

# Final project

## COVID segmentation based on Deep Learning

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Demo

# Demo

# 2.1

## Models implemented: Introduction

- **20 CT scans with ground-truth data to train**
- **Multi-class segmentation: Left and right lung and Disease**
- **Starting point: Unet-Model - state-of-the-art model**
- **Two approaches:**
  - Custom approach - model designed in code and modified
  - Model from Segmentation Models library - Unet - EfficientNet B7 - ImageNet weights
- **EfficientNet B7: Best accuracy and minimum amount of params (75M)**
- Optimizer used: Adaptive Moment Estimation - Adam
- Metrics implemented
  - Histogram of dice coefficient
  - Histogram of Hausdorff distance
  - Receiving Operating Characteristic (ROC) curve
  - Area under the ROC curve

# 2.2

## Models implemented: Architecture

```
def put_layer(self, previous_layer, no_nuerons):
    ...
    This Function Adds layer in CNN model
    ...
    first_conv = Conv2D(no_nuerons, (3, 3), activation='relu', padding='same')(previous_layer)
    layer_complete = Conv2D(no_nuerons, (3, 3), activation='relu', padding='same')(first_conv)

    return layer_complete
```

```
def inception_module(self, x,
                      filters_1x1,
                      filters_3x3_reduce,
                      filters_3x3,
                      filters_5x5_reduce,
                      filters_5x5,
                      filters_pool_proj,
                      name=None):

    from keras.initializers import glorot_uniform, Constant
    kernel_init = glorot_uniform()
    bias_init = Constant(value=0.2)

    conv_1x1 = Conv2D(filters_1x1, (1, 1), padding='same', activation='relu',
                      kernel_initializer=kernel_init, bias_initializer=bias_init)(x)

    conv_3x3 = Conv2D(filters_3x3_reduce, (1, 1), padding='same', activation='relu',
                      kernel_initializer=kernel_init, bias_initializer=bias_init)(x)

    conv_3x3 = Conv2D(filters_3x3, (3, 3), padding='same', activation='relu',
                      kernel_initializer=kernel_init, bias_initializer=bias_init)(conv_3x3)

    conv_5x5 = Conv2D(filters_5x5_reduce, (1, 1), padding='same', activation='relu',
                      kernel_initializer=kernel_init, bias_initializer=bias_init)(x)

    conv_5x5 = Conv2D(filters_5x5, (5, 5), padding='same', activation='relu',
                      kernel_initializer=kernel_init, bias_initializer=bias_init)(conv_5x5)

    pool_proj = MaxPooling2D((3, 3), strides=(1, 1), padding='same')(x)

    pool_proj = Conv2D(filters_pool_proj, (1, 1), padding='same', activation='relu',
                      kernel_initializer=kernel_init, bias_initializer=bias_init)(pool_proj)

    output = concatenate([conv_1x1, conv_3x3, conv_5x5, pool_proj], axis=3, name=name)

    return output
```



