

**Capstone Project – Pneumonia Detection Challenge**

**Final Report**

**Version 3.0**

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| --- | --- | --- |
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**GitHub repository**: [**https://github.com/bathibabud/Capstone\_Project\_RSNA\_Pneumonia\_detection**](https://github.com/bathibabud/Capstone_Project_RSNA_Pneumonia_detection)

**Report:**

**Interim:**

<https://github.com/bathibabud/Capstone_Project_RSNA_Pneumonia_detection/blob/main/AIML-Jun20A-Grp3A-InterimReport.docx>

**Final:**

<https://github.com/bathibabud/Capstone_Project_RSNA_Pneumonia_detection/blob/main/AIML-Jun20A-Group3A-FinalReport.docx>

**EDA**:

<https://github.com/bathibabud/Capstone_Project_RSNA_Pneumonia_detection/blob/main/capstone_project_object_detection_rsna_pneumonia_eda.html>

<https://github.com/bathibabud/Capstone_Project_RSNA_Pneumonia_detection/blob/main/capstone_project_object_detection_rsna_pneumonia_eda.ipynb>

**UNet VGG16**:

<https://github.com/bathibabud/Capstone_Project_RSNA_Pneumonia_detection/blob/main/capstone_project_object_detection_rsna_pneumonia_unet_vgg16.html>

<https://github.com/bathibabud/Capstone_Project_RSNA_Pneumonia_detection/blob/main/capstone_project_object_detection_rsna_pneumonia_unet_vgg16.ipynb>

**MASK RCNN**:

<https://github.com/bathibabud/Capstone_Project_RSNA_Pneumonia_detection/blob/main/capstone_project_object_detection_rsna_pneumonia_maskrcnn.html>

<https://github.com/bathibabud/Capstone_Project_RSNA_Pneumonia_detection/blob/main/capstone_project_object_detection_rsna_pneumonia_maskrcnn.ipynb>

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# Summary of the problem statement, Data and findings

## Problem Statement

Pneumonia is a global health problem that does not understand social or cultural strata, causing millions of deaths each year. In several conferences and articles, it is called "the silent killer", a nickname that reflects the little social and political awareness towards this disease that without receiving the same attention as other pathologies her number of affectations year after year are forceful. In developing countries, the affectation of this lung disease makes it one of the deadliest among children under 5 years of age, causing 15% of the deaths recorded each year

In the United States, pneumonia accounts for over 500,000 visits to emergency departments and over 50,000 deaths in 2015, keeping the ailment on the list of top 10 causes of death in the country. [Official Report](https://www.cdc.gov/nchs/data/nvsr/nvsr66/nvsr66_06_tables.pdf). Pneumonia diagnosing requires a review of a chest radiograph (CXR), clinical history, vital signs, and laboratory exams. When interpreting an X-ray by a professional, his capacity and experience are key.

Chest Radiographs basically is the process of taking an image, in other words the X-ray passes through the body and reaches a detector on the other side. Tissues with sparse material, such as lungs, which are full of air, do not absorb X-rays and appear black in the image. Dense tissues such as bones absorb X-rays and appear white in the image. (1) Black = Air (2) White = Bone (3) Grey = Tissue or Fluid. Pneumonia usually appears as an area of increased lung opacity on CXR.

The objective is to build an algorithm that can detect visual signals for pneumonia in medical images. Specifically, the algorithm needs to automatically locate lung opacities on chest radiographs, but only the opacities that look like pneumonia, and discard other types of opacities like the ones caused by fluid overload (pulmonary edema), bleeding, volume loss (atelectasis or collapse), lung cancer, post-radiation or surgical changes. Outside of the lungs, fluid in the pleural space (pleural effusion) also appears as increased opacity on CXR. A pneumonia opacity is a part of the lungs that looks darker on a radiograph and has a shape that indicates that pneumonia is (or may be) present.

As the objective is to detect and draw a bounding box on each of the pneumonia opacities, where each image can have 0 or many opacities, and the training set is already classified, it will be analyzed as a supervised classification. The neural networks seem to be the best bet, specially a FCNN (Fully Convolutional Neural Network).

To improve the efficiency and reach of diagnostic services, our aim is to build a deep learning model to detect a visual signal for pneumonia in medical images assisting medical practitioners. Model needs to automatically locate lung opacities on chest radiographs, and hence help clinicians to detect and diagnose pneumonia with high accuracy & efficiency.

## Data & Findings

One of the ways to diagnose pneumonia is to analyze the Chest X-Ray. The dataset available contains these CXR images in **DICOM** format.

DICOM – Digital Imaging and Communications in Medicine is known as an international standard for medical images and everything that is related to them. DICOM images are known to have high quality, since the diagnosis requires as clear information as possible. This format is used in a variety of medical domains like radiology, cardiology and so on.

It contains several important bits of information: for example, unique patient ID, position of the body when the scan was taken, patient gender, age and so on.

The input files provided are:

* stage\_2\_train\_labels.csv - the training set, contains patientIds and bounding box/target information.
* Data Fields in stage\_2\_train.csv are shown in the below table

|  |  |
| --- | --- |
| **Data Field** | **Description** |
| patientId | Patient Id. Each patient Id corresponds to a unique image |
| x | the upper-left x coordinate of the bounding box |
| y | the upper-left y coordinate of the bounding box |
| width | the width of the bounding box |
| height | the height of the bounding box |
| Target | the binary Target, indicating whether this sample has evidence of pneumonia |

* stage\_2\_detailed\_class\_info.csv - provides detailed information about the type of each image

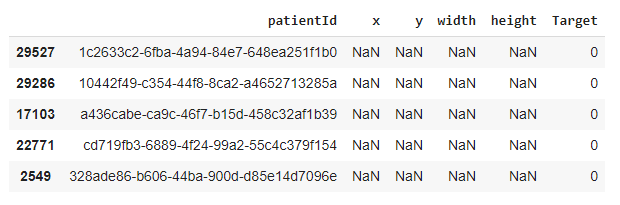
|  |  |
| --- | --- |
| **Data Field** | **Description** |
| patientId | Patient Id. Each patientId corresponds to a unique image |
| class | Contains one of the below values in each of the rows:   * No Lung Opacity / Not Normal * Normal * Lung Opacity |

* Training images are provided in stage\_2\_train\_images.zip
* Test images are provided as stage\_2\_test\_images.zip

The training data is provided as a set of patientIds and bounding boxes. Bounding boxes are defined as follows: x-min y-min width height. There is also a binary target column, Target, indicating pneumonia or normal. The objective is to identify if the patient is having pneumonia i.e., predict whether pneumonia exists in a given image. It is done by predicting bounding boxes around areas of the lung. Samples without bounding boxes are negative and contain no definitive evidence of pneumonia. Samples with bounding boxes indicate evidence of pneumonia.

When making predictions, as many bounding boxes as required are to be predicted. There should be only ONE predicted row per image. This row may include multiple bounding boxes.

**Sample Data:**



**Data Findings:**

1. stage\_2\_train\_labels.csv: The CSV file contains PatientId, bounding box details with (x, y) coordinates and width and height that encapsulates the box. It also contains the Target variable. For target variable 0, the bounding box values have NaN values.
2. There are only 26684 images in the image directory, but the csv file contains 30227 rows. There are more rows than the images, which indicates there are duplicate entries for the patientId.
3. Patient Ids are duplicated in different rows of stage\_2\_train\_labels.csv with multiple bounding boxes values. From the data it is clear that the patient is identified with pneumonia at multiple areas in the lungs.
4. We observe that of the total 30227 rows, 9555 rows have non null bounding box values. So, all bounding boxes are either defined or not defined.
5. The total number of patientIds that are identified with Pneumonia are 9555 and it matches the non-null values. It can be inferred from this that all pneumonia data set has bounding boxes defined and for normal patients, no bounding boxes exist
6. The “target” data field has two values “0” and “1” denoting “Normal” and “affected by Pneumonia” respectively.

