: 0.1095 - val_accuracy: 0.9620
Epoch 17/50
accuracy: 0.9948 - val_loss: 0.1209 - val_accuracy: 0.9681
Epoch 18/50
103/103 [================ ] - 12s 121ms/step - Loss: 0.0216 -
accuracy: 0.9924 - val_loss: 0.1031 - val_accuracy: 0.9694
Epoch 19/50
103/103 [=============== ] - 13s 130ms/step - Loss: 0.0180 -
```

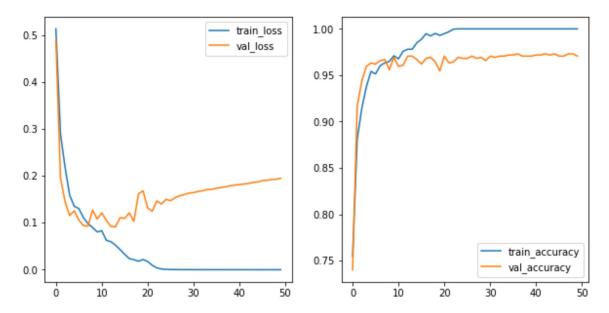
```
accuracy: 0.9951 - val_loss: 0.1614 - val_accuracy: 0.9645
Epoch 20/50
103/103 [============== ] - 13s 131ms/step - Loss: 0.0218 -
accuracy: 0.9930 - val_loss: 0.1682 - val_accuracy: 0.9547
Epoch 21/50
accuracy: 0.9948 - val_loss: 0.1314 - val_accuracy: 0.9706
Epoch 22/50
103/103 [=============== ] - 13s 124ms/step - Loss: 0.0095 -
accuracy: 0.9969 - val_loss: 0.1246 - val_accuracy: 0.9632
Epoch 23/50
accuracy: 0.9997 - val_loss: 0.1466 - val_accuracy: 0.9645
Epoch 24/50
103/103 [=============== ] - 13s 124ms/step - Loss: 0.0015 -
accuracy: 1.0000 - val_loss: 0.1398 - val_accuracy: 0.9694
Epoch 25/50
4 - accuracy: 1.0000 - val_loss: 0.1498 - val_accuracy: 0.9681
Epoch 26/50
4 - accuracy: 1.0000 - val_loss: 0.1469 - val_accuracy: 0.9681
Epoch 27/50
4 - accuracy: 1.0000 - val_loss: 0.1533 - val_accuracy: 0.9706
Epoch 28/50
4 - accuracy: 1.0000 - val_loss: 0.1576 - val_accuracy: 0.9681
Epoch 29/50
4 - accuracy: 1.0000 - val_loss: 0.1601 - val_accuracy: 0.9694
Epoch 30/50
4 - accuracy: 1.0000 - val_loss: 0.1632 - val_accuracy: 0.9657
Epoch 31/50
103/103 [================ ] - 13s 127ms/step - Loss: 1.8072e-0
4 - accuracy: 1.0000 - val loss: 0.1642 - val accuracy: 0.9706
Epoch 32/50
103/103 [================= ] - 12s 121ms/step - Loss: 1.6162e-0
4 - accuracy: 1.0000 - val_loss: 0.1667 - val_accuracy: 0.9694
Epoch 33/50
4 - accuracy: 1.0000 - val_loss: 0.1682 - val_accuracy: 0.9706
Epoch 34/50
103/103 [================= ] - 13s 124ms/step - Loss: 1.2853e-0
4 - accuracy: 1.0000 - val_loss: 0.1708 - val_accuracy: 0.9706
Epoch 35/50
4 - accuracy: 1.0000 - val_loss: 0.1711 - val_accuracy: 0.9718
Epoch 36/50
103/103 [================ ] - 13s 129ms/step - Loss: 1.0194e-0
4 - accuracy: 1.0000 - val_loss: 0.1735 - val_accuracy: 0.9718
Epoch 37/50
5 - accuracy: 1.0000 - val_loss: 0.1751 - val_accuracy: 0.9730
Epoch 38/50
```

```
5 - accuracy: 1.0000 - val_loss: 0.1767 - val_accuracy: 0.9706
      Epoch 39/50
      103/103 [=============== ] - 13s 126ms/step - Loss: 7.7741e-0
      5 - accuracy: 1.0000 - val_loss: 0.1787 - val_accuracy: 0.9706
      Epoch 40/50
      5 - accuracy: 1.0000 - val_loss: 0.1803 - val_accuracy: 0.9706
      Epoch 41/50
      5 - accuracy: 1.0000 - val_loss: 0.1813 - val_accuracy: 0.9718
      Epoch 42/50
      5 - accuracy: 1.0000 - val_loss: 0.1824 - val_accuracy: 0.9718
      Epoch 43/50
      5 - accuracy: 1.0000 - val_loss: 0.1838 - val_accuracy: 0.9730
      Epoch 44/50
      5 - accuracy: 1.0000 - val_loss: 0.1859 - val_accuracy: 0.9718
      Epoch 45/50
      103/103 [=============== ] - 12s 120ms/step - Loss: 4.7310e-0
      5 - accuracy: 1.0000 - val_loss: 0.1869 - val_accuracy: 0.9730
      Epoch 46/50
      5 - accuracy: 1.0000 - val_loss: 0.1897 - val_accuracy: 0.9706
      Epoch 47/50
      5 - accuracy: 1.0000 - val_loss: 0.1904 - val_accuracy: 0.9706
      Epoch 48/50
      5 - accuracy: 1.0000 - val_loss: 0.1923 - val_accuracy: 0.9730
      Epoch 49/50
      5 - accuracy: 1.0000 - val_loss: 0.1926 - val_accuracy: 0.9730
      Epoch 50/50
      5 - accuracy: 1.0000 - val_loss: 0.1947 - val_accuracy: 0.9706
In [49]:
     results_train = model 2. evaluate(train_images, train_y)
      103/103 [============= ] - 1s 10ms/step - loss: 2.7606e-05
      - accuracy: 1.0000
In [50]:
     results_test = model 2. evaluate(test_i mages, test_y)
      55/55 [============== ] - 1s 10ms/step - Loss: 0.3585 - accu
      racy: 0.9515
```

```
In [51]: train_loss = history2.history['loss']
    train_acc = history2.history['accuracy']
    val_loss = history2.history['val_loss']
    val_acc = history2.history['val_accuracy']

fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(10, 5))
    sns.lineplot(x=history2.epoch, y=train_loss, ax=ax1, label='train_loss')
    sns.lineplot(x=history2.epoch, y=train_acc, ax=ax2, label='train_accuracy')
    sns.lineplot(x=history2.epoch, y=val_loss, ax=ax1, label='val_loss')
    sns.lineplot(x=history2.epoch, y=val_acc, ax=ax2, label='val_accuracy')
```

Out[51]: <AxesSubpl ot: >



```
In [52]: def build_cnn2():
              model 2 = model s. Sequential ()
              model 2. add(layers. Conv2D(32, (3, 3), activation='relu',
                                   input_shape=(64,64,
              model 2. add(layers. MaxPooling2D((2, 2)))
              model 2. add(layers. Conv2D(32, (4, 4), activation='relu'))
              model 2. add(layers. MaxPooling2D((2, 2)))
              model 2. add(layers. Conv2D(64, (3, 3), activation='relu'))
              model 2. add(layers. MaxPooling2D((2, 2)))
              model 2. add(layers. Conv2D(96, (3, 3), activation='relu'))
              model 2. add(layers. MaxPooling2D((2, 2)))
              model 2. add(l ayers. Fl atten())
              model 2. add(layers. Dense(64, activation='relu'))
              model 2. add(layers. Dense(1, activation='sigmoid'))
              model 2. compile(loss='binary_crossentropy',
                         opti mi zer="Adam",
                         metri cs=['accuracy'])
              return model 2
In [53]: keras_model 3 = scikit_learn. KerasClassifier(build_cnn2,
                                                         epochs=50.
                                                         validation_data=(valid_images,
          valid_y),
                                                         validation_steps = validation_s
          ize)
```

Prediction for Confusion Matrix

```
In [54]: predictions = model 2. predict(x = test_images, steps = 10, verbose=0)
In [55]: pred_check = np. round(predictions)
In [56]: pred_check = pred_check[:]
    pred_check = pred_check. flatten()
    pred_check
Out[56]: array([1., 1., 1., ..., 0., 0.], dtype=float32)
In [57]: test_check = test_labels[:,0]
    test_check
Out[57]: array([1., 1., 1., ..., 0., 0.], dtype=float32)
In [58]: cm = confusion_matrix(y_true=test_check, y_pred=pred_check)
```

```
In [59]: save_result(cm, 'CNN #2', )

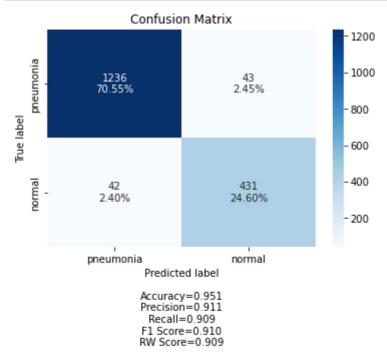
Out[59]:

Model RW Score F1 Recall Precision Accuracy

O CNN #2 0 909 0 910 0 909 0 911 0 951
```

	Model	RW Score	F1	Recall	Precision	Accuracy
0	CNN #2	0.909	0.910	0.909	0.911	0.951
0	CNN #1	0.866	0.874	0.864	0.884	0.931
0	baseline_model	0.861	0.856	0.863	0.850	0.923

```
In [60]: cm_plot_labels = ['pneumonia', 'normal']
    make_confusion_matrix(cm, categories = cm_plot_labels, title='Confusion Mat
    rix')
```



CNN Model 3

For this model, I add two layers of batch normalization and a 10% dropout. Also added one more layer of Conv2D and MaxPooling.

```
In [61]: model 3 = model s. Sequential()
          model 3. add(layers. Conv2D(32, (3, 3), activation='relu', input_shape=(64,64,
          3)))
          model 3. add(l ayers. MaxPooling2D((2, 2)))
          model 3. add(layers. Conv2D(32, (4, 4), activation='relu'))
          model 3. add(BatchNormalization())
          model 3. add(layers. MaxPooling2D((2, 2)))
          model 3. add(layers. Conv2D(64, (3, 3), activation='relu'))
          model 3. add(layers. MaxPooling2D((2, 2)))
          model 3. add(layers. Conv2D(128, (3, 3), activation='relu'))
          model 3. add(BatchNormalization())
          model 3. add(layers. MaxPooling2D((2, 2)))
          model 3. add(l ayers. Fl atten())
          model 3. add(layers. Dense(64, activation='relu'))
          model 3. add(Dropout(0.1))
          model 3. add(l ayers. Dense(1, activation='sigmoid'))
          model 3. compile(loss='binary_crossentropy',
                         opti mi zer="adam",
                         metri cs=['accuracy'])
```

In [62]: model 3. summary()

Model: "sequential_3"

Layer (type)	Output	Shape	Param #
conv2d_7 (Conv2D)	(None,	62, 62, 32)	896
max_pooling2d_7 (MaxPooling2	(None,	31, 31, 32)	0
conv2d_8 (Conv2D)	(None,	28, 28, 32)	16416
batch_normalization (BatchNo	(None,	28, 28, 32)	128
max_pooling2d_8 (MaxPooling2	(None,	14, 14, 32)	0
conv2d_9 (Conv2D)	(None,	12, 12, 64)	18496
max_pooling2d_9 (MaxPooling2	(None,	6, 6, 64)	0
conv2d_10 (Conv2D)	(None,	4, 4, 128)	73856
batch_normalization_1 (Batch	(None,	4, 4, 128)	512
max_pooling2d_10 (MaxPooling	(None,	2, 2, 128)	0
flatten_2 (Flatten)	(None,	512)	0
dense_8 (Dense)	(None,	64)	32832
dropout (Dropout)	(None,	64)	0
dense_9 (Dense)	(None,	1)	65

Total params: 143,201 Trainable params: 142,881 Non-trainable params: 320

In [63]: print(f' trainsize {train_size}, validation size {validation_size}')

trainsize 3272, validation size 816

In [64]: history3 = model 3. fit(train_images, #Make sure that your dataset can gener ate at least `steps_per_epoch * epochs` batches

train_y, #Integer or None. Number of samples per gradi ent update. default to 32. Do not specify the batch_size if your data is in the form of datasets, generators, or keras. utils. Sequence instances (since they generate batches).

#steps_per_epoch = 100, #The steps per epoch determine s how many steps are done before the model is updated.

epochs=50, # Integer. Number of e pochs to train the model. An epoch is an iteration over the entire x and y data provided (unless the steps per epoch flag is set to something other th an None). Note that in conjunction with initial_epoch, epochs is to be unde rstood as "final epoch". The model is not trained for a number of iteration s given by epochs, but merely until the epoch of index epochs is reached.

> validation_data=(valid_images, valid_y), validation_steps = validation_size)