# 2.2

## Models implemented: Architecture

```
def first_sketch_model(self, classes):
    from keras.losses import BinaryCrossentropy, CategoricalCrossentropy
    from keras.optimizers import Adam

    optimizer = Adam(lr = 0.0001)
    loss_func = CategoricalCrossentropy()
    activation_func = 'sigmoid'

    inputs = Input((None, None, 1))

    layer_1 = self.inception_module(inputs,
        64, 96,
        128, 16,
        32 , 32,
        name = 'inception_1')      You, a few seconds ago • Uncommitted ch
pool1 = MaxPooling2D(pool_size=(2, 2))(layer_1)
drop = Dropout(0.3)(pool1)

    layer_2 = self.put_layer(drop,64)
    pool2 = MaxPooling2D(pool_size=(2, 2))(layer_2)

    layer_3 = self.put_layer(pool2,128)
    pool3 = MaxPooling2D(pool_size=(2, 2))(layer_3)

    layer_4 = self.put_layer(pool3, 256)
    pool4 = MaxPooling2D(pool_size=(2, 2))(layer_4)

    layer_5 = self.put_layer(pool4, 512)

    up6 = concatenate([Conv2DTranspose(256,
        (2, 2),
        strides=(2, 2),
        padding='same')(layer_5), layer_4], axis=3)
    layer_6 = self.put_layer(up6, 256)

    up7 = concatenate([Conv2DTranspose(128,
        (2, 2),
        strides=(2, 2),
        padding='same')(layer_6), layer_3], axis=3)
    layer_7 = self.put_layer(up7, 128)

    up8 = concatenate([Conv2DTranspose(64,
        (2, 2),
        strides=(2, 2),
        padding='same')(layer_7), layer_2], axis=3)
    layer_8 = self.put_layer(up8, 64)

    up9 = concatenate([Conv2DTranspose(32,
        (2, 2),
        strides=(2, 2),
        padding='same')(layer_8), layer_1], axis=3)
    layer_9 = self.put_layer(up9, 32)

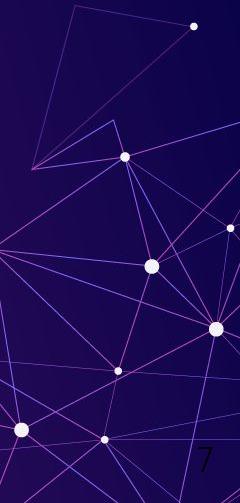
    layer_10 = Conv2D(classes, (1, 1), activation=activation_func)(layer_9)

model = Model(inputs=[inputs], outputs=[layer_10])
model.compile(optimizer = optimizer, loss=loss_func, metrics = ['accuracy'])
return model
```