# Summary of the Approach to EDA and Pre-processing

## Approach to EDA and Pre-processing

Major steps taken as part of EDA are:

* The DICOM data is explored
* The meta information from the DICOM files are extracted
* Various features of the DICOM images grouped by age, sex are visualized

## EDA and Pre-processing – Steps and Results

**Loading the data**

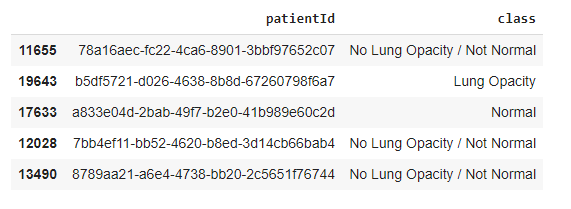
The tabular data provided in the form of csv files are loaded. There are two files:

* Detailed class info - stage\_2\_detailed\_class\_info.csv
* Train labels - stage\_2\_train\_labels.csv

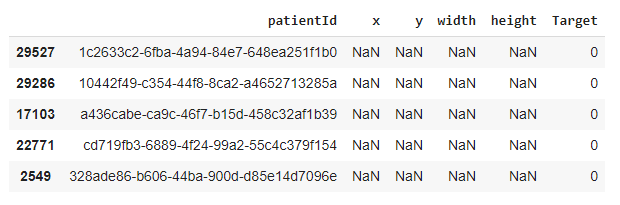
Number of rows and columns in these csv files are:

* Detailed class info - rows: 30227, columns: 2
* Train labels - rows: 30227, columns: 6

In **Detailed class info** file, the detailed information about the type of class associated with a certain patient are given. Sample data is given below:



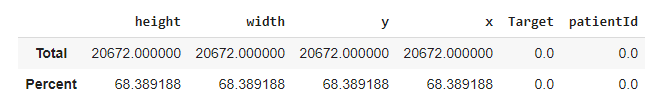
In **train labels** file, the patient ID and the window (x min, y min, width and height of the) containing evidence of pneumonia are given. Below image shows sample data.



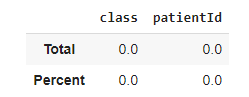
**Checking Missing values**

Missing information is checked in both the input files.

**Train labels file:** There are 20672 rows that have NaN values, of the total 30227 entries.

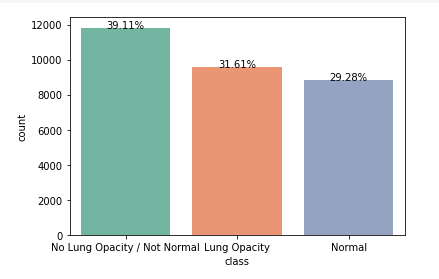
68.38% of values are missing for x, y, height and width in train labels for target 0 (not Lung opacity)

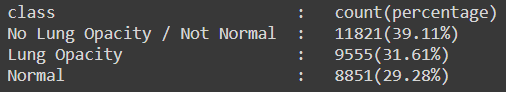
Detailed class infofile: There are no missing values



**Checking class distribution in Detailed class info file**

The class distribution of the three classes – No Lung Opacity / Not Normal, Lung Opacity and Normal is depicted below:



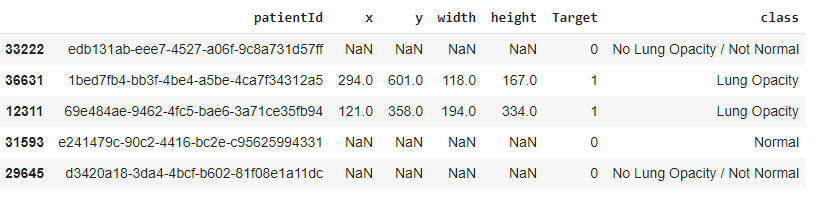


**Observations:**

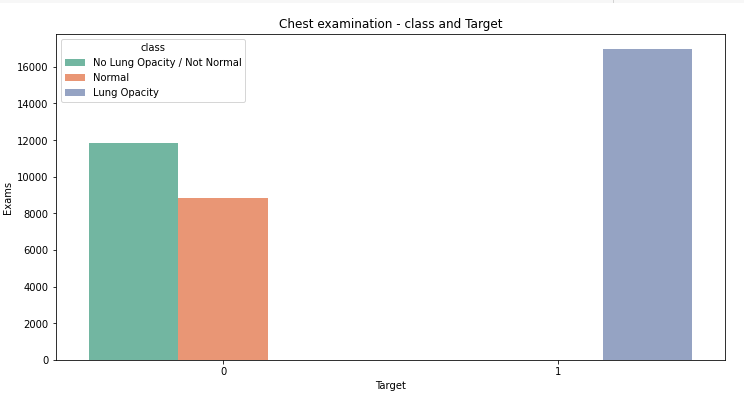
* The count of “No Lung Opacity / Not Normal” is higher than the other two classes.
* Count of “Lung Opacity” class is 31.61% of the total count 30227 which is **9555**
* No Lung Opacity / Not Normal and Normal have together the same percent (68.39%) as the percent of missing values for target window in class details information.
* In the train set, the percent of data with value for Target = 1 is therefore 30.9%.

**Merging train labels and Detailed class info datasets to get more insights**

The two datasets (train labels and Detailed class info) are merged using Patient ID as the merge criteria. The training dataset looks as below after merging.



Number of examinations for each class detected, grouped by Target value is plotted



**Inferences:**

* All chest examinations with Target = 1 (pathology detected) associated with class: Lung Opacity.
* The chest examinations with Target = 0 (no pathology detected) are either of class: Normal or class: No Lung Opacity / Not Normal.

**Exploring DICOM image files - Reading training & test files**

The input DICOM images provided in the folders - stage\_2\_train\_images, stage\_2\_test\_images - are read



**Observations**

* The files names are the patient’s ID
* Only a reduced number of images are present in the training set (26684), compared with data in train labels dataset (30227)



* Number of unique patientIds are equal to the number of DICOM images in the train set

**Extracting the inner details of single DICOM image and processing the information**

Single Image is processed for extracting below DICOM information

Dataset.file\_meta -------------------------------

(0002, 0000) File Meta Information Group Length UL: 202

(0002, 0001) File Meta Information Version OB: b'\x00\x01'

(0002, 0002) Media Storage SOP Class UID UI: Secondary Capture Image Storage

(0002, 0003) Media Storage SOP Instance UID UI: 1.2.276.0.7230010.3.1.4.8323329.28530.1517874485.775526

(0002, 0010) Transfer Syntax UID UI: JPEG Baseline (Process 1)

(0002, 0012) Implementation Class UID UI: 1.2.276.0.7230010.3.0.3.6.0

(0002, 0013) Implementation Version Name SH: 'OFFIS\_DCMTK\_360'

-------------------------------------------------

(0008, 0005) Specific Character Set CS: 'ISO\_IR 100'

(0008, 0016) SOP Class UID UI: Secondary Capture Image Storage

(0008, 0018) SOP Instance UID UI: 1.2.276.0.7230010.3.1.4.8323329.28530.1517874485.775526

(0008, 0020) Study Date DA: '19010101'

(0008, 0030) Study Time TM: '000000.00'

(0008, 0050) Accession Number SH: ''

(0008, 0060) Modality CS: 'CR'

(0008, 0064) Conversion Type CS: 'WSD'

(0008, 0090) Referring Physician's Name PN: ''

(0008, 103e) Series Description LO: 'view: PA'

(0010, 0010) Patient's Name PN: '0004cfab-14fd-4e49-80ba-63a80b6bddd6'

(0010, 0020) Patient ID LO: '0004cfab-14fd-4e49-80ba-63a80b6bddd6'

(0010, 0030) Patient's Birth Date DA: ''

(0010, 0040) Patient's Sex CS: 'F'

(0010, 1010) Patient's Age AS: '51'

(0018, 0015) Body Part Examined CS: 'CHEST'

(0018, 5101) View Position CS: 'PA'

(0020, 000d) Study Instance UID UI: 1.2.276.0.7230010.3.1.2.8323329.28530.1517874485.775525

(0020, 000e) Series Instance UID UI: 1.2.276.0.7230010.3.1.3.8323329.28530.1517874485.775524

(0020, 0010) Study ID SH: ''

(0020, 0011) Series Number IS: "1"

(0020, 0013) Instance Number IS: "1"

(0020, 0020) Patient Orientation CS: ''

(0028, 0002) Samples per Pixel US: 1

(0028, 0004) Photometric Interpretation CS: 'MONOCHROME2'

(0028, 0010) Rows US: 1024

(0028, 0011) Columns US: 1024

(0028, 0030) Pixel Spacing DS: [0.14300000000000002, 0.14300000000000002]

(0028, 0100) Bits Allocated US: 8

(0028, 0101) Bits Stored US: 8

(0028, 0102) High Bit US: 7

(0028, 0103) Pixel Representation US: 0

(0028, 2110) Lossy Image Compression CS: '01'