```
Epoch 1/50
103/103 [=============== ] - 15s 141ms/step - Loss: 0.2536 -
accuracy: 0.9028 - val_loss: 0.5468 - val_accuracy: 0.7243
Epoch 2/50
103/103 [============ ] - 14s 136ms/step - Loss: 0.1478 -
accuracy: 0.9450 - val_loss: 1.0107 - val_accuracy: 0.3039
accuracy: 0.9578 - val_loss: 0.6858 - val_accuracy: 0.6054
Epoch 4/50
accuracy: 0.9575 - val_loss: 0.7674 - val_accuracy: 0.4657
Epoch 5/50
accuracy: 0.9639 - val_loss: 0.2213 - val_accuracy: 0.9142
Epoch 6/50
accuracy: 0.9667 - val_loss: 4.6814 - val_accuracy: 0.7304
Epoch 7/50
accuracy: 0.9688 - val_loss: 1.0449 - val_accuracy: 0.5012
Epoch 8/50
accuracy: 0.9765 - val_loss: 0.1416 - val_accuracy: 0.9498
accuracy: 0.9832 - val_loss: 0.3669 - val_accuracy: 0.8971
Epoch 10/50
accuracy: 0.9853 - val_loss: 0.2769 - val_accuracy: 0.8934
Epoch 11/50
103/103 [=============== ] - 15s 141ms/step - Loss: 0.0301 -
accuracy: 0.9884 - val_loss: 0.2698 - val_accuracy: 0.9179
Epoch 12/50
103/103 [============ ] - 15s 142ms/step - Loss: 0.0366 -
accuracy: 0.9875 - val_loss: 0.5107 - val_accuracy: 0.8064
Epoch 13/50
accuracy: 0.9841 - val_loss: 0.3394 - val_accuracy: 0.9044
Epoch 14/50
accuracy: 0.9917 - val_loss: 0.1878 - val_accuracy: 0.9583
Epoch 15/50
accuracy: 0.9939 - val_loss: 0.1614 - val_accuracy: 0.9596
Epoch 16/50
103/103 [=========== ] - 14s 135ms/step - Loss: 0.0151 -
accuracy: 0.9942 - val_loss: 0.6765 - val_accuracy: 0.8529
Epoch 17/50
accuracy: 0.9988 - val_loss: 0.4380 - val_accuracy: 0.9375
Epoch 18/50
103/103 [================= ] - 13s 125ms/step - Loss: 0.0026 -
accuracy: 0.9991 - val_loss: 0.2337 - val_accuracy: 0.9436
Epoch 19/50
```

```
4 - accuracy: 1.0000 - val_loss: 0.2085 - val_accuracy: 0.9596
Epoch 20/50
4 - accuracy: 1.0000 - val_loss: 0.2149 - val_accuracy: 0.9620
Epoch 21/50
4 - accuracy: 1.0000 - val_loss: 0.2286 - val_accuracy: 0.9596
Epoch 22/50
103/103 [================= ] - 14s 141ms/step - Loss: 1.5958e-0
4 - accuracy: 1.0000 - val_loss: 0.2406 - val_accuracy: 0.9596
Epoch 23/50
4 - accuracy: 1.0000 - val_loss: 0.2444 - val_accuracy: 0.9620
Epoch 24/50
4 - accuracy: 1.0000 - val_loss: 0.2453 - val_accuracy: 0.9645
Epoch 25/50
4 - accuracy: 1.0000 - val_loss: 0.2621 - val_accuracy: 0.9596
Epoch 26/50
4 - accuracy: 1.0000 - val_loss: 0.2512 - val_accuracy: 0.9632
Epoch 27/50
103/103 [=============== ] - 14s 132ms/step - Loss: 7.1218e-0
5 - accuracy: 1.0000 - val_loss: 0.2486 - val_accuracy: 0.9669
Epoch 28/50
5 - accuracy: 1.0000 - val_loss: 0.2525 - val_accuracy: 0.9620
Epoch 29/50
5 - accuracy: 1.0000 - val_loss: 0.2549 - val_accuracy: 0.9657
Epoch 30/50
103/103 [=============== ] - 14s 132ms/step - Loss: 4.9241e-0
5 - accuracy: 1.0000 - val_loss: 0.2531 - val_accuracy: 0.9645
Epoch 31/50
5 - accuracy: 1.0000 - val loss: 0.2625 - val accuracy: 0.9657
Epoch 32/50
5 - accuracy: 1.0000 - val_loss: 0.2690 - val_accuracy: 0.9645
Epoch 33/50
5 - accuracy: 1.0000 - val_loss: 0.2643 - val_accuracy: 0.9632
Epoch 34/50
5 - accuracy: 1.0000 - val_loss: 0.2674 - val_accuracy: 0.9657
Epoch 35/50
5 - accuracy: 1.0000 - val_loss: 0.2649 - val_accuracy: 0.9657
Epoch 36/50
103/103 [================= ] - 14s 139ms/step - Loss: 1.5808e-0
5 - accuracy: 1.0000 - val_loss: 0.2709 - val_accuracy: 0.9632
Epoch 37/50
103/103 [================ ] - 15s 142ms/step - Loss: 1.7154e-0
5 - accuracy: 1.0000 - val_loss: 0.2751 - val_accuracy: 0.9645
Epoch 38/50
```

```
5 - accuracy: 1.0000 - val_loss: 0.2856 - val_accuracy: 0.9632
       Epoch 39/50
       103/103 [================= ] - 15s 143ms/step - Loss: 1.5362e-0
       5 - accuracy: 1.0000 - val_loss: 0.2826 - val_accuracy: 0.9632
       Epoch 40/50
       103/103 [=============== ] - 17s 168ms/step - Loss: 1.2962e-0
       5 - accuracy: 1.0000 - val_loss: 0.2853 - val_accuracy: 0.9645
       Epoch 41/50
       103/103 [================ ] - 17s 165ms/step - Loss: 1.5056e-0
       5 - accuracy: 1.0000 - val_loss: 0.2838 - val_accuracy: 0.9632
       Epoch 42/50
       103/103 [================ ] - 17s 162ms/step - Loss: 1.6323e-0
       5 - accuracy: 1.0000 - val_loss: 0.2876 - val_accuracy: 0.9645
       Epoch 43/50
       103/103 [================ ] - 16s 160ms/step - Loss: 1.5799e-0
       5 - accuracy: 1.0000 - val_loss: 0.2908 - val_accuracy: 0.9620
       Epoch 44/50
       5 - accuracy: 1.0000 - val_loss: 0.2947 - val_accuracy: 0.9632
       Epoch 45/50
       103/103 [=============== ] - 17s 166ms/step - Loss: 1.9991e-0
       5 - accuracy: 1.0000 - val_loss: 0.2951 - val_accuracy: 0.9645
       Epoch 46/50
       103/103 [================ ] - 16s 157ms/step - Loss: 1.0660e-0
       5 - accuracy: 1.0000 - val_loss: 0.3021 - val_accuracy: 0.9620
       Epoch 47/50
       6 - accuracy: 1.0000 - val_loss: 0.2995 - val_accuracy: 0.9632
       Epoch 48/50
       5 - accuracy: 1.0000 - val_loss: 0.3077 - val_accuracy: 0.9620
       Epoch 49/50
       6 - accuracy: 1.0000 - val_loss: 0.3027 - val_accuracy: 0.9632
       Epoch 50/50
       103/103 [================ ] - 17s 165ms/step - Loss: 7.8112e-0
       6 - accuracy: 1.0000 - val_loss: 0.3017 - val_accuracy: 0.9632
In [65]: results_train = model 3. evaluate(train_images, train_y)
       103/103 [============ ] - 1s 13ms/step - loss: 1.2746e-06
       - accuracy: 1.0000
In [66]: results_test = model 3. evaluate(test_i mages, test_y)
       55/55 [============= ] - 1s 13ms/step - Loss: 0.4020 - accu
       racy: 0.9475
In [67]: results_train
Out[67]: [1.2745512094625155e-06, 1.0]
```

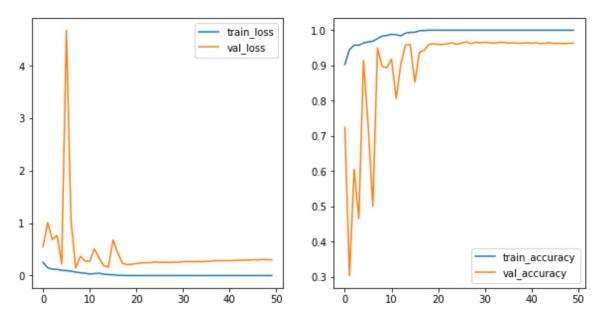
```
In [68]: results_test
```

Out [68]: [0. 4019615948200226, 0. 9474886059761047]

```
In [69]: train_loss = history3.history['loss']
    train_acc = history3.history['accuracy']
    val_loss = history3.history['val_loss']
    val_acc = history3.history['val_accuracy']

fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(10, 5))
    sns.lineplot(x=history3.epoch, y=train_loss, ax=ax1, label='train_loss')
    sns.lineplot(x=history3.epoch, y=train_acc, ax=ax2, label='train_accuracy')
    sns.lineplot(x=history3.epoch, y=val_loss, ax=ax1, label='val_loss')
    sns.lineplot(x=history3.epoch, y=val_acc, ax=ax2, label='val_accuracy')
```

Out[69]: <AxesSubpl ot: >



In [70]: #model 3. save('model s/model _3. h5')

```
In [71]: def build_cnn3():
              model 3 = model s. Sequential ()
              model 3. add(layers. Conv2D(32, (3, 3), activation='relu', input_shape=(64
          , 64,
              model 3. add(layers. MaxPooling2D((2, 2)))
              model 3. add(layers. Conv2D(32, (4, 4), activation='relu'))
              model 3. add(BatchNormalization())
              model 3. add(layers. MaxPooling2D((2, 2)))
              model 3. add(layers. Conv2D(64, (3, 3), activation='relu'))
              model 3. add(layers. MaxPooling2D((2, 2)))
              model 3. add(layers. Conv2D(128, (3, 3), activation='relu'))
              model 3. add(BatchNormalization())
              model 3. add(layers. MaxPooling2D((2, 2)))
              model 3. add(l ayers. Fl atten())
              model 3. add(layers. Dense(64, activation='relu'))
              model 3. add(Dropout(0.1))
              model 3. add(layers. Dense(1, activation='sigmoid'))
              model 3. compile(loss='binary_crossentropy',
                         opti mi zer="adam",
                         metri cs=['accuracy'])
              return model 3
```

```
In [72]: keras_model 4 = scikit_learn. KerasClassifier(build_cnn3, epochs=50,