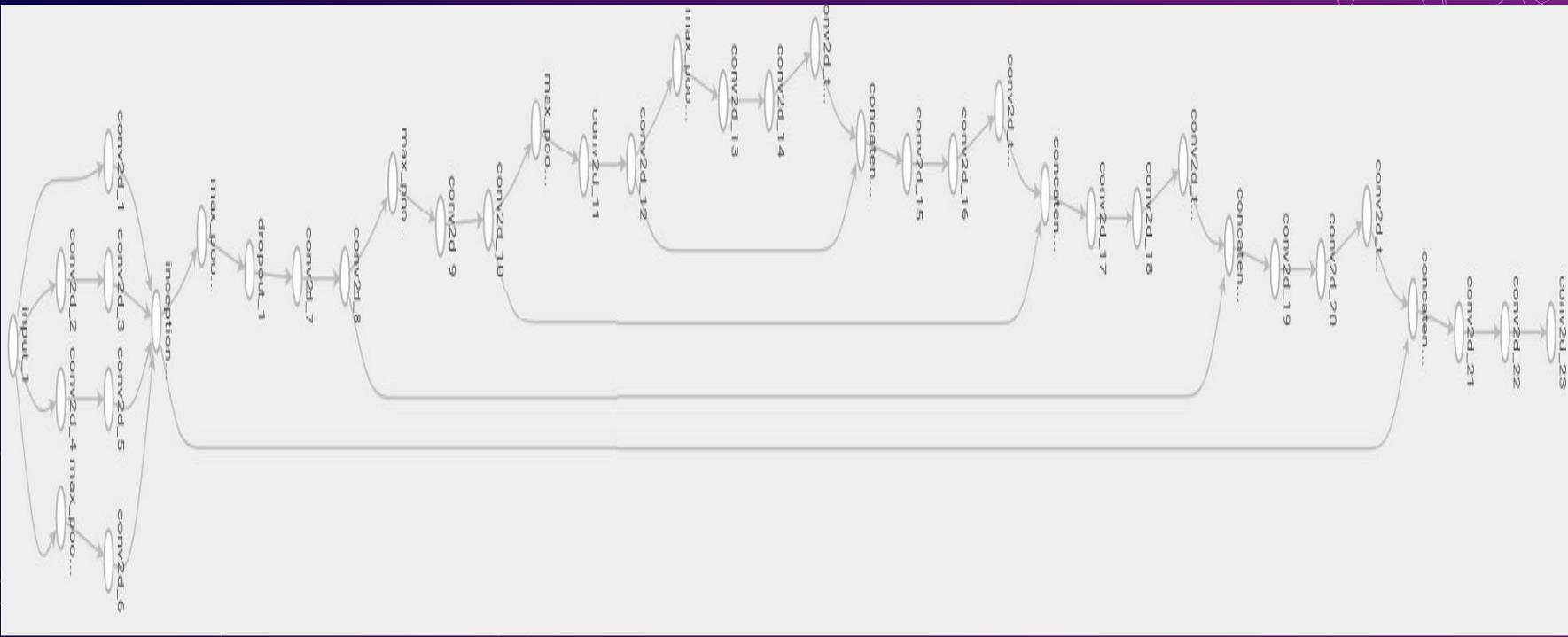
## 2.2

### Models implemented: Architecture



# 2.2

## Models implemented: Architecture



# 2.2

## Models implemented: Architecture

### Key Features of our proposed Hybrid Model

- Primary Architecture: U-net
- Inception module embedded in U-net
- Additional dropout layer to avoid overfitting
- Categorical Cross entropy
- Adam optimizer: Initial LR 0.0001
- Amount of epochs = 8
- Batch size = 8
- Activation function for last layer: Sigmoid

# 3

## Experimentation & results

- **Test realized:**

- **Convolution layers added**
- **Single Drop out layer - Change in the dropout rate : 10, 30 and 50%**
- **Multiple dropout layers (Encoder section, DO rate: 30 and 50%)**
- **Inception modules added**
  - Single module added as first layer
  - Multiple modules added in the first and second layer

- **Combination of dropout and inception module**

- **Changing epochs and batch size**

- **Use of different loss function**

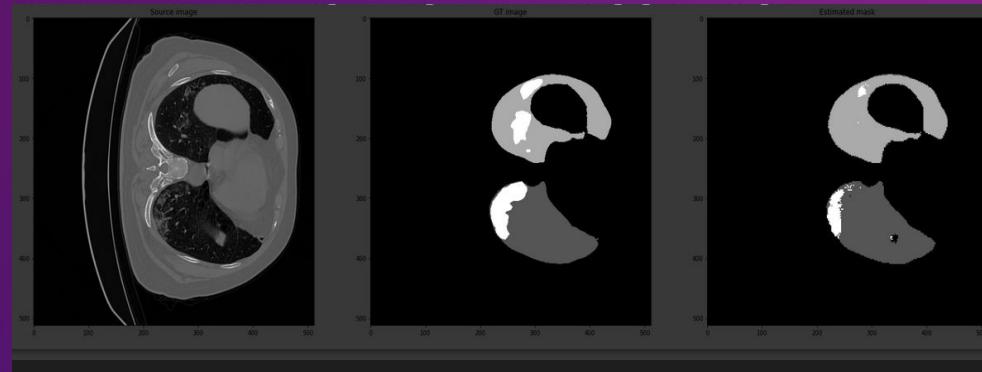
- **Binary cross entropy**
- **Categorical cross entropy**
- **Poisson**
- **MSE - MAE**
- **Kullback-Leibler Divergence**

# 3

## Experimentation & results

### Initial experimental setup

- **Results of the initial model:**
  - **Optimizer = Adam**
  - **Learning rate = 0.0001**
  - **Loss function = Binary Cross entropy**
  - **Activation function in last layer: softmax**
- - **No drop out layer used**
  - **No Data Augmentation used**