(0028, 2114) Lossy Image Compression Method CS: 'ISO\_10918\_1'

(7fe0, 0010) Pixel Data OB: Array of 142006 elements

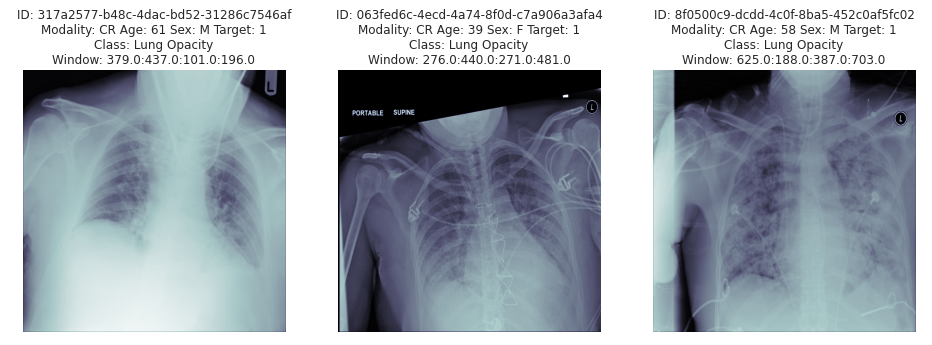
It is observed that some useful information is available in the DICOM metadata with predictive values, for example:

* Patient sex
* Patient age
* Modality
* Body part examined
* View position
* Rows & Columns
* Pixel Spacing

After extracting DICOM information, DICOM images are plotted for further study

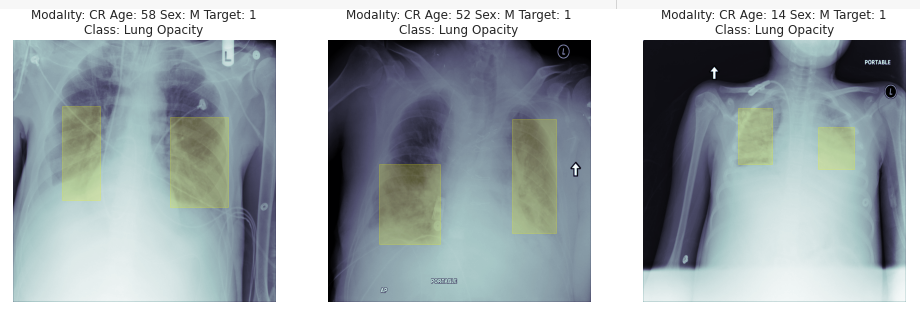
**Plotting DICOM images with Target = 1(with Pneumonia)**

Sample images plotted with Target = 1 are as below:



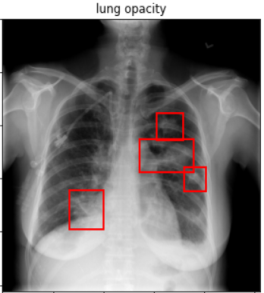
Next step is to represent the images with the overlay boxes superposed. For this, the whole dataset with Target = 1 has to be parsed and all coordinates of the windows showing a Lung Opacity on the same image have to be gathered.

Sample DICOM Images with boxes superposed showing chest areas affected by Pneumonia is as below:



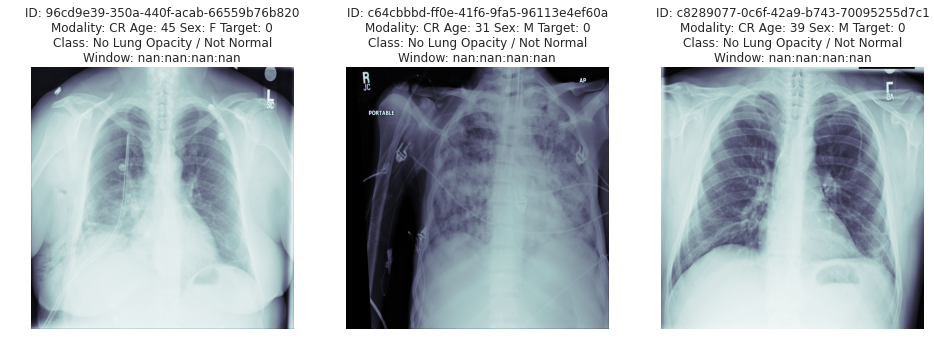
For some of the images with Target=1, we could see multiple areas (boxes) with Lung Opacity.

In the given dataset, the lung opacity is identified in lungs at a maximum of 4 areas.



**Plotting DICOM images with Target = 0 (with out Pneumonia)**

Sample images plotted with Target = 0 are as below:



**Adding metadata information from DICOM data to training and test datasets**

Once DICOM images are plotted and checked, the next step is to parse the DICOM meta information and add it to the train and test datasets

**Inferences from DICOM Metadata extracted from given images**

* Only one **modality**, “CR” - Computer Radiography is used
* As per the data given, **body part examined** is only “Chest”
* **View Position** is a radiographic view associated with the Patient Position. Both AP and PA body positions are present in the data. The meaning of these view positions is: AP - Anterior/Posterior; PA - Posterior/Anterior.

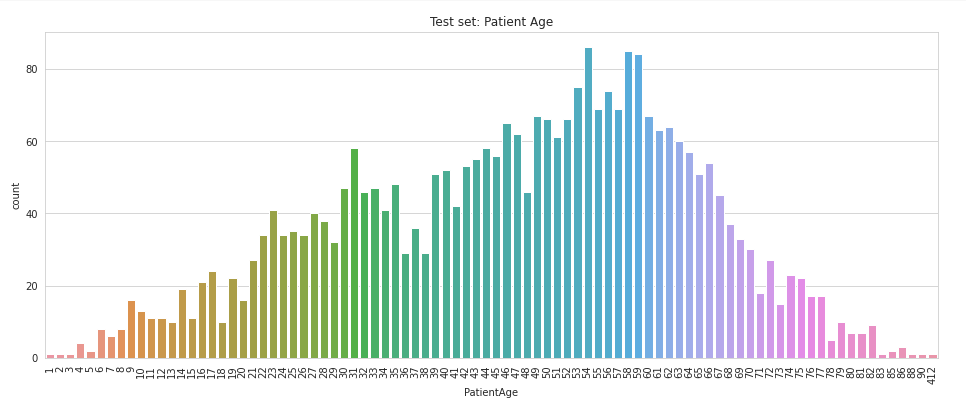
While checking the View Positions distribution in the training dataset, we got below inference:

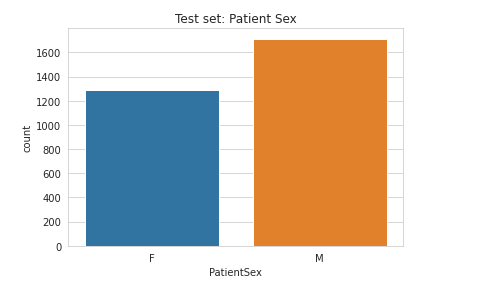


While checking the View Positions distribution in the test dataset, we got below inference:



* Both train and test have only WSD Conversion Type Data. The meaning of this Conversion Type is WSD: Workstation
* Only {Rows: Columns}, {1024:1024} are present in both train and test datasets
* Distribution of patient age for the test data set is as below:



* We can observe a few datasets has the age equal to 412. This could be a typo or a mistake in the value. As the dataset size is small compared to the total data and is not used for model training, we ignore this typo and use the data for training.
* Distribution of Patient Sex for the test data is shown as below:
* stage\_2\_sample\_submission.csv - Contains patientIds for the test set. Each row in sample submission represents one bounding box per image. This is the input .csv file for creating test dataset

## EDA - Conclusions

After exploring both the tabular and DICOM data, we draw following conclusions.

1. Discovered duplicate patientIds in the tabular data, an indication that the patient infected with Pneumonia at more places in the lungs.
2. There are patients who are infected with Pneumonia at more than one place in the lungs
3. Able to extract meta information from the DICOM data for more detailed analysis
4. Further analyze the distribution of the data with the newly added features from DICOM metadata

All these findings are useful for building a model.

# Deciding Models and Model Building

## Model Approach

The objective of our model is to detect the object classification along with identification of the object location. In simple words, the aim of [detection technique](https://analyticsindiamag.com/this-is-how-i-created-an-object-detection-model-in-less-than-5-minutes/) is to determine the classification as well as the localization of the infection. The solution requires model building based on the Classification model approach by using Computer Vision - Object Detection techniques.