# Integer. Number of epochs to train the model. An epoch is an iteration ov er the entire x and y data provided (unless the steps_per_epoch flag is set to something other than None). Note that in conjunction with initial_epoch, epochs is to be understood as "final epoch". The model is not trained for a number of iterations given by epochs, but merely until the epoch of index e pochs is reached.

validation_data=(valid_images, valid_y), validation_steps = validation_size)
```

Prediction for Confusion Matrix

```
In [73]: predictions = model 3. predict(x = test_i mages, steps = 10, verbose=0)
In [74]: pred_check = np. round(predictions)
```

pred_check = pred_check[:] In [75]: pred_check = pred_check.flatten() pred_check

Out[75]: array([1., 1., 1., ..., 0., 1., 0.], dtype=float32)

In [76]: test_check = test_labels[:,0] test_check

Out[76]: array([1., 1., 1., ..., 0., 0., 0.], dtype=float32)

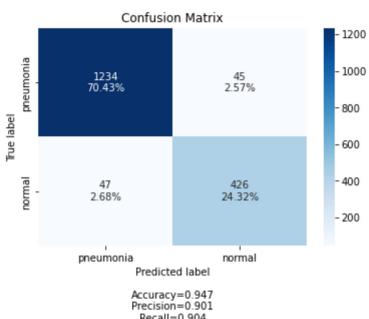
In [77]: cm = confusion_matrix(y_true=test_check, y_pred=pred_check)

In [78]: save_result(cm, 'CNN #3')

Out[78]:

	Model	RW Score	F1	Recall	Precision	Accuracy
0	CNN #2	0.909	0.910	0.909	0.911	0.951
0	CNN #3	0.904	0.903	0.904	0.901	0.947
0	CNN #1	0.866	0.874	0.864	0.884	0.931
0	baseline_model	0.861	0.856	0.863	0.850	0.923

cm_plot_labels = ['pneumonia', 'normal'] In [79]: make_confusion_matrix(cm, categories = cm_plot_labels, title='Confusion Mat rix')



Recall=0.904 F1 Score=0.903 RW Score=0.904

CNN Model 4

For this model, I will add weights

```
In [80]:
         neg = len(os.listdir('re-split data/train/normal'))
          pos = len(os.listdir('re-split_data/train/pneumonia'))
          total = neg + pos
In [81]: | # Scaling by total/2 helps keep the loss to a similar magnitude.
          # The sum of the weights of all examples stays the same.
          weight_for_0 = (1 / neg) * (total / 2.0)
          weight_for_1 = (1 / pos) * (total / 2.0)
          class_weight = {0: weight_for_0, 1: weight_for_1} #
          print('Weight for class 0: {:.2f}'.format(weight_for_0))
          print('Weight for class 1: {:.2f}'.format(weight_for_1))
          #Weight for class 0: 0.50 --- This is from https://www.tensorflow.org/tutor
          ials/structured_data/imbalanced_data
          #Weight for class 1: 289.44
         Weight for class 0: 1.85
         Weight for class 1: 0.68
In [82]: model 4 = model s. Sequential()
          model 4. add(layers. Conv2D(32, (3, 3), activation='relu', input_shape=(64, 64,
          3)))
          model 4. add(l ayers. MaxPool i ng2D((2, 2)))
          model 4. add(layers. Conv2D(32, (4, 4), activation='relu'))
          model 4. add(BatchNormalization())
          model 4. add(l ayers. MaxPool i ng2D((2, 2)))
          model 4. add(layers. Conv2D(64, (3, 3), activation='relu'))
          model 4. add(l ayers. MaxPooling2D((2, 2)))
          model 4. add(layers. Conv2D(128, (3, 3), activation='relu'))
          model 4. add(BatchNormalization())
          model 4. add(layers. MaxPooling2D((2, 2)))
          model 4. add(l avers. Fl atten())
          model 4. add(layers. Dense(64, activation='relu'))
          model 4. add(Dropout(0.1))
          model 4. add(layers. Dense(1, activation='sigmoid'))
          model 4. compile(loss='binary_crossentropy',
                        opti mi zer="adam",
                        metri cs=['accuracy'])
```

```
Epoch 1/50
103/103 [=============== ] - 17s 163ms/step - Loss: 0.2515 -
accuracy: 0.8994 - val_loss: 0.5301 - val_accuracy: 0.7966
Epoch 2/50
accuracy: 0.9389 - val_loss: 1.4481 - val_accuracy: 0.2696
accuracy: 0.9453 - val_loss: 0.3075 - val_accuracy: 0.8370
Epoch 4/50
103/103 [================ ] - 18s 172ms/step - Loss: 0.1224 -
accuracy: 0.9514 - val_loss: 0.3145 - val_accuracy: 0.8934
Epoch 5/50
accuracy: 0.9597 - val_loss: 0.2012 - val_accuracy: 0.9179
Epoch 6/50
accuracy: 0.9612 - val_loss: 0.1865 - val_accuracy: 0.9375
Epoch 7/50
103/103 [================= ] - 17s 164ms/step - Loss: 0.0882 -
accuracy: 0.9685 - val_loss: 2.7809 - val_accuracy: 0.3125
Epoch 8/50
accuracy: 0.9658 - val_loss: 0.2949 - val_accuracy: 0.8701
accuracy: 0.9774 - val_loss: 2.1309 - val_accuracy: 0.7328
Epoch 10/50
accuracy: 0.9798 - val_loss: 2.9822 - val_accuracy: 0.7304
Epoch 11/50
accuracy: 0.9762 - val_loss: 0.1526 - val_accuracy: 0.9534
Epoch 12/50
103/103 [=========== ] - 17s 162ms/step - Loss: 0.0508 -
accuracy: 0.9807 - val_loss: 0.3456 - val_accuracy: 0.8542
Epoch 13/50
accuracy: 0.9780 - val_loss: 0.2764 - val_accuracy: 0.8860
Epoch 14/50
accuracy: 0.9804 - val_loss: 0.7378 - val_accuracy: 0.6838
Epoch 15/50
accuracy: 0.9887 - val_loss: 0.1075 - val_accuracy: 0.9657
Epoch 16/50
103/103 [=========== ] - 17s 163ms/step - Loss: 0.0396 -
accuracy: 0.9859 - val_loss: 0.5260 - val_accuracy: 0.8186
Epoch 17/50
accuracy: 0.9850 - val_loss: 0.1430 - val_accuracy: 0.9534
Epoch 18/50
accuracy: 0.9890 - val_loss: 0.5616 - val_accuracy: 0.7794
Epoch 19/50
103/103 [=============== ] - 17s 166ms/step - Loss: 0.0180 -
```

```
accuracy: 0.9930 - val_loss: 2.5463 - val_accuracy: 0.7328
Epoch 20/50
103/103 [=============== ] - 17s 165ms/step - Loss: 0.0559 -
accuracy: 0.9786 - val_loss: 1.8068 - val_accuracy: 0.4547
Epoch 21/50
accuracy: 0.9814 - val_loss: 0.2521 - val_accuracy: 0.9081
Epoch 22/50
103/103 [============== ] - 18s 171ms/step - Loss: 0.0203 -
accuracy: 0.9921 - val_loss: 0.1832 - val_accuracy: 0.9473
Epoch 23/50
103/103 [============== ] - 17s 168ms/step - Loss: 0.0263 -
accuracy: 0.9911 - val_loss: 0.1444 - val_accuracy: 0.9657
Epoch 24/50
103/103 [=============== ] - 17s 166ms/step - Loss: 0.0346 -
accuracy: 0.9862 - val_loss: 7.7792 - val_accuracy: 0.7304
Epoch 25/50
103/103 [=============== ] - 16s 160ms/step - Loss: 0.0474 -
accuracy: 0.9823 - val_loss: 0.2823 - val_accuracy: 0.9142
Epoch 26/50
103/103 [============ ] - 17s 162ms/step - Loss: 0.0296 -
accuracy: 0.9887 - val_loss: 1.2618 - val_accuracy: 0.5723
Epoch 27/50
103/103 [=============== ] - 17s 162ms/step - Loss: 0.0205 -
accuracy: 0.9921 - val_loss: 0.1313 - val_accuracy: 0.9559
Epoch 28/50
103/103 [============ ] - 17s 165ms/step - Loss: 0.0187 -
accuracy: 0.9927 - val_loss: 0.5204 - val_accuracy: 0.9020
Epoch 29/50
103/103 [=============== ] - 17s 165ms/step - Loss: 0.0382 -
accuracy: 0.9859 - val_loss: 0.3172 - val_accuracy: 0.9265
Epoch 30/50
accuracy: 0.9945 - val_loss: 0.2207 - val_accuracy: 0.9375
Epoch 31/50
103/103 [=========== ] - 17s 166ms/step - Loss: 0.0119 -
accuracy: 0.9957 - val loss: 0.2333 - val accuracy: 0.9510
Epoch 32/50
103/103 [=============== ] - 16s 159ms/step - Loss: 0.0024 -
accuracy: 0.9994 - val_loss: 0.1631 - val_accuracy: 0.9694
Epoch 33/50
4 - accuracy: 1.0000 - val_loss: 0.1756 - val_accuracy: 0.9632
Epoch 34/50
103/103 [================ ] - 17s 164ms/step - Loss: 5.8622e-0
4 - accuracy: 1.0000 - val_loss: 0.1803 - val_accuracy: 0.9645
Epoch 35/50
4 - accuracy: 1.0000 - val_loss: 0.1902 - val_accuracy: 0.9669
Epoch 36/50
4 - accuracy: 1.0000 - val_loss: 0.2028 - val_accuracy: 0.9608
Epoch 37/50
4 - accuracy: 1.0000 - val_loss: 0.2226 - val_accuracy: 0.9620
Epoch 38/50
```

```
4 - accuracy: 1.0000 - val_loss: 0.2254 - val_accuracy: 0.9571
      Epoch 39/50
      4 - accuracy: 1.0000 - val_loss: 0.2243 - val_accuracy: 0.9620
      Epoch 40/50
      4 - accuracy: 1.0000 - val_loss: 0.3107 - val_accuracy: 0.9632
      Epoch 41/50
      4 - accuracy: 1.0000 - val_loss: 0.2245 - val_accuracy: 0.9608
      Epoch 42/50
      5 - accuracy: 1.0000 - val_loss: 0.2263 - val_accuracy: 0.9657
      Epoch 43/50
      5 - accuracy: 1.0000 - val_loss: 0.2320 - val_accuracy: 0.9632
      Epoch 44/50
      5 - accuracy: 1.0000 - val_loss: 0.2294 - val_accuracy: 0.9669
      Epoch 45/50
      103/103 [=============== ] - 18s 177ms/step - Loss: 7.3191e-0
      5 - accuracy: 1.0000 - val_loss: 0.2306 - val_accuracy: 0.9620
      Epoch 46/50
      5 - accuracy: 1.0000 - val_loss: 0.2369 - val_accuracy: 0.9645
      Epoch 47/50
      5 - accuracy: 1.0000 - val_loss: 0.2387 - val_accuracy: 0.9645
      Epoch 48/50
      5 - accuracy: 1.0000 - val_loss: 0.2393 - val_accuracy: 0.9645
      Epoch 49/50
      103/103 [================ ] - 18s 171ms/step - Loss: 3.0723e-0
      5 - accuracy: 1.0000 - val_loss: 0.2411 - val_accuracy: 0.9657
      Epoch 50/50
      5 - accuracy: 1.0000 - val_loss: 0.2465 - val_accuracy: 0.9632
In [84]:
     results_train = model 4. evaluate(train_images, train_y)
      103/103 [============ ] - 1s 12ms/step - loss: 8.2286e-06
      - accuracy: 1.0000
In [85]:
     results_test = model 4. evaluate(test_i mages, test_y)
      55/55 [============= ] - 1s 12ms/step - Loss: 0.3108 - accu
      racy: 0.9566
In [86]: results_train
Out[86]: [8. 228609658544883e-06, 1. 0]
```

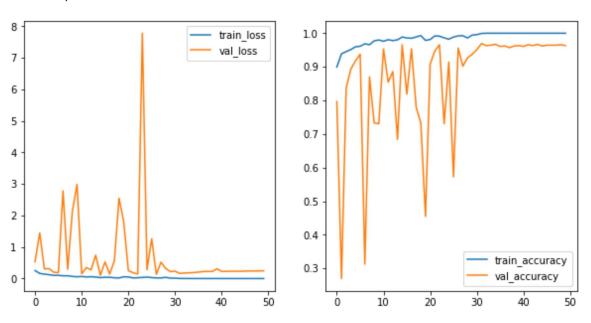
```
In [87]: results_test
```

Out[87]: [0. 3107980787754059, 0. 956620991230011]