# 3

## Experimentation & results

- **Experiment 1: Adding convolutional layer**



- **Adding one convolutional layer**



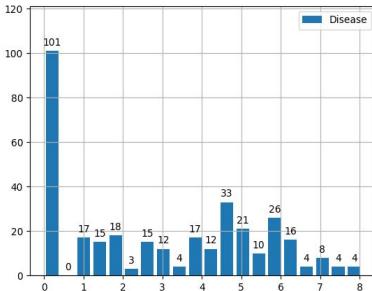
- **Adding two convolutional layer**

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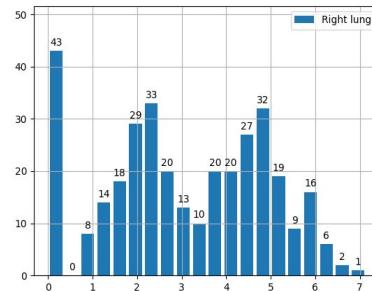
## Experimentation & results

- Experiment 1: Statistical Analysis adding one convolution layer

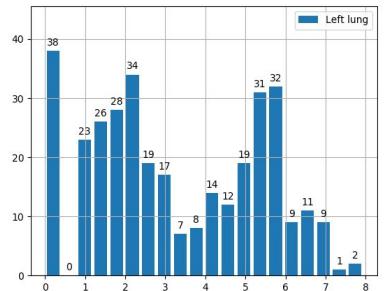
Hausdorff - Disease



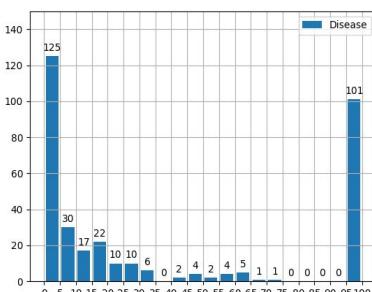
Hausdorff - RL



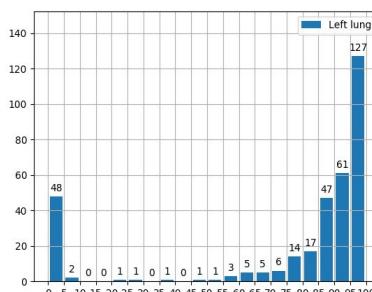
Hausdorff - LL



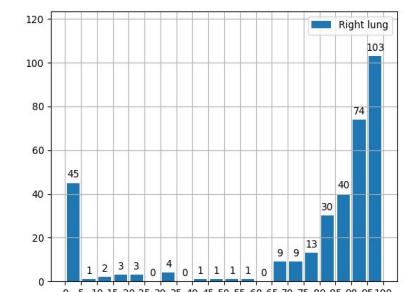
Dice coefficient - Disease



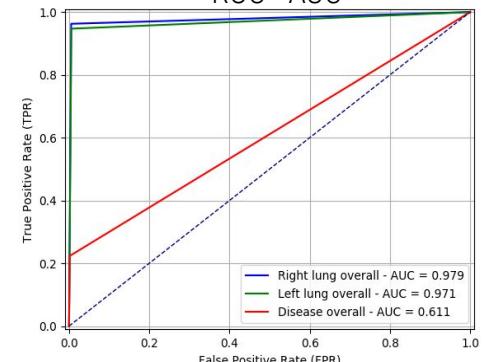
Dice coefficient - RL



Dice coefficient - LL



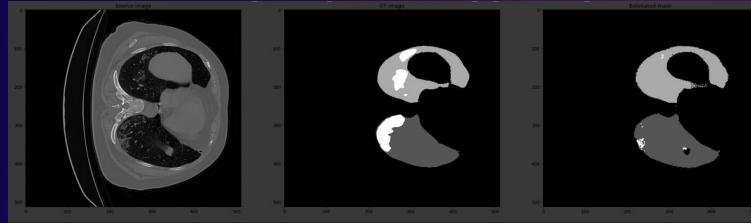
ROC - AUC



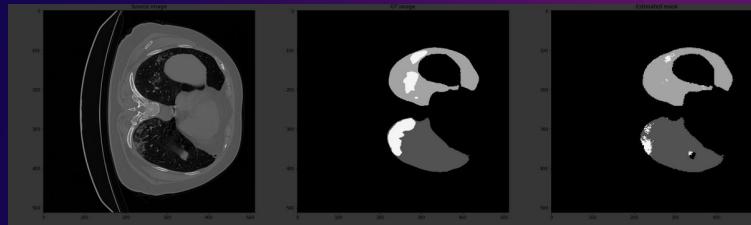
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## Experimentation & results

- Experiment 2: Adding dropout layer to initial model



- Dropout of 10% on layer 4



- Dropout of 30% on layer 4



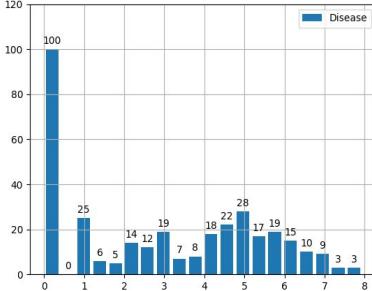
- Adding DO (30%) on first 5 layers

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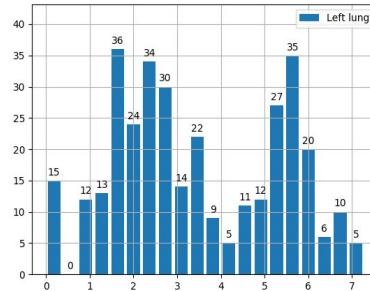
## Experimentation & results

