

# FDA Submission

**Your Name::** Kofi Ofuafor

**Name of your Device::** Pneumonia detection algorithm

## Algorithm Description

### 1. General Information

**Intended Use Statement:** This algorithm is intended for use in *assisting the clinician* in acute care setting with a speedy detection of pneumonia within a chest x-ray. The predicate device for this algorithm is a CADx device.

**Indications for Use:** This algorithm is used as a Pneumonia classifier in a chest x-ray taken in acute medical or surgical admission units, accident and emergency and acute care wards. It is intended for use in people aged 20-70 years with *no prior history* of presence of edema, pleural effusion and pulmonary infiltrations

**Device Limitations:** The device study was based on images from people of age 20- 70 and was significantly less accurate in the presence of concurrent disease and therefore not recommended for use in the category of patients where there is prior presence of disease in a column num\_prior\_positive in the demographic data. The algorithm was not trained in patients over 80 years old.

**Clinical Impact of Performance:** \* Performance statistics \* Accuracy: 81% \* Precision: 0.16666666666666666 \* Sensitivity: 1 \* Specificity: 0

For this algorithm, missing a detection of Pneumonia is not acceptable as human life depends on it. False negative may slow down clinicians' urgency to look at the patient's x-ray for a physical examination. Therefore False Negative results have more impact than a False positive. Accordingly, the threshold for the algorithm has been weighed in favour of recall (sensitivity).

When a test with high recall returns a negative result, you can be pretty confident that the result is truly negative. Recall does not take into account FP though, so you may still be labelling a lot of negative cases as positive. So recall is good for screening tests.

### 2. Algorithm Design and Function

**Algorithm Flow chart** The check\_dicom function reads in a .dcm file, checks the important fields for our device, and returns a numpy array of just the imaging data.

The function, check\_dicom(), is required to check the image type (MODALITY), Body Part Examined (Chest), and Image position (PA or AP) for each DICOM image and check if the input to the algorithm is valid to be predicted by our algorithm or not.

**DICOM Checking Steps:** There are three steps: 1. Extract the desired attributes \*Modality: DX The image type DX is a digital radiography \*Body Part Examined: Must be Chest \*Patient Position: AP or PA The image position could be AP or PA \*Study Description: label This is the label for the chest x-ray.

2. Next the check\_dicom function checks each image if the input to the algorithm is valid to be predicted upon by our device.
3. Finally, valid images are included as input and a numpy array of the image data is returned.

**Preprocessing Steps:** The preprocess\_image function used here takes in the image value returned from check\_dicom, and the image mean, standard deviation as well as image size corresponding with that of VGG16 input image size of (1,224,224, 3). These inputs are used to standardize and resize the image which is returned as a processed image.

Image augmentation techniques were also applied prior to training the model especially to training data.

**CNN Architecture:** The pretrained model was loaded and the first 17 layers were rendered untrainable. Here is the pretrained model

Loaded Model

Seven fully connected layers were added to the pretrained model as classifiers. Of the 7 layers, three were Dropout layers to prevent overfitting.

My model

### 3. Algorithm Training

**Parameters:** \* Types of augmentation used during training \* 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 \* Training: 16 \* Validation: 32 \* Performance evaluation: 100 \* Optimizer learning rate \* Adam \* Learning rate: 5e-6. For 5 trainable layers. Decaying rate per epoch: 0.5

**Layers of pre-existing architecture that were frozen** 17 Layers of pretrained model were frozen and therefore not trainable

Frozen Architecture

**Layers of pre-existing architecture that were fine-tuned** \* Fully connected layers were fine-tuned.

**Layers added to pre-existing architecture**

*One one convoluted was layers were fine tune and four layers were added for training*

Fine-Tuned Layers

**Algorithm training performance visualization**

*Train and validation loss/accuracy versus epochs.*

Training performance

*The plot of how the model distinguishes between the classes.*

Roc curve

*This is plot of two important parameters for the calculation of F1 Score.*

Precision recall curve

*This is a plot of F1 Scores v Thresholds.*

F1 Scores plot

**Final Threshold and Explanation:** \* Best\_threshold = 40 The decision model for this threshold was based the calculation two thresholds, from F1 scored weighted by a precision value of 0.8 and recall value of 0.8.

The threshold value from the for recall of 0.8 was chosed because, it is important for the algorithm to pick all postive cases of pneumonia. Missing cases of pneumonia is not accepetable. Recall is more valuable under the circumstance where a negative test means you have a high chance of not having the disease.

## 4. Databases

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

### Description of Training Dataset:

*Data set was split 80:20 percent in favour of training data* By using the len of positive cases of pneumonia to randomly select equal number of negative cases.

Training data after split with 4:1

*Sizes of train and validation data after discarding excess data*

Training data after discarding excess negative data

### Description of Validation Dataset:

*Validation after split 80:20 with 20% of the entire data in the validation dataset* By finding using random sample to add a negative data to the in the ration of 1:4 postive to negative cases

Validation data after split

*Validation set must contain 20:80 ratio (positive to negative)*

Validation data after split

## 5. Ground Truth

The dataset were labeled by qualified radiologist. In AI for medical imaging, using a radiologist's labels as our "true" label is often the standard of practice, and the algorithm performance is judged based on performance against an expert human.

## 6. FDA Validation Plan

### Patient Population Description for FDA Validation Dataset:

Gender distribution

gender

Ags Distribution

Age

Concurrent disease

comorbidity

**Ground Truth Acquisiion methodology** The label was obtained by 3 radiologist by vote. The approach of using radiologist was optimal for my project becuase the data is from NIH.

**Algorithm Performance Standard:** Recall weighted threshold, was used because It is crucial not to miss positive cases.