```
In [88]: train_loss = history4.history['loss']
    train_acc = history4.history['accuracy']
    val_loss = history4.history['val_loss']
    val_acc = history4.history['val_accuracy']

fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(10, 5))
    sns.lineplot(x=history4.epoch, y=train_loss, ax=ax1, label='train_loss')
    sns.lineplot(x=history4.epoch, y=train_acc, ax=ax2, label='train_accuracy')
    sns.lineplot(x=history4.epoch, y=val_loss, ax=ax1, label='val_loss')
    sns.lineplot(x=history4.epoch, y=val_acc, ax=ax2, label='val_accuracy')
```

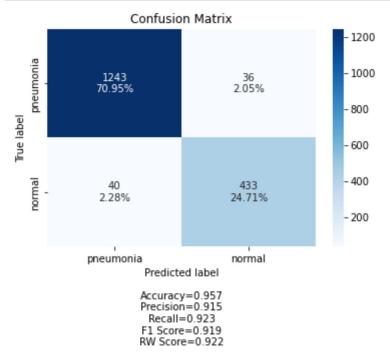
Out[88]: <AxesSubplot:>



```
In [89]:
          def build_cnn5():
              model 4 = model s. Sequential ()
              model 4. add(layers. Conv2D(32, (3, 3), activation='relu', input_shape=(64
          , 64,
              model 4. add(layers. MaxPooling2D((2, 2)))
              model 4. add(layers. Conv2D(32, (4, 4), activation='relu'))
              model 4. add(BatchNormalization())
              model 4. add(layers. MaxPooling2D((2, 2)))
              model 4. add(layers. Conv2D(64, (3, 3), activation='relu'))
              model 4. add(layers. MaxPooling2D((2, 2)))
              model 4. add(layers. Conv2D(128, (3, 3), activation='relu'))
              model 4. add(BatchNormalization())
              model 4. add(layers. MaxPooling2D((2, 2)))
              model 4. add(l ayers. Fl atten())
              model 4. add(layers. Dense(64, activation='relu'))
              model 4. add(Dropout(0.1))
              model 4. add(layers. Dense(1, activation='sigmoid'))
              model 4. compile(loss='binary_crossentropy',
                         opti mi zer="adam",
                         metri cs=['accuracy'])
              return model 4
In [90]:
          keras_model 5 = sci ki t_l earn. KerasCl assi fi er (bui l d_cnn5,
                                                         epochs=50,
                                                         validation_data=(valid_images,
          valid_y),
                                                         validation_steps = validation_s
          i ze,
                                                         class_weight = class_weight)
In [91]: predictions = model 4. predict(x = test_images, steps = 10, verbose=0)
In [92]: pred_check = np. round(predictions)
In [93]:
          pred_check = pred_check[:]
          pred_check = pred_check.flatten()
          pred_check
Out[93]: array([1., 1., 1., ..., 0., 0., 0.], dtype=float32)
In [94]: | test_check = test_labels[:,0]
          test_check
Out[94]: array([1., 1., 1., ..., 0., 0., 0.], dtype=float32)
```

```
In [95]:
           cm = confusi on_matri x(y_true=test_check, y_pred=pred_check)
In [96]:
           save_result(cm, 'CNN #4')
Out[96]:
                      Model
                              RW Score
                                               Recall
                                                      Precision Accuracy
            0
                     CNN #4
                                  0.922
                                        0.919
                                                0.923
                                                          0.915
                                                                    0.957
            0
                     CNN #2
                                        0.910
                                                0.909
                                                          0.911
                                  0.909
                                                                    0.951
            0
                     CNN #3
                                  0.904 0.903
                                                0.904
                                                          0.901
                                                                    0.947
            0
                     CNN #1
                                  0.866
                                       0.874
                                                0.864
                                                          0.884
                                                                    0.931
              baseline_model
                                  0.861
                                        0.856
                                                0.863
                                                          0.850
                                                                    0.923
```

In [97]: cm_plot_labels = ['pneumonia', 'normal']
 make_confusion_matrix(cm, categories = cm_plot_labels, title='Confusion Mat
 rix')



CNN Model 5: Model 3 with added layer

After observing how the first five models ran, CNN #2 was the best model due to low standard deviation for the cross validation and high test and train accuracy. I will try adding dropout to help the little bit of overtraining that is occurring

```
In [98]:
          model 6 = model s. Sequential ()
          model 6. add(layers. Conv2D(32, (3, 3), activation='relu', input_shape=(64,64,
          3)))
          model 6. add(l ayers. MaxPool i ng2D((2, 2)))
          model 6. add(layers. Conv2D(32, (4, 4), activation='relu'))
          model 6. add(BatchNormalization())
          model 6. add(layers. MaxPooling2D((2, 2)))
          model 6. add(layers. Conv2D(64, (3, 3), activation='relu'))
          model 6. add(BatchNormalization())
          model 6. add(layers. MaxPooling2D((2, 2)))
          model 6. add(layers. Conv2D(96, (3, 3), activation='relu', padding='same')) #
          model 6. add(BatchNormalization())
                                                                                        # n
          model 6. add(Dropout(0.1))
                                                                                        # n
          model 6. add(l ayers. MaxPool i ng2D((2, 2)))
          # new
          model 6. add(layers. Conv2D(128, (3, 3), activation='relu', padding='same'))
          model 6. add(BatchNormalization())
          model 6. add(layers. MaxPooling2D((2, 2)))
          model 6. add(l ayers. Fl atten())
          model 6. add(layers. Dense(64, activation='relu'))
          model 6. add(Dropout (0. 1))
          model 6. add(layers. Dense(1, activation='sigmoid'))
          model 6. compile(loss='binary_crossentropy',
                         opti mi zer="adam",
                         metri cs=['accuracy'])
```

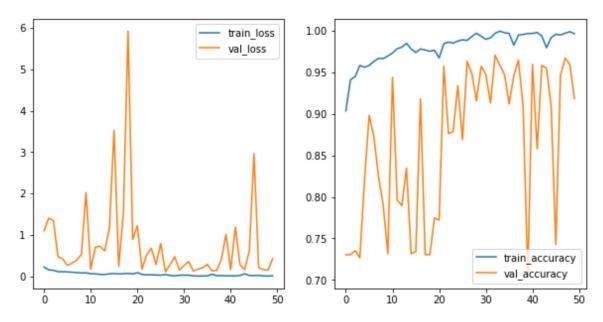
```
Epoch 1/50
103/103 [=============== ] - 15s 144ms/step - Loss: 0.2241 -
accuracy: 0.9034 - val_loss: 1.0929 - val_accuracy: 0.7304
Epoch 2/50
accuracy: 0.9410 - val_loss: 1.4036 - val_accuracy: 0.7304
accuracy: 0.9447 - val_loss: 1.3484 - val_accuracy: 0.7353
Epoch 4/50
103/103 [================ ] - 15s 143ms/step - Loss: 0.1083 -
accuracy: 0.9581 - val_loss: 0.4689 - val_accuracy: 0.7267
Epoch 5/50
accuracy: 0.9560 - val_loss: 0.4136 - val_accuracy: 0.8199
Epoch 6/50
accuracy: 0.9581 - val_loss: 0.2618 - val_accuracy: 0.8983
Epoch 7/50
accuracy: 0.9630 - val_loss: 0.3147 - val_accuracy: 0.8725
Epoch 8/50
accuracy: 0.9667 - val_loss: 0.3895 - val_accuracy: 0.8248
accuracy: 0.9664 - val_loss: 0.5231 - val_accuracy: 0.7904
Epoch 10/50
accuracy: 0.9694 - val_loss: 2.0211 - val_accuracy: 0.7316
Epoch 11/50
accuracy: 0.9731 - val_loss: 0.1677 - val_accuracy: 0.9436
Epoch 12/50
103/103 [============ ] - 15s 145ms/step - Loss: 0.0552 -
accuracy: 0.9783 - val_loss: 0.7018 - val_accuracy: 0.7966
Epoch 13/50
accuracy: 0.9801 - val_loss: 0.7240 - val_accuracy: 0.7892
Epoch 14/50
accuracy: 0.9847 - val_loss: 0.6096 - val_accuracy: 0.8346
Epoch 15/50
accuracy: 0.9780 - val_loss: 1.1437 - val_accuracy: 0.7316
Epoch 16/50
103/103 [=========== ] - 15s 147ms/step - Loss: 0.0628 -
accuracy: 0.9737 - val_loss: 3.5257 - val_accuracy: 0.7341
Epoch 17/50
accuracy: 0.9780 - val_loss: 0.2381 - val_accuracy: 0.9179
Epoch 18/50
accuracy: 0.9768 - val_loss: 1.5093 - val_accuracy: 0.7304
Epoch 19/50
103/103 [=============== ] - 16s 154ms/step - Loss: 0.0677 -
```

```
accuracy: 0.9752 - val_loss: 5.9237 - val_accuracy: 0.7304
Epoch 20/50
103/103 [=============== ] - 16s 151ms/step - Loss: 0.0545 -
accuracy: 0.9765 - val_loss: 0.8913 - val_accuracy: 0.7745
Epoch 21/50
accuracy: 0.9673 - val_loss: 1.2214 - val_accuracy: 0.7721
Epoch 22/50
103/103 [=============== ] - 16s 152ms/step - Loss: 0.0431 -
accuracy: 0.9844 - val_loss: 0.1675 - val_accuracy: 0.9571
Epoch 23/50
accuracy: 0.9862 - val_loss: 0.5166 - val_accuracy: 0.8762
Epoch 24/50
103/103 [=============== ] - 16s 154ms/step - Loss: 0.0354 -
accuracy: 0.9850 - val_loss: 0.6751 - val_accuracy: 0.8787
Epoch 25/50
103/103 [============ ] - 16s 151ms/step - Loss: 0.0300 -
accuracy: 0.9875 - val_loss: 0.2796 - val_accuracy: 0.9338
Epoch 26/50
103/103 [============ ] - 16s 155ms/step - Loss: 0.0232 -
accuracy: 0.9890 - val_loss: 0.7921 - val_accuracy: 0.8689
Epoch 27/50
103/103 [=============== ] - 16s 157ms/step - Loss: 0.0404 -
accuracy: 0.9884 - val_loss: 0.1049 - val_accuracy: 0.9632
Epoch 28/50
103/103 [============ ] - 15s 148ms/step - Loss: 0.0170 -
accuracy: 0.9930 - val_loss: 0.2773 - val_accuracy: 0.9485
Epoch 29/50
103/103 [=============== ] - 16s 161ms/step - Loss: 0.0086 -
accuracy: 0.9969 - val_loss: 0.4706 - val_accuracy: 0.9154
Epoch 30/50
accuracy: 0.9933 - val_loss: 0.1458 - val_accuracy: 0.9571
Epoch 31/50
103/103 [================ ] - 16s 151ms/step - Loss: 0.0254 -
accuracy: 0.9896 - val loss: 0.2570 - val accuracy: 0.9473
Epoch 32/50
accuracy: 0.9914 - val_loss: 0.3531 - val_accuracy: 0.9130
Epoch 33/50
103/103 [=========== ] - 16s 152ms/step - Loss: 0.0080 -
accuracy: 0.9969 - val_loss: 0.1255 - val_accuracy: 0.9706
Epoch 34/50
accuracy: 0.9994 - val_loss: 0.1689 - val_accuracy: 0.9583
Epoch 35/50
103/103 [=============== ] - 15s 144ms/step - Loss: 0.0048 -
accuracy: 0.9976 - val_loss: 0.2054 - val_accuracy: 0.9473
Epoch 36/50
103/103 [=============== ] - 16s 152ms/step - Loss: 0.0078 -
accuracy: 0.9966 - val_loss: 0.2838 - val_accuracy: 0.9118
Epoch 37/50
103/103 [============ ] - 17s 167ms/step - Loss: 0.0464 -
accuracy: 0.9826 - val_loss: 0.1300 - val_accuracy: 0.9461
Epoch 38/50
```