- Experiment 2: Statistical Analysis for DO 10% on first layer

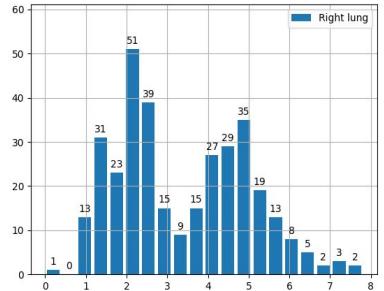
Hausdorff - Disease



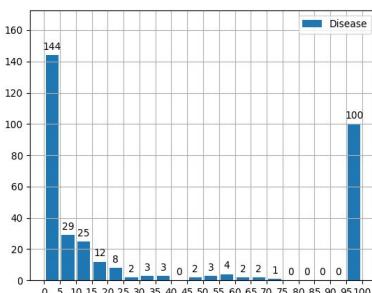
Hausdorff - RL



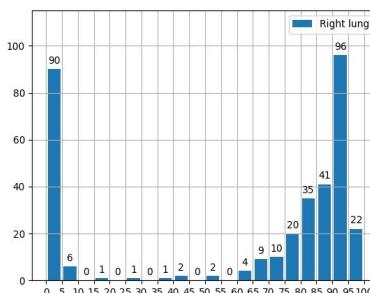
Hausdorff - LL



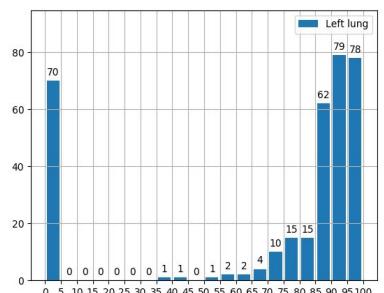
Dice coefficient - Disease



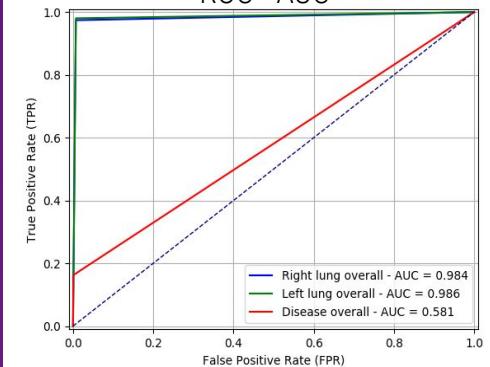
Dice coefficient - RL



Dice coefficient - LL



ROC - AUC

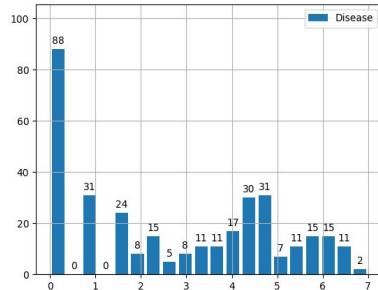


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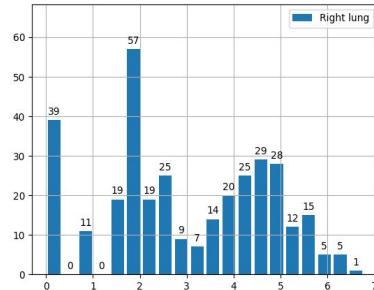