The data set is huge and due to the restricted hardware resources, time taken for model training, memory consumption etc., of 26684 images data set, we have used 2000 images for training (Opacity - 1500 and normal - 500) and 500 images for validation (Opacity - 350, normal - 150).

**Bounding Box for Object Classification**

The primary goal will be the detection of bounding boxes consisting of a binary classification e.g. the presence or absence of pneumonia. However, in addition to the binary classification, the dataset without pneumonia is further categorized into normal or no lung opacity / not normal. This extra third class indicates that while pneumonia was determined not to be present, there was nonetheless some type of abnormality in the image; and oftentimes this finding may mimic the appearance of true pneumonia. This extra class is provided as supplemental information to help improve algorithm accuracy if needed; generation of this separate class will not be a formal metric used to evaluate performance.

## Model Identification

There are multiple approaches to solve an object detection problem. They are broadly classified into a 2-stage, a single stage without anchor boxes and a single stage with anchor boxes. There are numerous algorithms available for object detection with subtle variations. The popular models in this area are Mask RCNN, YOLO, SSD. As part of the project our aim is to work on 1 state-of-the-art model along with the base U-Net model.

**Base Model**: As a base model we choose to utilize the very popular UNET architecture for semantic segmentation. The UNET was developed by Olaf Ranneberger et al. for Bio Medical Image Segmentation. The architecture contains two parts. First part is the encoder, which is used to capture the context in the image. It is crucial for extracting the right features and hence we have used the most popular ImageNet competition winner, VGG16. The second part is the decoder, symmetric to the encoder, which is used to enable precise location using transposed convolutions technique. The encoder uses transfer learning while the model will be trained for decoder block.

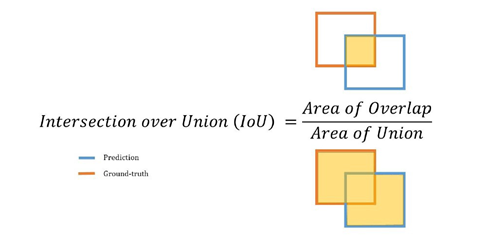
**YOLO V3**: You Only Look Once is one of the most popular and a defacto model used for object detection. Darknet-53 model is used as the backbone which consists of 53 layers. It is further added 53 more layers making YOLO a 106-layer deep convolutional neural network. YOLO uses only convolutional layers and hence it is a Fully Convolutional network. We are researching to replace the backbone of the Darknet53 with DenseNet121 as a feature extractor for YOLO and train the model with YOLO architecture creating another version of state-of-the-art model.

## Model Metrics

While there are different model metrics available to measure the model performance, two of them are widely used for object detection models. Intersection Over Union or simply IOU and Mean Average Precision, called mAP.

We have used mean IOU as our metric

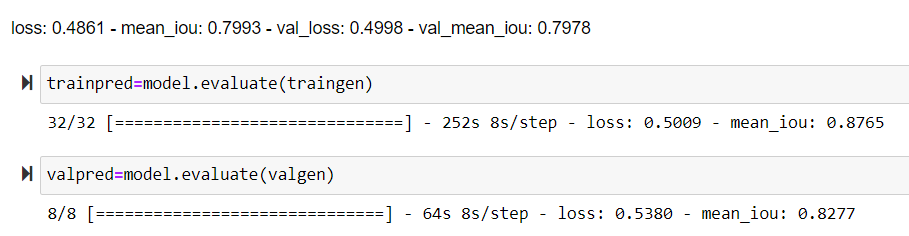
**Intersection Over Union**: IOU is a measure of the magnitude of the overlap between two bounding boxes. It calculates the size of the overlap between the two objects, divided by the total area of the two objects combined. It can be visualized as follows.



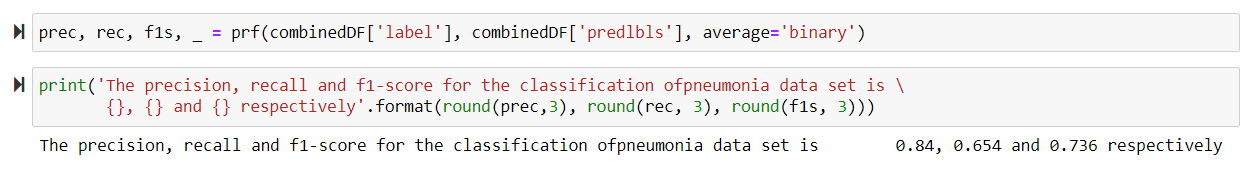
## Model Training Summary

Summary of Model performance during model training is as follows.

UNET model: With initial hyperparameters, the model had performed decently. The training and validation mean IOU during model training are 79.93% and 79.78% respectively. And during the model evaluation, the mean IOU on the training and validation data are 87.65% and 82.77% respectively. The mean IOU for both train and validation are deviated with 5%; the model is an overfit.



For classification, we calculated the precision, recall and F1 score the predictions and have 0.84, 0.654 and 0.736 respectively.



# Improving Model Performance

## Feature Extraction

In computer vision, a feature is a measurable piece of data in the image which is a unique to that specific object. It may be a specific shape such as a line, edge or a distinct color in an image or an image segment. A good feature is used to distinguish objects from one another. Hence, feature extraction is a crucial task in the success for model performance.

Although the features are extracted with different variants of pre-trained CNN models, we are trying with different approaches for better model performance and faster results. For UNET, we used the VGG16 model as our feature extractor and trained the last few layers for feature extraction. Considering the number of parameters and faster performance, we plan to use DenseNet model architecture for feature extraction.

## Data Manipulation/Augmentation

Data augmentation implies increasing the amount of training data by applying transformations to both image and contour (so we could calculate the true bounding box). Data augmentation is typically applied as a pre-processing step to the model.

There are several data augmentation techniques like random cropping (with constraints), expansion, horizontal ﬂip, image shearing, support bounding boxes, resize (with random interpolation), and color jittering (including brightness, hue, saturation, and contrast).

Data augmentation techniques can also be used to improve object detection models, although they improve single-stage detectors more than the multi-stage detectors as multi-stage detectors like Faster-RCNN use candidate object proposals that are sampled from large pool of generated ROIs, the detection results are produced by repetitive cropping of regions of feature maps. Due to this cropping, multi-stage models do not use random cropped input images, thereby they do not require detailed geometric augmentations to be applied during training.

There are several predefined image augmentation packages available for python like ImageDataGenerator, imgaug, Albumentations etc. that help us quickly perform the required augmentations.

We have considered performing image augmentation using the “Albumentations” package. This package is based on OpenCV, NumPy and imgaug.

This package efficiently implements a rich variety of image transform operations that are optimized for performance. Albumentations supports creating multiple images for computer vision tasks. We had used the Albumentations package and performed some techniques like rotation, horizontal flip, blur, brightness etc., on the images for model training.

## Model Improvements

Model improvements can be achieved in fine tuning various hyper parameters. These hyper parameters vary from model to model. There are few basic parameters that are valid for every model like Learning rate, weight decay, optimizers selection and so on.

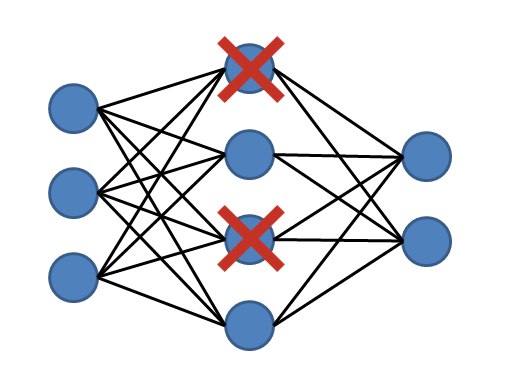
Hyperparameters are the variables which determine the network structure (Eg: Number of Hidden Units) and the variables which determine how the network is trained (Eg: Learning Rate Scheduler). Hyperparameters are set before training (before optimizing the weights and bias).

**Hyperparameters related to Network Structure**

**Number of Hidden Layers and units**

Hidden layers are the layers between input layer and output layer. Many hidden units within a layer with regularization techniques can increase accuracy. Smaller number of units may cause underfitting.

**Dropout**



Dropout is a regularization technique to avoid overfitting (increase the validation accuracy) thus increasing the generalizing power. Random neurons are cancelled

Generally, use a small dropout value of 20%-50% of neurons with 20% providing a good starting point. A probability too low has minimal effect and a value too high results in under-learning by the network.