```
accuracy: 0.9948 - val_loss: 0.1358 - val_accuracy: 0.9645
       Epoch 39/50
       103/103 [================ ] - 16s 152ms/step - Loss: 0.0137 -
       accuracy: 0.9954 - val_loss: 0.4000 - val_accuracy: 0.9081
       Epoch 40/50
       accuracy: 0.9966 - val_loss: 1.0111 - val_accuracy: 0.7047
       Epoch 41/50
       103/103 [=============== ] - 16s 154ms/step - Loss: 0.0082 -
       accuracy: 0.9966 - val_loss: 0.1587 - val_accuracy: 0.9596
       Epoch 42/50
       accuracy: 0.9979 - val_loss: 1.1864 - val_accuracy: 0.8578
       Epoch 43/50
       103/103 [================ ] - 15s 145ms/step - Loss: 0.0167 -
       accuracy: 0.9936 - val_loss: 0.2773 - val_accuracy: 0.9583
       Epoch 44/50
       103/103 [============ ] - 16s 155ms/step - Loss: 0.0563 -
       accuracy: 0.9795 - val_loss: 0.1572 - val_accuracy: 0.9547
       Epoch 45/50
       103/103 [=============== ] - 16s 151ms/step - Loss: 0.0167 -
       accuracy: 0.9917 - val_loss: 0.6077 - val_accuracy: 0.9093
       Epoch 46/50
       accuracy: 0.9957 - val_loss: 2.9611 - val_accuracy: 0.7426
       Epoch 47/50
       accuracy: 0.9948 - val_loss: 0.2072 - val_accuracy: 0.9461
       Epoch 48/50
       103/103 [================ ] - 16s 158ms/step - Loss: 0.0080 -
       accuracy: 0.9969 - val_loss: 0.1592 - val_accuracy: 0.9669
       Epoch 49/50
       103/103 [=============== ] - 16s 159ms/step - Loss: 0.0028 -
       accuracy: 0.9988 - val_loss: 0.1479 - val_accuracy: 0.9596
       Epoch 50/50
       accuracy: 0.9963 - val_loss: 0.4236 - val_accuracy: 0.9179
In [100]:
       results_train = model 6. evaluate(train_images, train_y)
       curacy: 0.9364
In [101]: results_test = model 6. evaluate(test_images, test_y)
       55/55 [============= ] - 1s 12ms/step - Loss: 0.4768 - accu
       racy: 0.9138
```

In [102]: train_loss = history6.history['loss'] train_acc = history6.history['accuracy'] val_loss = history6.history['val_loss'] val_acc = history6.history['val_accuracy'] fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(10, 5)) sns.lineplot(x=history6.epoch, y=train_loss, ax=ax1, label='train_loss') sns.lineplot(x=history6.epoch, y=train_acc, ax=ax2, label='train_accuracy') sns.lineplot(x=history6.epoch, y=val_loss, ax=ax1, label='val_loss') sns.lineplot(x=history6.epoch, y=val_acc, ax=ax2, label='val_accuracy')

Out[102]: <AxesSubplot:>



```
In [103]: def build_cnn6():
               model 6 = model s. Sequential ()
               model 6. add(layers. Conv2D(32, (3, 3), activation='relu', input_shape=(64
           , 64,
               model 6. add(layers. MaxPooling2D((2, 2)))
               model 6. add(layers. Conv2D(32, (4, 4), activation='relu'))
               model 6. add(BatchNormalization())
               model 6. add(layers. MaxPooling2D((2, 2)))
               model 6. add(layers. Conv2D(64, (3, 3), activation='relu'))
               model 6. add(BatchNormalization())
               model 6. add(layers. MaxPooling2D((2, 2)))
               model 6. add(layers. Conv2D(96, (3, 3), activation='relu', padding='same
           ')) # new
               model 6. add(BatchNormalization())
           # new
               model 6. add(Dropout(0.1))
           # new
               model 6. add(layers. MaxPooling2D((2, 2)))
           # new
               model 6. add(layers. Conv2D(128, (3, 3), activation='relu', padding='same
           '))
               model 6. add(BatchNormalization())
               model 6. add(layers. MaxPooling2D((2, 2)))
               model 6. add(layers. Flatten())
               model 6. add(layers. Dense(64, activation='relu'))
               model 6. add(Dropout(0.1))
               model 6. add(l ayers. Dense(1, activation='sigmoid'))
               model 6. compile(loss='binary_crossentropy',
                          optimizer="adam",
                          metri cs=['accuracy'])
               return model 6
In [104]:
           keras_model 6 = sci ki t_l earn. KerasCl assi fi er(build_cnn6,
                                                          epochs=50,
                                                          validation_data=(valid_images,
           valid_y),
                                                          validation_steps = validation_s
           i ze,
                                                          class_weight = class_weight)
```

Prediction for Confusion Matrix

```
In [105]: predictions = model 6. predict(x = test_i mages, steps = 10, verbose=0)
```

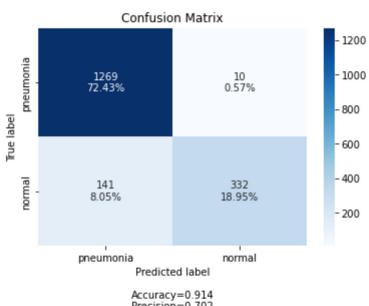
```
In [106]: pred_check = np. round(predictions)
In [107]:
          pred_check = pred_check[:]
          pred_check = pred_check.flatten()
          pred_check
Out[107]:
          array([0., 1., 0., ..., 0., 0., 0.], dtype=float32)
In [108]:
          test_check = test_labels[:,0]
          test_check
Out[108]:
          array([1., 1., 1., ..., 0., 0., 0.], dtype=float32)
In [109]:
          cm = confusi on_matri x(y_true=test_check, y_pred=pred_check)
```

In [110]: save_result(cm, 'CNN #6')

Out[110]:

	Model	RW Score	F1	Recall	Precision	Accuracy
0	CNN #4	0.922	0.919	0.923	0.915	0.957
0	CNN #2	0.909	0.910	0.909	0.911	0.951
0	CNN #3	0.904	0.903	0.904	0.901	0.947
0	CNN #1	0.866	0.874	0.864	0.884	0.931
0	baseline_model	0.861	0.856	0.863	0.850	0.923
0	CNN #6	0.935	0.815	0.971	0.702	0.914

cm_plot_labels = ['pneumonia', 'normal'] In [111]: make_confusion_matrix(cm, categories = cm_plot_labels, title='Confusion Mat rix')



Precision=0.702 Recall=0.971 F1 Score=0.815 RW Score=0.935

VGG Model

(32, 1)

```
In [237]:
          train_path = 're-split_data/train'
          valid_path = 're-split_data/validation'
           test_path = 're-split_data/test'
          train_batches2 = ImageDataGenerator(preprocessing_function=tf.keras.applica
In [238]:
          tions.vgg16.preprocess_input) \
               .flow_from_directory(directory=train_path, target_size=(224, 224), batc
          h si ze=32)
          valid_batches2 = ImageDataGenerator(preprocessing_function=tf.keras.applica
          tions.vgg16.preprocess_input) \
               .flow_from_directory(directory=valid_path, target_size=(224, 224),
                                                                                     bat
          ch_si ze=32)
          test_batches2 = ImageDataGenerator(preprocessing_function=tf.keras.applicat
          ions. vgg16. preprocess_i nput) \
               .flow_from_directory(directory=test_path, target_size=(224, 224),
          h_si ze=32, shuffl e=Fal se)
          Found 3272 images belonging to 2 classes.
          Found 816 images belonging to 2 classes.
          Found 1752 images belonging to 2 classes.
In [239]:
          # create the data sets
          train_i mages2, train_labels2 = next(train_batches2)
           test_i mages2, test_l abel s2 = next(test_batches2)
          valid_images2, valid_labels2 = next(valid_batches2)
          train_i mg2 = train_i mages2. reshape(train_i mages2. shape[0], -1)
          test img2 = test images2.reshape(test images2.shape[0], -1)
          valid_img2 = valid_images2.reshape(valid_images2.shape[0], -1)
          print(train_i mg2. shape)
          print(test_i mg2. shape)
          print(valid_img2.shape)
           (32, 150528)
           (32, 150528)
           (32, 150528)
In [240]:
          train_y2 = np. reshape(train_labels2[:,0], (32,1))
          test_y2 = np. reshape(test_labels2[:, 0], (32, 1))
          valid_y2 = np. reshape(valid_labels2[:, 0], (32, 1))
          print(train_y2. shape)
          print(test y2. shape)
          print(valid_y2. shape)
           (32, 1)
           (32, 1)
```

Out[241]: "\ntrain_generator = ImageDataGenerator(rescal e=1./255).flow_from_directory ('re-split_data/train',\n target_size=(64, 64), batch_size = train_s ize) \n\ntest_generator = ImageDataGenerator(rescal e=1./255).flow_from_directory('re-split_data/test',\n target_size=(64, 64), batch_size = test _size, shuffle= False) \n\nvalid_generator = ImageDataGenerator(rescal e=1./2 55).flow_from_directory('re-split_data/validation',\n target_size=(64, 64), batch_size = validation_size)\n"

```
In [242]: # create the data sets
    train_images2, train_labels2 = next(train_generator)
    test_images2, test_labels2 = next(test_generator)
    valid_images2, valid_labels2 = next(valid_generator)
    valid_images2, valid_labels2 = next(valid_generator)
```

In [243]: show_i mages(train_i mages2)

Clipping input data to the valid range for imshow with RGB data ([0..1] for floats or [0..255] for integers). Clipping input data to the valid range for imshow with RGB data ([0..1] for

floats or [0..255] for integers).

Clipping input data to the valid range for imshow with RGB data ([0..1] for floats or [0..255] for integers).

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Clipping input data to the valid range for imshow with RGB data ([0..1] for floats or [0..255] for integers).

Clipping input data to the valid range for imshow with RGB data ([0..1] for floats or [0..255] for integers).



