# **FDA Submission**

Your Name: Le Ngoc Anh

Name of your Device: X-Ray Pneumonia Analyzer

# Algorithm Description

### 1. General Information

**Intended Use Statement:** Support radiologist to detect the presence or absence of pneumonia from Chest X-ray image

**Indications for Use:** This algorithm is intended for use both man and woman from the ages of 1-95 who have taken X-Rays image of Chest on PA (Posteroanterior) or AP (Anteroposterior) position.

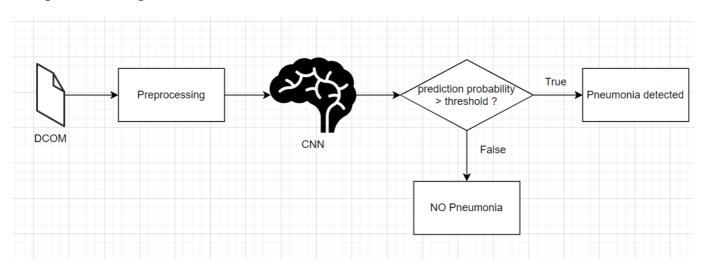
#### **Device Limitations:**

- This device is intended only for the analysis of chest X-ray images PA (Posteroanterior) or AP (Anteroposterior) position
- Image resolution is 224x224 pixels
- The model is designed specifically to detect pneumonia only and may not reliably detect other diseases on the chest X-Rays image

### **Clinical Impact of Performance:**

• This model support radiologist to reduce the diagnostic time to detect pneumonia on chest x-rays image

# 2. Algorithm Design and Function



## **DICOM Checking Steps:**

- The file extention is .dcm
- Can read the file using pydicom.dcmread
- the image is at pixel\_array

### **Preprocessing Steps:**

• Use ImageDataGenerator for Preprocessing

#### **CNN Architecture:**

• refer to "my\_model.json"

# 3. Algorithm Training

#### **Parameters:**

 Types of augmentation used during training: ImageDataGenerator with horizontal\_flip = True, vertical\_flip = False, height\_shift\_range= 0.1, width\_shift\_range=0.1,

rotation\_range=20,

shear\_range = 0.1,
zoom\_range=0.1

- Batch size: 16
- Optimizer learning rate: Adam(lr=1e-4)
- Layers of pre-existing architecture that were frozen: vgg\_model.layers[0:15]
- Layers of pre-existing architecture that were fine-tuned: vgg\_model.layers[15:]
- Layers added to pre-existing architecture new\_model.add(Flatten())

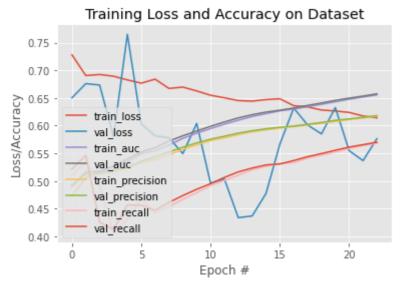
new\_model.add(Dropout(0.5))
new\_model.add(Dense(1024, activation='relu'))
new\_model.add(Dropout(0.5))

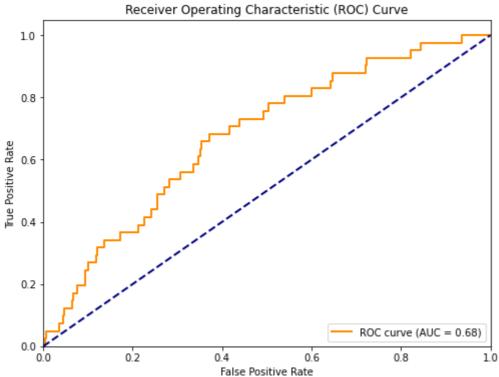
new\_model.add(Dense(512, activation='relu'))

 $new\_model.add(Dropout(0.5))$ 

new\_model.add(Dense(256, activation='relu'))

 $new\_model.add(Dense(1,\,activation='sigmoid'))$ 





Final Threshold and Explanation: The Threshold is caculated to have the best F1 score

# 4. Databases

(For the below, include visualizations as they are useful and relevant)

# **Description of Training Dataset:**

- Trainging dataset has 2290 records
- 50% records in training dataset is labeled pneumonia

# **Description of Validation Dataset:**

- Validation dataset has 22424 records
- 286 record is labeled pneumonia

### 5. Ground Truth

The authors used Natural Language Processing to text-mine disease classifications from the associated radiological reports. The labels are expected to be >90% accurate

### 6. FDA Validation Plan

# **Patient Population Description for FDA Validation Dataset:**

- Patient age is 1-95 years
- Balance distribution between male and female
- Balance PA and AP view position

**Ground Truth Acquisition Methodology:** The ground truth for pneumonia use Natural Language Processing to text-mine disease classifications from the associated radiological reports. The labels are expected to be >90% accurate

**Algorithm Performance Standard:** The best threshold is 0.7964839 has precision 5% and recall 7%