Use a larger network. You are likely to get better performance when dropout is used on a larger network, giving the model more of an opportunity to learn independent representations.

**Network Weight Initialization**

Ideally, it may be better to use different weight initialization schemes according to the activation function used on each layer. Mostly uniform distribution is used.

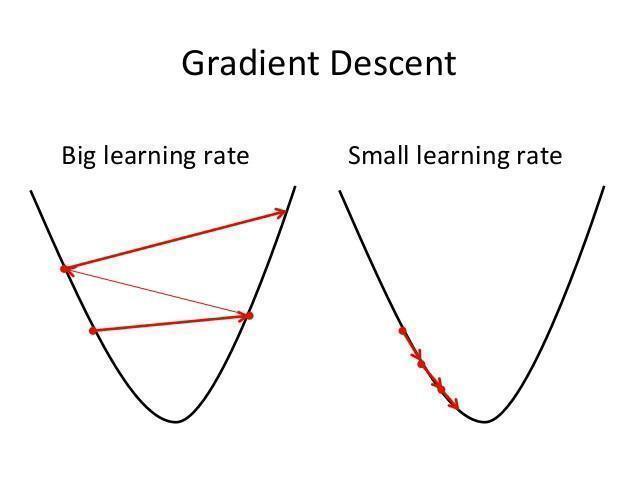
**Activation function**

Activation functions are used to introduce nonlinearity to models, which allows deep learning models to learn nonlinear prediction boundaries.

Sigmoid is used in the output layer while making binary predictions. Softmax is used in the output layer while making multi-class predictions.

**Hyperparameters related to Training Algorithm**

**Learning Rate**



The learning rate defines how quickly a network updates its parameters. Low learning rate slows down the learning process but converges smoothly. Larger learning rate speeds up the learning but may not converge. Usually a decaying Learning rate is preferred.

Sharp learning rate transition may cause the optimizer to re-stabilize the learning momentum in the following iterations.

Using a cosine scheduler (where the learning rate decreases slowly) with proper warmup (two epochs) can give better validation accuracy than using a step scheduler.

The learning rate is one of the hyperparameters that most affects performance. If we can only adjust one hyperparameter to obtain better results, then the best choice is the learning rate.

**Momentum**

Momentum helps to know the direction of the next step with the knowledge of the previous steps. It helps to prevent oscillations. A typical choice of momentum is between 0.5 to 0.9.

**Number of epochs**

Number of epochs is the number of times the whole training data is shown to the network while training. Increase the number of epochs until the validation accuracy starts decreasing even when training accuracy is increasing(overfitting).

**Batch Size Normalization**

Mini batch size is the number of sub samples given to the network after which parameter update happens. A good default for batch size might be 32. Batch size of 64, 128, 256 can also be tried

**Optimizer**

Adam is chosen as the optimizer since it’s just the best one to minimize loss value in most neural network tasks.

**Custom Loss Function**

We have used Binary Cross Entropy loss in conjunction with the Mean IOU Loss.

We will also evaluate the model by modifying the loss function parameters, using only BCE loss or using only IOU loss and compare the model performance.

**Hyper parameters used for tuning:**

Model specific parameters that are used to tune in the project are provided below.

**UNET VGG16:**

* Adding additional dropout or batch normalization layer(s)
* Play around with the last few layers for training with the custom images
* Reduced Learning Rate: Initial model used 0.01
* Use different optimizers
  + Adam optimizer which is one of the best optimizers currently
  + SGD optimizer
* Number of epochs, Batch size

**Yolo V3:**

* Custom architecture – evaluate different backbones
* Add additional dropout or batch normalization layer(s)
* Learning Rate and Optimizer hyper parameters will be modified
* Create Custom Loss functions
* Number of epochs, Batch size

**Hyperparameter Tuning evaluation**

To compare model performance, we used the followingmetrics:

* Mean IOU
* IOU loss
* Binary cross entropy loss
* Precision/Recall
* F1 Score

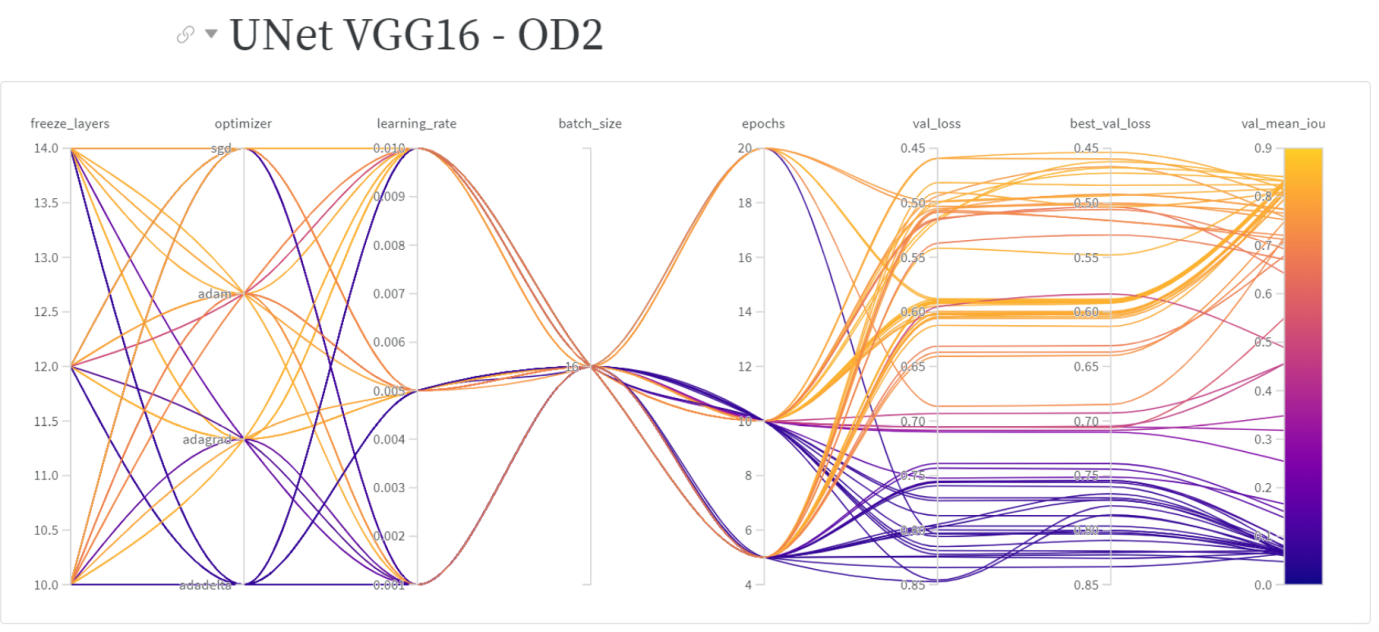
# Comparison of Models, Benchmarking and Visualization of Results of models

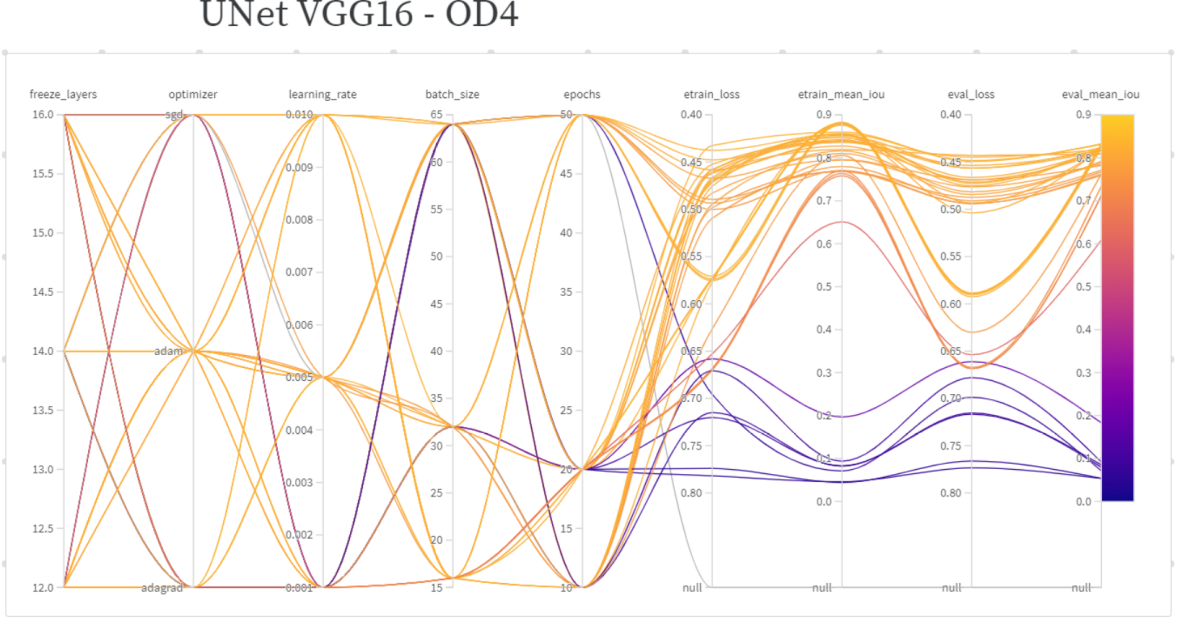
## 5.1 Approach to select final model based on outcomes