```
In [244]:
           train_img2 = train_images2.reshape(train_images2.shape[0], -1)
           test_img2 = test_images2. reshape(test_images2. shape[0], -1)
           valid_img2 = valid_images2.reshape(valid_images2.shape[0], -1)
           pri nt(trai n_i mg2. shape)
           print(test_i mg2. shape)
           print(valid_img2.shape)
```

Out[244]:

'\ntrain_i mg2 = train_i mages2. reshape(train_i mages2. shape[0], -1)\ntest_i mg 2 = test_i mages2. reshape(test_i mages2. shape[0], -1)\nvalid_i mg2 = valid_i ma ges2. reshape(valid_i mages2. shape[0], -1)\n\nprint(train_i mg2. shape)\nprint (test_i mg2. shape)\npri nt(val i d_i mg2. shape)\n'

```
In [245]:
             train_y2 = np. reshape(train_labels2[:,0], (train_size, 1))
             test_y2 = np. reshape(test_labels2[:, 0], (test_size, 1))
             valid_y2 = np. reshape(valid_labels2[:,0], (validation_size, 1))
             print(train_y2. shape)
             pri nt (test_y2. shape)
             pri nt (val i d_y2. shape)
  Out[245]:
            ' ntrain_y2 = np. reshape(train_labels2[:, 0], (train_size, 1)) ntest_y2 = np.
             reshape(test_labels2[:,0], (test_size,1))\nvalid_y2 = np.reshape(valid_labe
             Is2[:,0], (validation\_size,1))\n\print(train\_y2.shape)\nprint(test\_y2.shape)
             e)\npri nt (val i d_y2. shape)\n'
             vgg16_model = tf. keras. applications. vgg16. VGG16()
  In [246]:
vgg16_model.summary()
  In [247]: # create new model of type sequential, then iterate over each of layers in
             vgg model (save last),
             # add layers to sequential.
             #model_VGG = Sequential()
             #for layer in vgg16_model.layers[:-1]:
                # model_VGG. add(layer)
  In [248]:
            #for layer in model_VGG.layers:
                # layer. trainable = False
  In [249]: | #model_VGG. add(Dense(units=2, activation='sigmoid'))
             #model_VGG.compile(optimizer=Adam(learning_rate=0.001), loss='binary_crosse
  In [250]:
             ntropy', metrics=['accuracy'])
             #vgg_hist = model_VGG.fit(x=train_batches,
  In [251]:
                                       steps_per_epoch = len(train_batches),
              #
                                       val i dati on_data=val i d_batches,
                                       validation_steps = len(valid_batches),
               #
                                       epochs=7)
  In [252]:
             #os. mkdi r(' model s')
  In [253]:
             #model_VGG. save('model s/VGG_model . h5')
  In [254]:
             # Load model
             from tensorflow.keras.models import load_model
             new_model = load_model('models/VGG_model.h5')
  In [255]: model_VGG = new_model
```

```
In [256]:
            test_i mages2, test_l abel s2= next(test_batches2)
In [257]:
            predictions = model_VGG.predict(x=test_batches2, steps=len(test_batches2),
            verbose = 0)
In [258]:
            y_pred = np. argmax(predictions, axis = 1)
In [259]:
            y_true = test_batches2. classes
In [260]:
            # Confusion Matrix
            cm = confusion_matrix(y_true = y_true, y_pred = y_pred)
            CM
Out[260]:
            array([[ 421,
                              52],
                      40, 1239]], dtype=i nt64)
In [261]:
            cm_plot_labels = ['normal', 'pneumonia']
            make_confusion_matrix(cm, categories=cm_plot_labels, title='Confusion Matri
            x')
                              Confusion Matrix
                                                              - 1200
                                                              - 1000
               normal
                                             52
2.97%
                          421
                         24.03%
                                                              800
            True label
                                                              600
               pneumonia
                                                              - 400
                                             1239
                          40
                         2.28%
                                             70.72%
                                                              - 200
                                           pneumonia
                         normal
                                Predicted label
                               Accuracy=0.947
                               Precision=0.969
                                Recall=0.960
                               F1 Score=0.964
                               RW Score=0.961
In [262]:
           # Sani ty Check
            tp = cm[1, 1]
            fn = cm[0, 1]
            recall = tp / (tp + fn)
            recal I
```

Out[262]: 0. 959721146398141

Expaining The Final Model With LIME

```
In [263]: import lime
from lime import lime_image
In [264]: explainer = lime_image.LimeImageExplainer()
```

Here, I am checking the labels and predictions of various images, then viewing the decision-making weights using LIME Explinations.

An explanation is a local linear approximation of the models behavior. While the model may be complex globally, it is simple(r) to approximate it around the vicinity of a particular insance (github.com/marcotcr/lime)