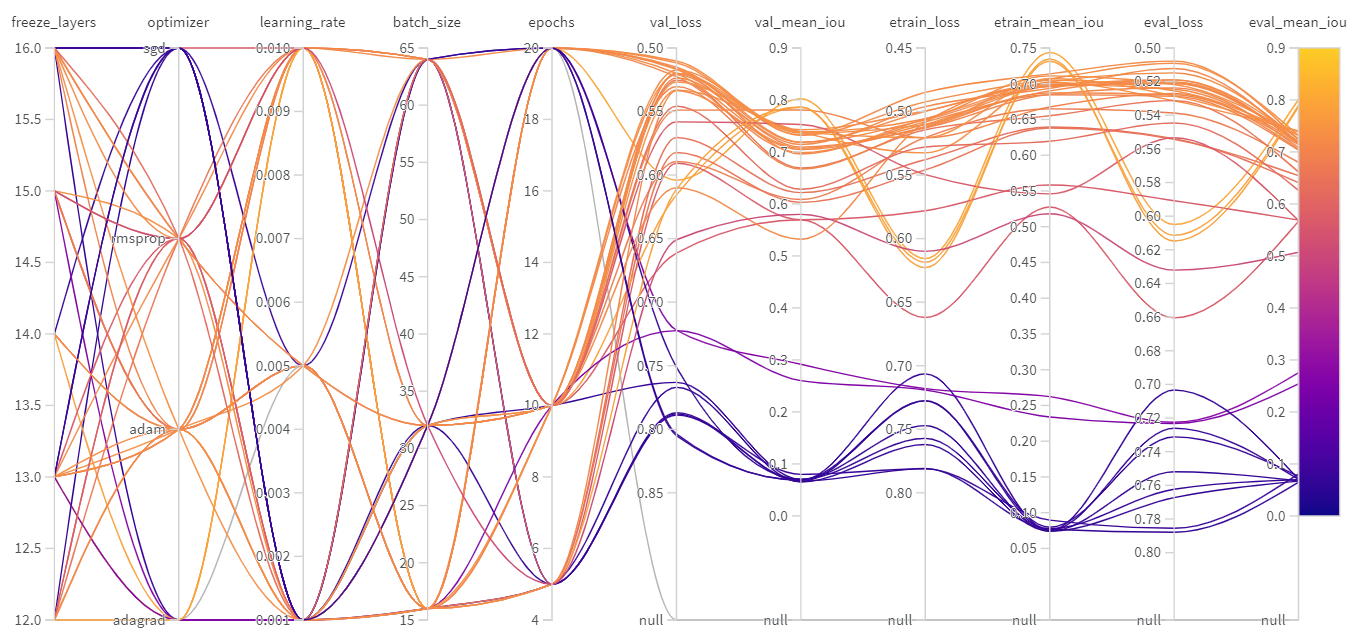
We have run hyper parameter tuning for the three models.

**UNet VGG16:**

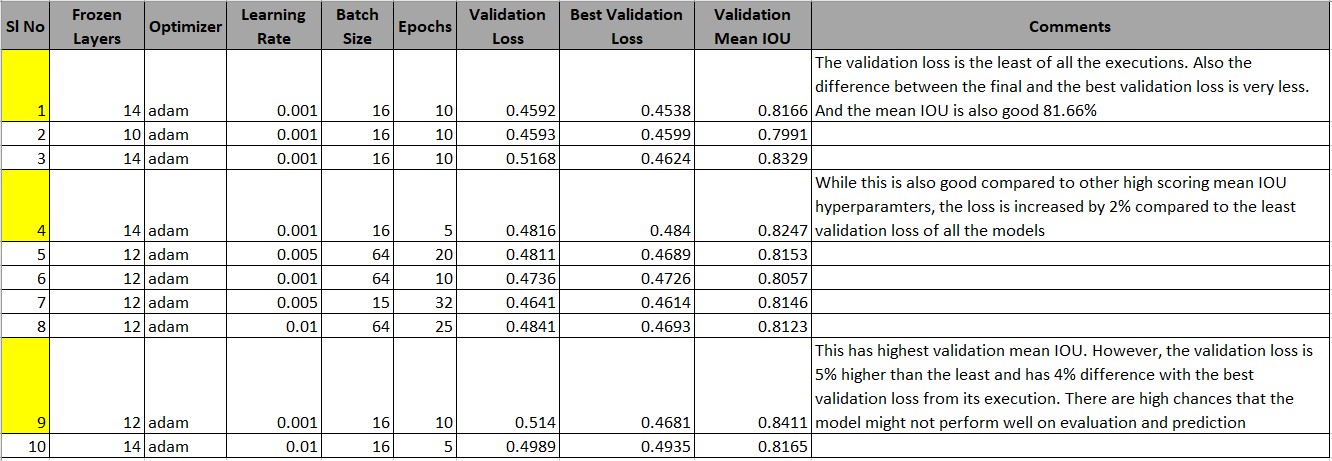
The hyper parameters considered for fine tuning the UNet VGG16 model are freeze layers of VGG16, optimizer, learning rate, batch size and epochs. We tried for multiple runs and with multiple values for each parameter. Following are some visualizations of the run. As part of training the MASK RCNN, we changed the approach in dataset selection, preprocessing techniques.