```
In [265]:
           from skimage.segmentation import mark_boundaries
In [266]:
           def check_label(train_images_num):
               img1 = train_images[train_images_num]
               img = train_labels[train_images_num][0]
               if imq == 0:
                   return 'pneumonia'
               el se:
                   return 'normal'
In [268]:
           check_l abel (0)
Out [268]: 'pneumoni a'
In [271]:
          img = train_images[3].reshape(1,64, 64, 3)
           pred = model 4. predict(i mg)
           pred = np. round(pred)
           pred
Out[271]: array([[0.]], dtype=float32)
           img = train_i mages[20]. reshape(1, 64, 64, 3)
In [273]:
           pred = model 4. predict(img)
           pred
Out[273]: array([[6.1722814e-11]], dtype=float32)
In [274]: def check_prediction(train_images_num):
               img = train_images[train_images_num].reshape(1,64,64,3)
               pred = model 4. predict(img)
               if pred[0][0] < 0.5:
                   return 'pneumonia'
               el se:
                   return 'normal'
```

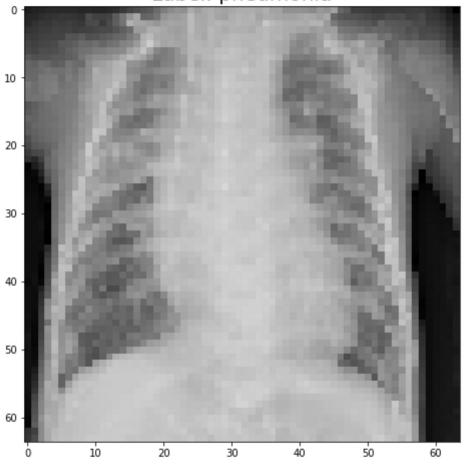
```
In [275]: def plot_image_preds(train_images_num):
               img = train_images[train_images_num].reshape(1,64,64,3)
              pred = model 4. predict(i mg)
              f, ax = plt.subplots(figsize=(10,8))
              plt.imshow(img[0])
              plt.title('Prediction: ' + check_prediction(train_images_num) + '\n' +
          'Label: ' + check_label(train_images_num), fontsize=20)
              plt.show()
In [276]: def plot_explanation(exp):
              temp, mask = exp.get_image_and_mask(exp.top_labels[0], positive_only=Tr
          ue, hi de_rest=Fal se)
              f, ax = plt.subplots(figsize=(8, 8))
              plt.title('Explanation for classification', fontsize=20)
              plt.imshow(mark_boundaries(temp / 2 + 0.5, mask))
In [277]: # Plots the positive and negative explanations for the label
          def plot_pos_neg_explanation(exp):
              temp, mask = exp.get_image_and_mask(exp.top_labels[0], positive_only=Fa
          lse, hide_rest=False)
              f, ax = plt.subplots(figsize=(8, 8))
              plt.title('Explanation for classification', fontsize=20)
              plt.imshow(mark_boundaries(temp / 2 + 0.5, mask))
In [278]: #Plots explanation with minimum weights
          def plot_with_weights(exp, mw): # mw is the minimum weight
              temp, mask = exp.get_i mage_and_mask(exp. top_l abels[0], positive_only=Fa
          lse, hide_rest=False, min_weight = mw)
              f, ax = plt.subplots(figsize=(8, 8))
              plt.title('Explanation - with weights - for classification', fontsize=
          18)
              plt.imshow(mark_boundaries(temp / 2 + 0.5, mask))
In [279]:
          # Plots the heatmap of the explanation
          def plot_explanation_heatmap(exp):
              ind = exp. top_labels[0]
               #Map each explanation weight to the corresponding superpixel
              dict_heatmap = dict(exp.local_exp[ind])
              heatmap = np. vectori ze(dict_heatmap.get) (exp. segments)
              f, ax = plt.subplots(figsize=(8, 8))
              #Plot. The visualization makes more sense if a symmetrical colorbar is
          used.
              plt.imshow(heatmap, cmap = 'RdBu', vmin = -heatmap.max(), vmax = heatm
          ap. max())
              plt.title('Heatmap for classification', fontsize=20)
              plt.colorbar()
```

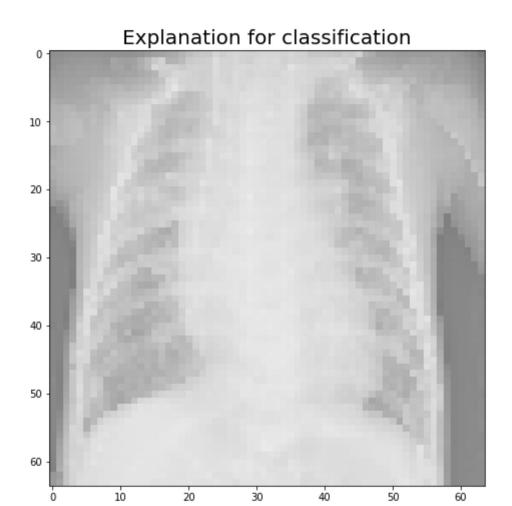
In [285]: # Hide color is the color for a superpixel turned OFF. Alternatively, if it is NONE, the superpixel will be replaced by the average of its pixels expl anation1 = expl ai ner. expl ai n_i nstance(trai n_i mages[1]. astype("double"), model 4. predict, top_label s=2, hide_color=0, num_samples=10000)

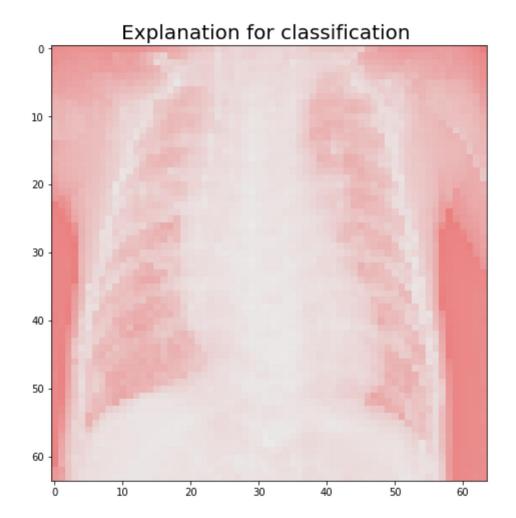
100%| 100%| 10000/10000 [00: 45<00: 00, 221. 94i t/s]

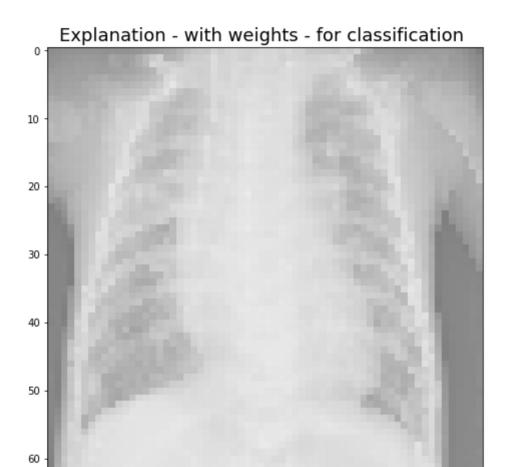
```
In [286]: plot_i mage_preds(1)
plot_expl anati on(expl anati on1)
plot_pos_neg_expl anati on(expl anati on1)
plot_wi th_wei ghts(expl anati on1, 0.5)
plot_expl anati on_heatmap(expl anati on1)
```

Prediction: pneumonia Label: pneumonia









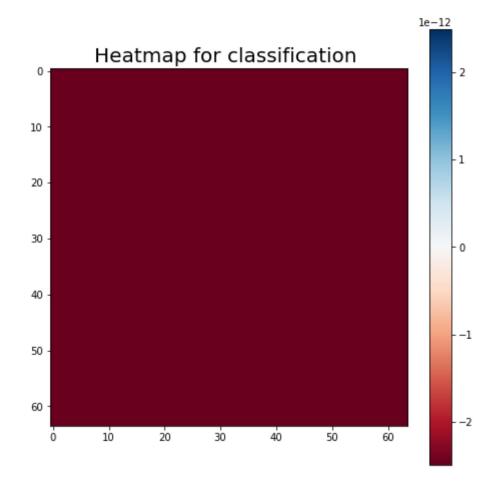
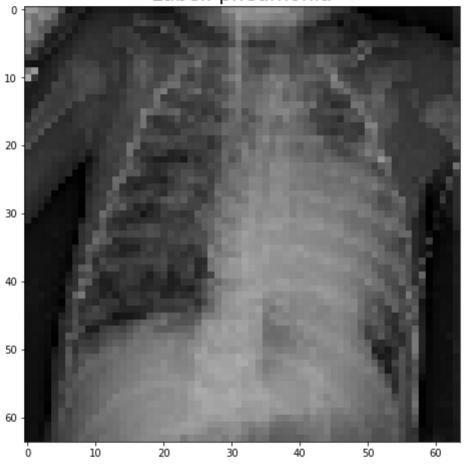


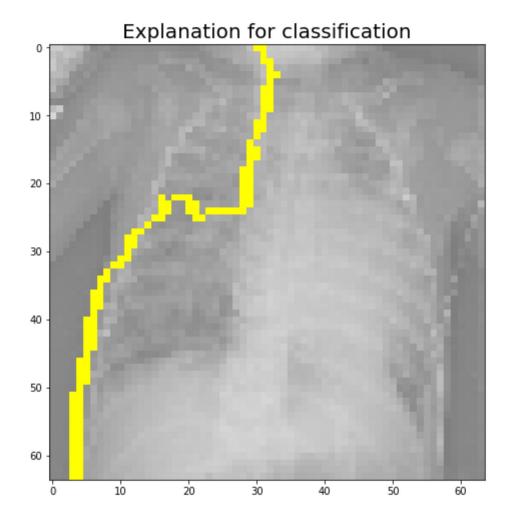
Image 2

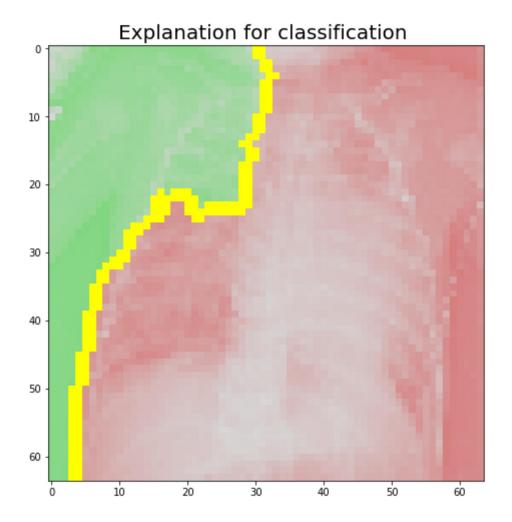
In []: # Hide color is the color for a superpixel turned OFF. Alternatively, if it is NONE, the superpixel will be replaced by the average of its pixels explanation2 = explainer.explain_instance(train_images[12].astype("double"), model 4. predict, top_labels = 2, hide_color=0, num_samples=10000)

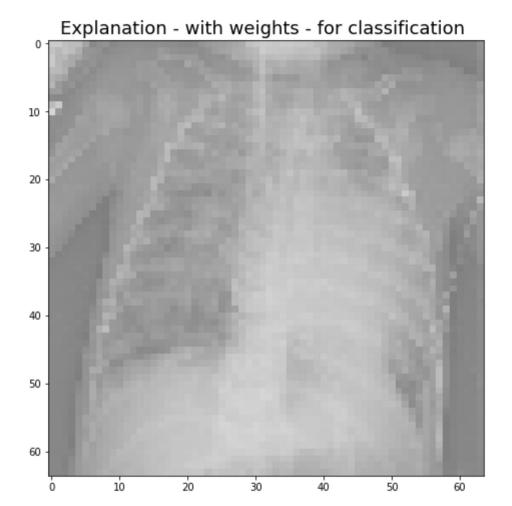
```
In []: plot_i mage_preds(12)
    plot_expl anati on(expl anati on2)
    plot_pos_neg_expl anati on(expl anati on2)
    plot_wi th_wei ghts(expl anati on2, 0.2)
    plot_expl anati on_heatmap(expl anati on2)
```

Prediction: pneumonia Label: pneumonia









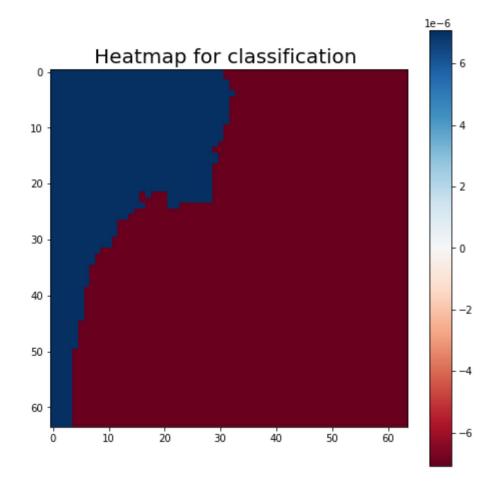
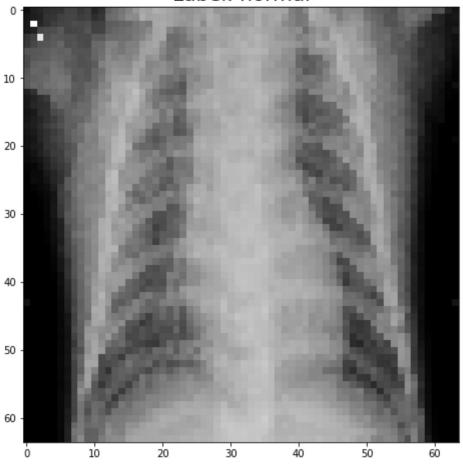


Image 3: A Positive Image

The first one I find

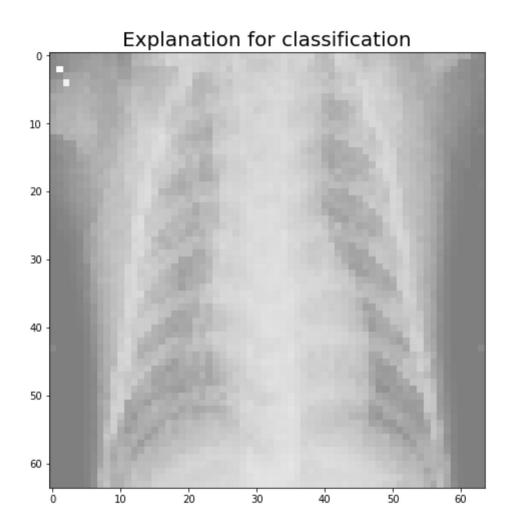
Prediction: normal Label: normal

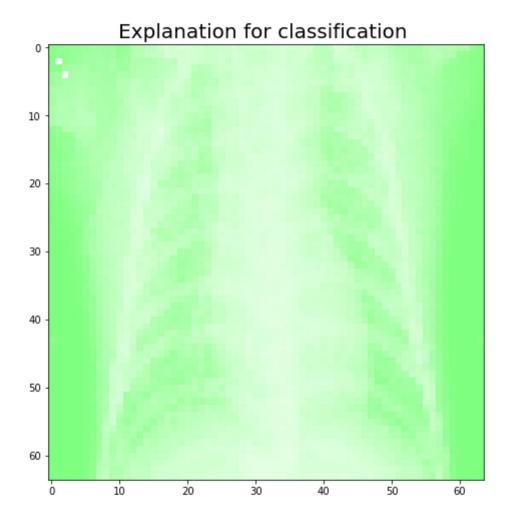


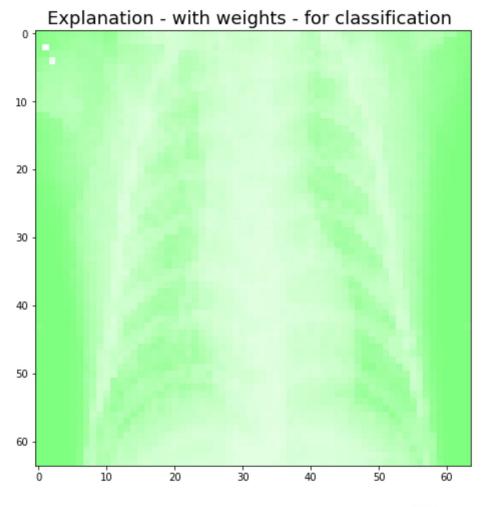
In []: explanation3 = explainer.explain_instance(train_images[723].astype("double"), model 4. predict, top_labels = 2, hide_color=0, num_samples=10000)

100%|| 100%|| 10000/10000 [00: 44<00: 00, 225. 18i t/s]

```
In []: plot_expl anation(expl anation3)
plot_pos_neg_expl anation(expl anation3)
plot_with_weights(expl anation3, 0.2)
plot_expl anation_heatmap(expl anation3)
```







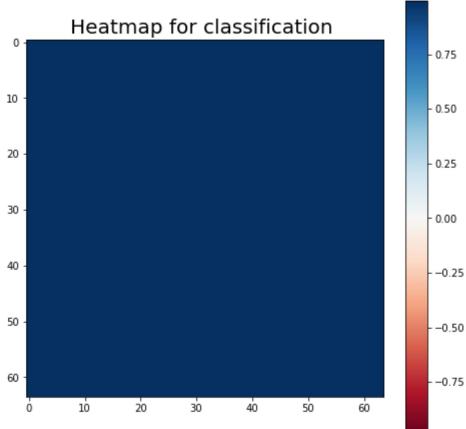
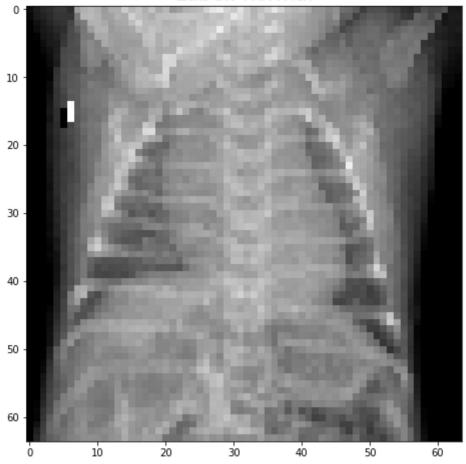
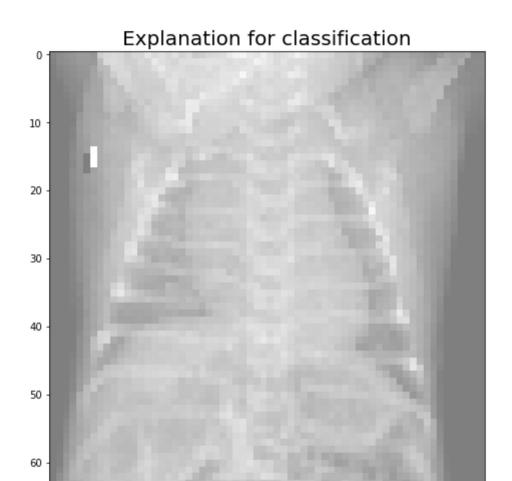


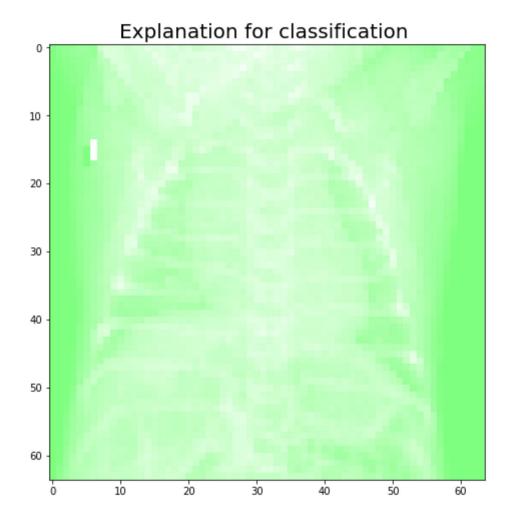
Image 4

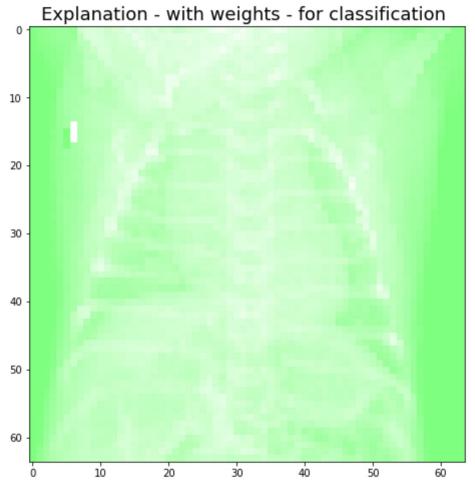
Prediction: normal Label: normal