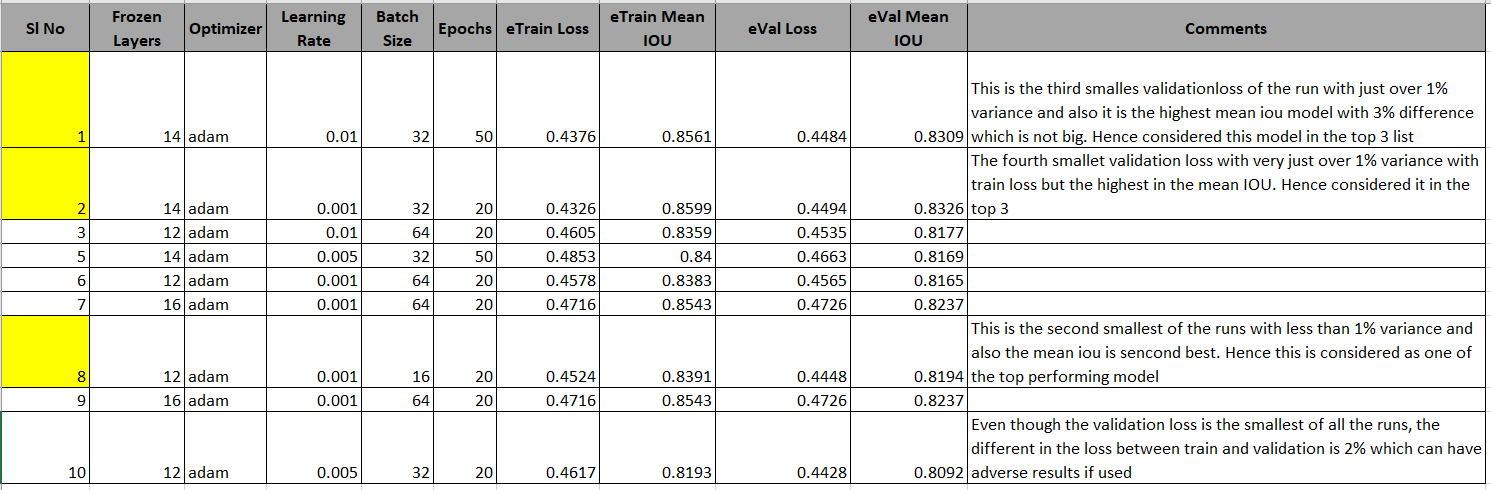


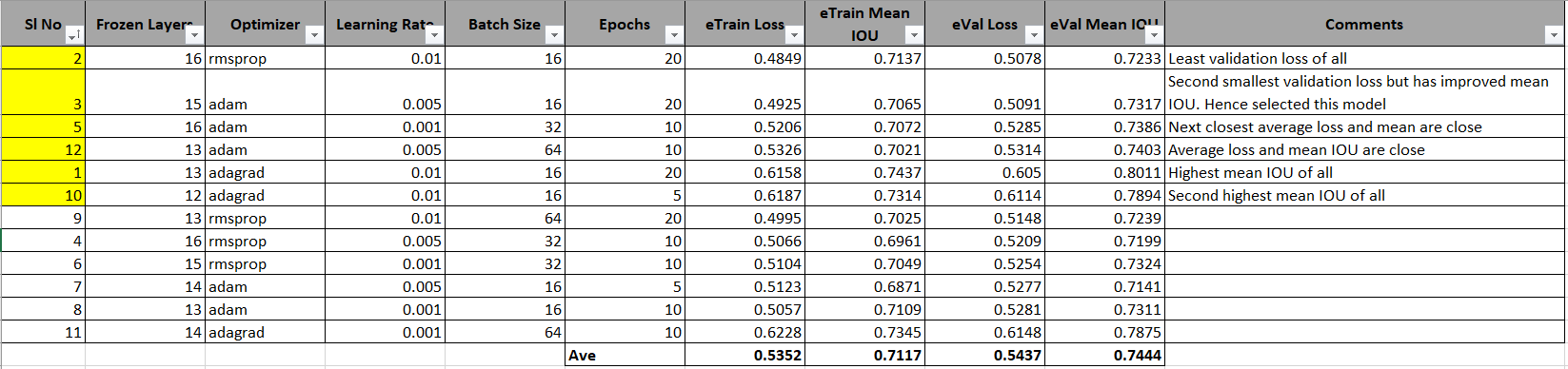




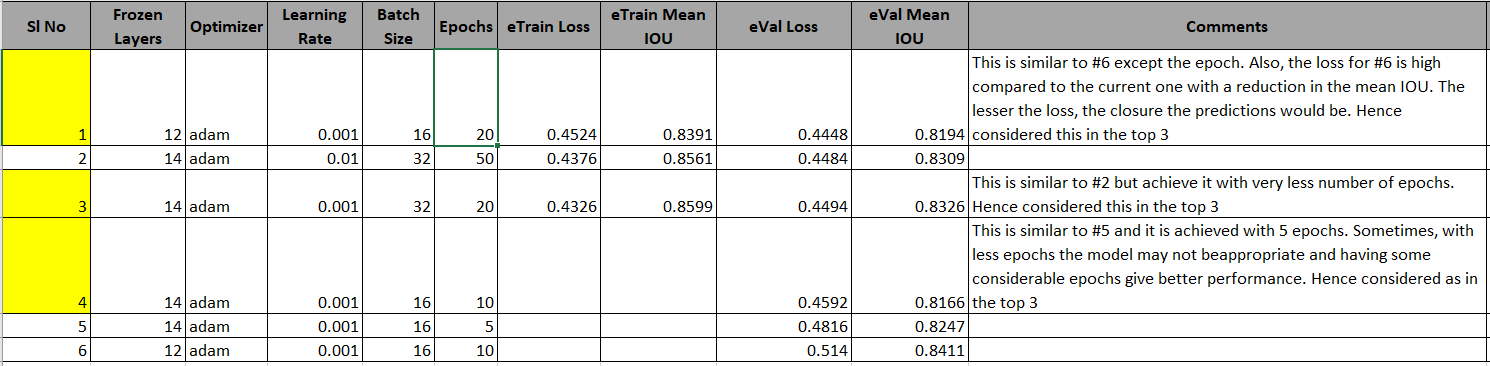
We performed 5 different runs with many different hype parameter combinations (200+), and took top 30 models that are performing well.



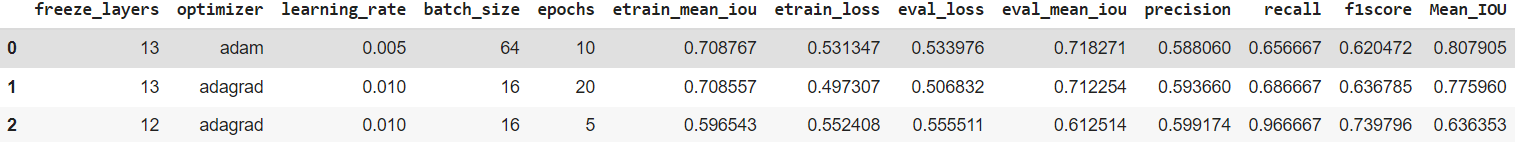




Of all 30, we reduced to 12 then to 3 best performing models.



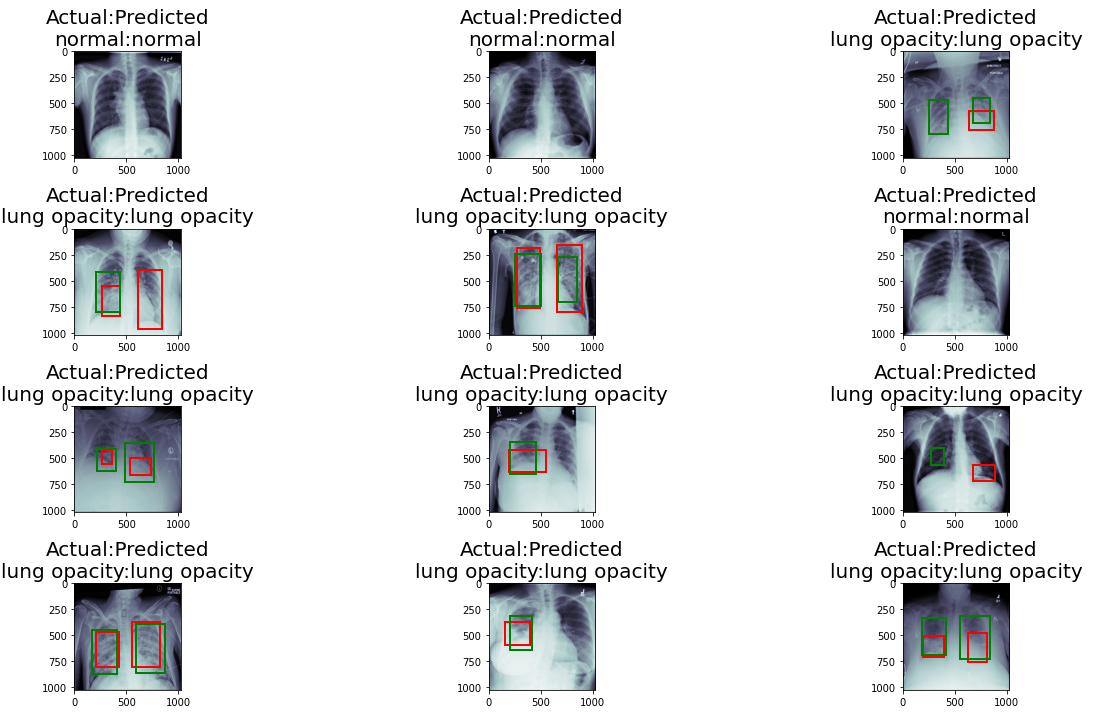
We ran the final run with the selected 3 models to identify the final best model for UNet VGG16 model. Following are the predictions for the top 3 chosen models of UNet VGG16.



From the results captured in the above table, we have the best model performance with a mean IOU of 0.808 and an F1 score of 0.621 when below hyper parameters are selected.