```
In []: plot_expl anation(expl anation4)
plot_pos_neg_expl anation(expl anation4)
plot_with_weights(expl anation4, 0.2)
plot_expl anation_heatmap(expl anation4)
```







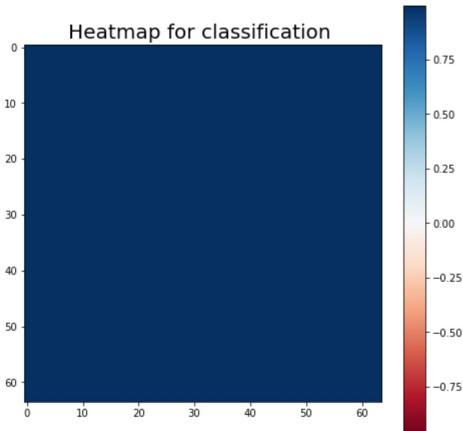
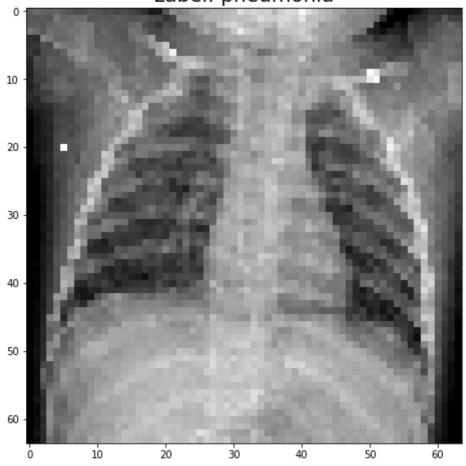


Image 5

In []: plot_i mage_preds(233)

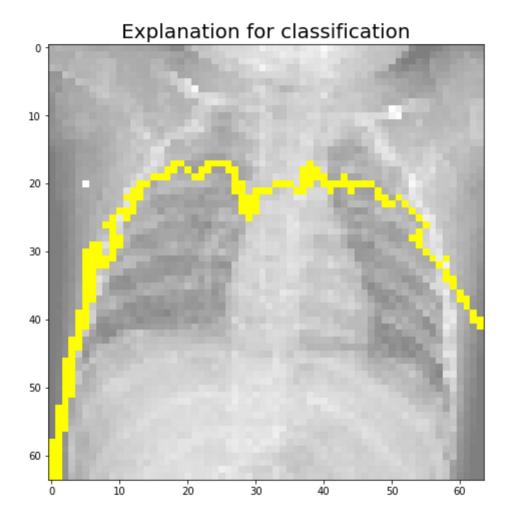
Prediction: pneumonia Label: pneumonia

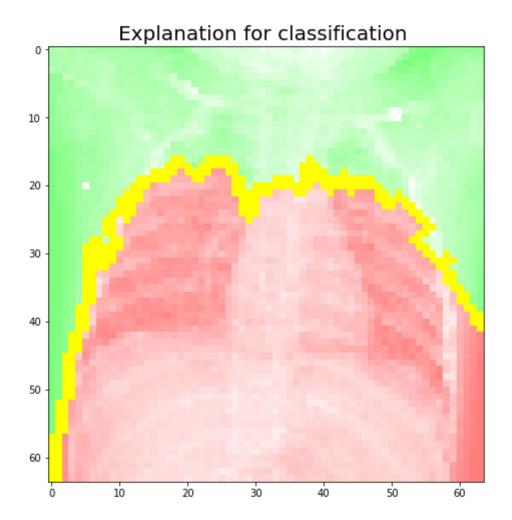


In []: explanation5 = explainer.explain_instance(train_images[233].astype("doubl
e"), model 4. predict, top_labels = 2, hide_color=0, num_samples=10000)

100%|| 100%|| 10000/10000 [00: 42<00: 00, 235. 76i t/s]

```
In []: plot_expl anation(expl anation5)
plot_pos_neg_expl anation(expl anation5)
plot_with_weights(expl anation5, 0.2)
plot_expl anation_heatmap(expl anation5)
```





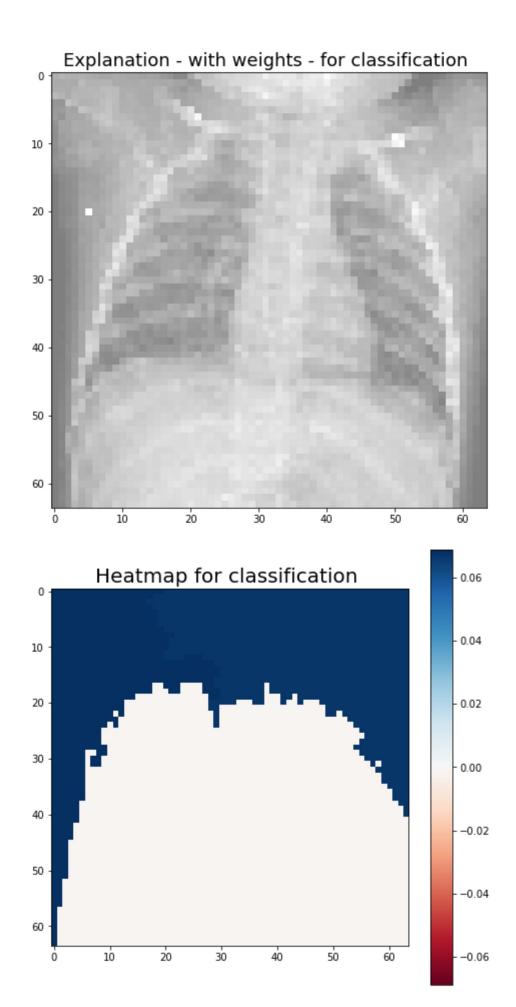
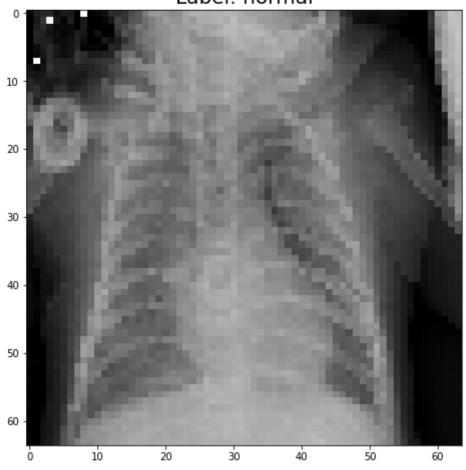


Image 6

In []: plot_i mage_preds(414)

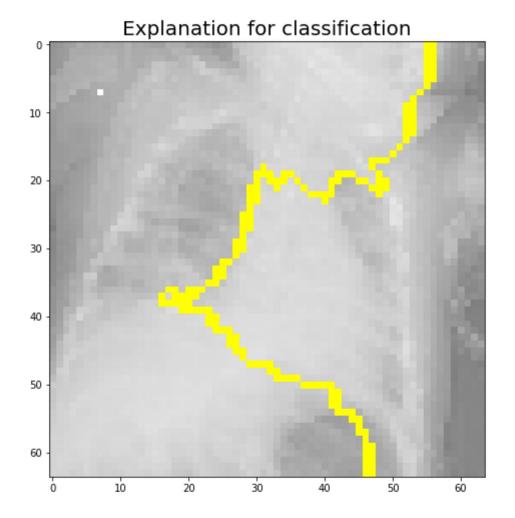
Prediction: normal Label: normal

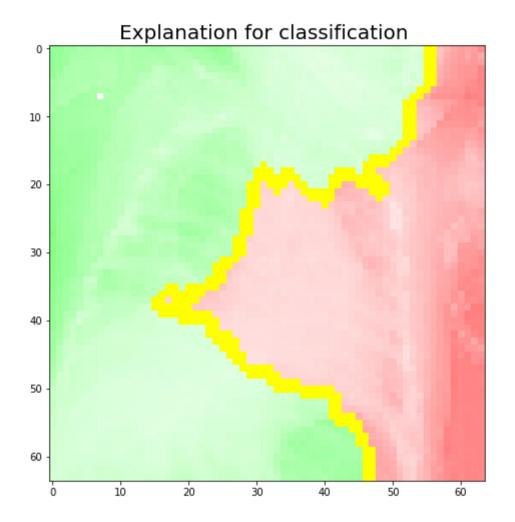


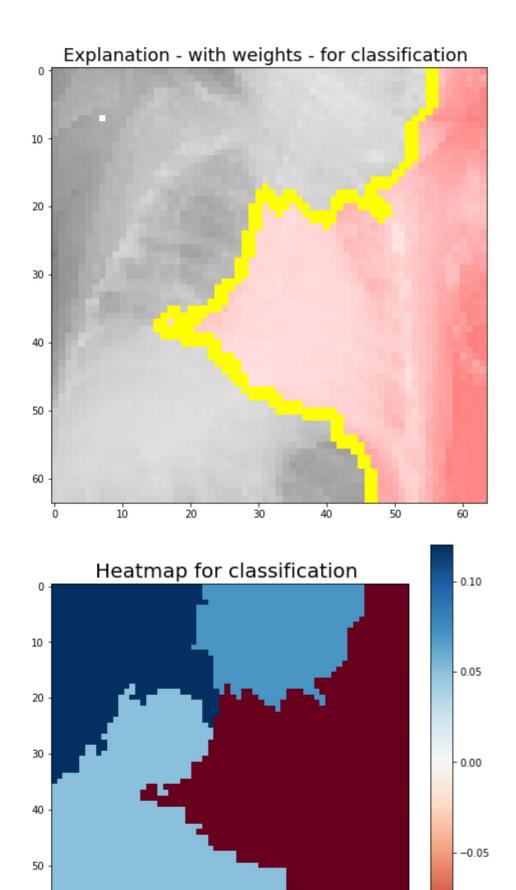
In []: # Image 123 is pneumonia
 expl anati on6 = expl ai ner. expl ai n_i nstance(trai n_i mages[123]. astype("doubl
e"), model 4. predict, top_l abels = 2, hi de_col or=0, num_sampl es=10000)

100%| 100%| 10000/10000 [00:41<00:00, 240.08it/s]

```
In []: plot_expl anation(expl anation6)
plot_pos_neg_expl anation(expl anation6)
plot_with_weights(expl anation6, 0.2)
plot_expl anation_heatmap(expl anation6)
```







-0.10

Conclusion

The VGG Model had the best F-beta score, making it the best model for this project.

Final Results

The final result included 1,239 true positives, 421 true negatives, 40 false negatives, and 52 false positives.

The final recall-weighted F-score (or, F-beta score) was .961.

The total accuracy was 94.7 percent.

Recall is the number of true positives divided by the total number of elements that actually belong to the positive class -i.e., true positives plus false negatives.

- Recall equaled .960.
- Precision equaled .969.
- The F1 Score equaled .964.

In product terms, this means that we could expect the model to correctly pick if an individual has pneumonia based on their x-ray 91.4 percent of the time. Further, it has a much higher false positive rate than false negative, as it was designed to.

If I could further this project, I would attempt further data augmentation. The data augmentation I attempted did not improve the performance of the model, although that portion of the model was cut out for brevity, along with numerous other versions of the CNNs.

Thank you.