* Freeze layers = 13
* optimizer = Adam
* LR = 0.005
* batch size = 64
* epoch = 10

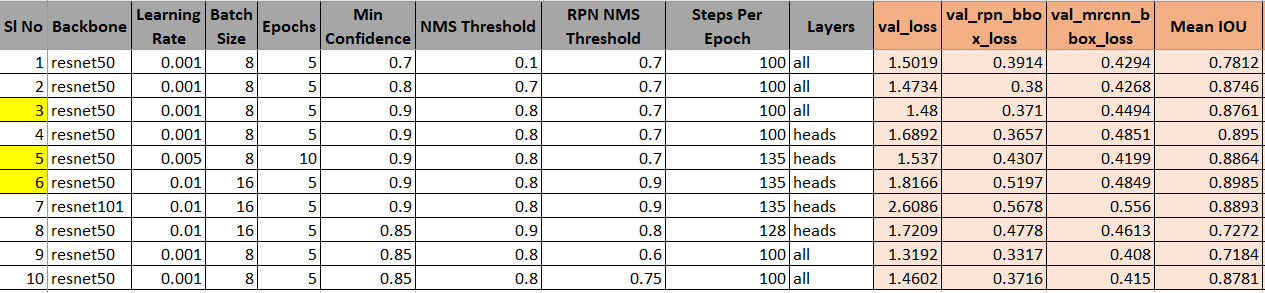
Hence, we choose the best model for UNet VGG16 with the above hyper parameters

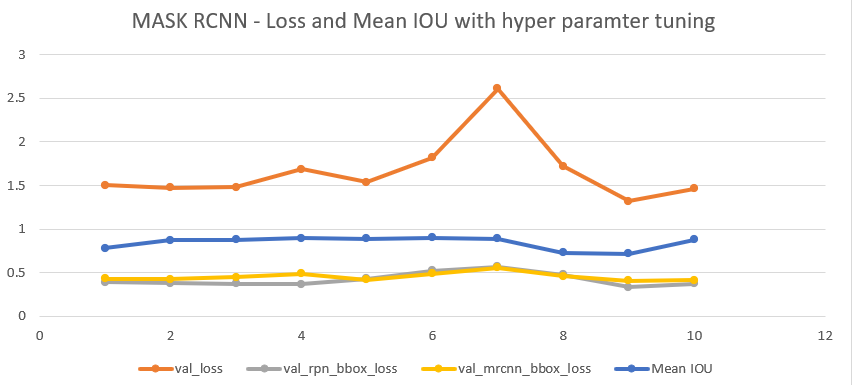


**MASK RCNN:**

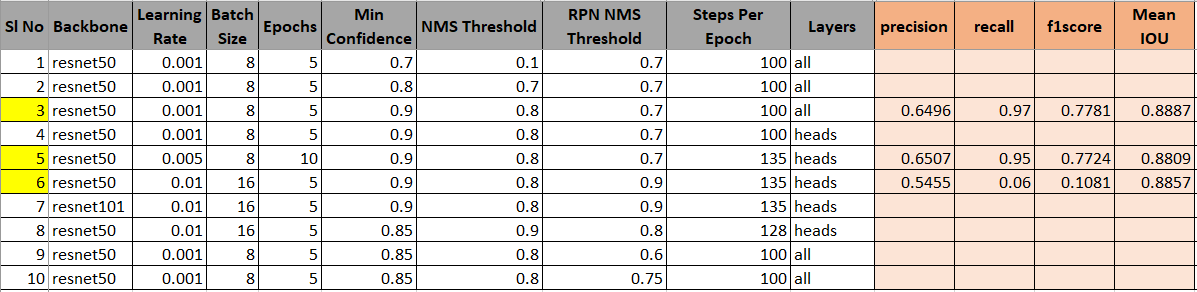
The hyper parameters considered for fine tuning the Mask RCNN model are freeze layers, learning rate, batch size, epochs, Detection Minimum Confidence, Detection NMS threshold, RPN NMS threshold and Steps Per Epoch. We tried 10 different combinations of hyper parameter tuning for the above list.

For different values for hyper parameters, the mean IOU and validation loss for all combinations are very close to one another. After close observation, with row 5 hyper parameters, the model had performed well with an overall validation loss of 1.53 and a mean IOU of 0.886.





From the top 3 models we had chosen above, we calculated Precision/Recall and F1 scores which have below results.



Based on the evaluation of the hyper parameters, the Precision, Recall and F1 score is high for model #5 which are 0.651, 0.95 and 0.772 respectively. Hence we choose this as our best model for MASK RCNN and following are the corresponding hyper parameter values.

* backbone = resnet50
* learning\_rate = 0.005
* batch\_size = 8
* epochs = 10
* det\_min\_conf = 0.9
* det\_nms\_th = 0.8
* rpn\_nms\_th = 0.7
* steps\_per\_epoch = 135
* layers = heads

**Yolo V3 – Output:**

Initial model UNET VGG16 model was created using:

* + dropout
  + batch normalization layer(s)
  + Learning Rate: 0.01
  + Optimizer = Adam optimizer
  + Number of epochs
  + Batch size

Modifications done:

* + dropout
  + batch normalization layer(s)
  + Learning Rate: 0.01
  + Optimizer = Adam optimizer
  + Number of epochs
  + Batch size

**Model Evaluation:**

On evaluating the models, we observe that Mean IOU for MASK RCNN is **0.88** which is higher than UNet VGG16 whose mean IOU is 0.80.

The precision, recall and f1 score for UNet VGG16 are 0.59, 0.66 and 0.62 respectively and compare it with MASK RCNN, Mask RCNN has high recall of **0.95** and a precision of **0.65** with an f1 score is **0.77**. The recall is high which means the model is able to predict most of the positive samples (both opacity and normal samples) correctly, however, the precision is low which indicates the true positive samples of all the positive samples is low.

**Best Model**: From the above comparison we can clearly say **MASK RCNN** is the BEST model.

# Limitations of our solution and Suggested Improvements

## Limitations of our solution

* The input data that was provided is challenging one to predict as all the images are similar, unable to differentiate between opacity and normal.
* The data set is imbalanced with respect to the classes, we had chosen to use a negative sampling approach for our model building.
* Team had a short time working on different CV models. Model executions are taking a lot of resources (GPU) and longer time, hence have considered 2000 images for training, 500 for validation and inference.
* While using the MASK RCNN model, we find it tough to look for the right github repository and the appropriate mapping of tensorflow/keras version. It didn’t work on google collab due to its nature of use of latest software versions. Hence we chose to work on kaggle, which provides the ability to select the correct versions of the software needed to build and train the model.
* As part of MASK RCNN training, the predictions are not coming correct even though the mean IOU is above 0.85. When we had a closure look, we identified that the model is predicting all images irrespective of the opacity or sometimes they are not predicting at all. Thus we had to change the data selection technique (negative sampling where we choose opacity as our main data and include negative samples with normal label data), preprocessing methods (data augmentation technique to restrict rotations, flips to horizontal) etc., for the model and adopt similar things for other models as well.

## Suggested Improvements

* Dataset selection can be improved, by trying with different combinations
* Pre-processing can be improved, by trying with different image shapes
* Model training with different hyper parameters and values

# Our Learnings

* Had an excellent opportunity to learn Computer Vision with SOTA models.
* Learned and used state of art models like UNet, MASK RCNN, Yolo v3 along with Transfer learning approach.
* Adopt different visualization techniques like line charts, bar charts, parallel coordinates, loading and displaying dicom images with bounding box representation.
* Handling imbalanced data and its implication, by adopting Negative sampling, using data augmentation technique (albumenations, imgaug) and generating more opacity images.
* Use of other external tools like Weights and Biases to train models with defined hyper parameter values, and visualize data of model parameter executions.
* It has helped to look and gain awareness on the SOTA models like MASK RCNN, YOLO V3, SSD.

# 

# Business Value Derived

* Automating Pneumonia screening in chest radiographs, providing affected area details through the bounding box.
* Assist physicians to make better clinical decisions or even replace human intervention in certain functional areas of healthcare (eg, radiology).
* Guided by relevant clinical questions, powerful AI techniques can unlock clinically relevant information hidden in the massive amount of data, which in turn can assist clinical decision making.

# Final Note

Many thanks to our Mentor – Imran for his expert guidance for completing this Capstone Project

Thanks to our Program Manager – Divya for her dedication and support in conducting AIML program smoothly as scheduled

Finally, Thanks to Great Learning for offering this wonderful course - AIML

# Citations:

**UNet**:

**Mask RCNN**: Thanks to [Thanh Bui and Matter Plot](https://www.kaggle.com/tbui001) Inc for providing the repositories that helped to execute the project smoothly

Matter Plot: <https://github.com/matterport/Mask_RCNN>

[Thanh Bui](https://www.kaggle.com/tbui001): <https://www.kaggle.com/tbui001/lung-opacity-classification-maskrcnn>

@misc{matterport\_maskrcnn\_2017,

title={Mask R-CNN for object detection and instance segmentation on Keras and TensorFlow},

author={Waleed Abdulla},

year={2017},

publisher={Github},

journal={GitHub repository},

howpublished={\url{https://github.com/matterport/Mask\_RCNN}},

}