## Experimentation & results

- Experiment 2: Statistical Analysis for DO 30% on first layer

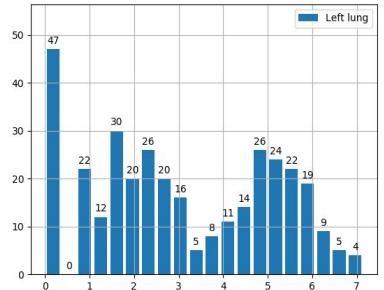
Hausdorff - Disease



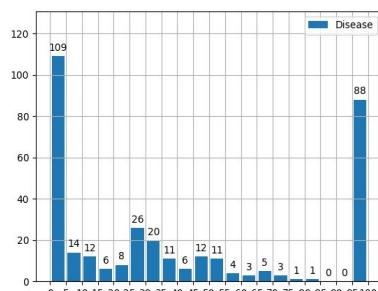
Hausdorff - RL



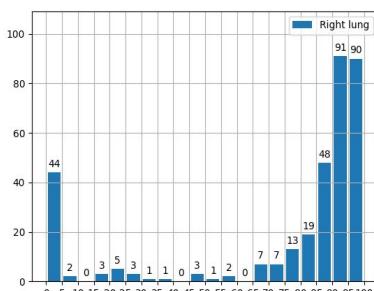
Hausdorff - LL



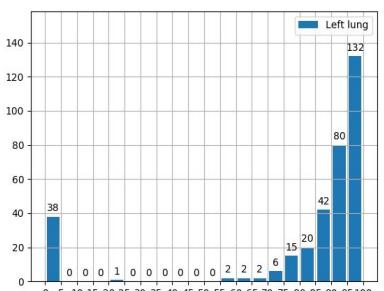
Dice coefficient - Disease



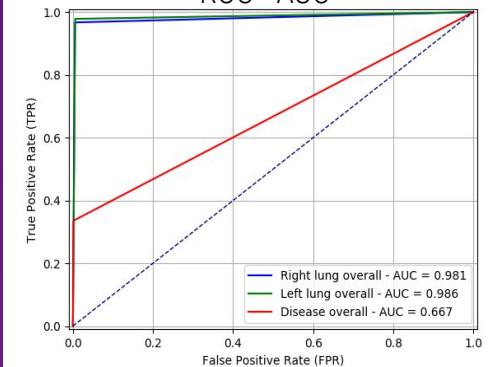
Dice coefficient - RL



Dice coefficient - LL



ROC - AUC

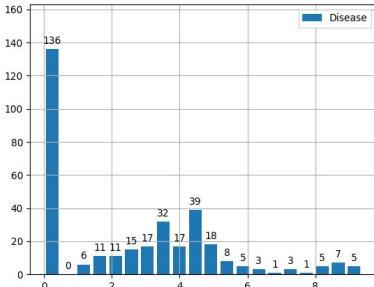


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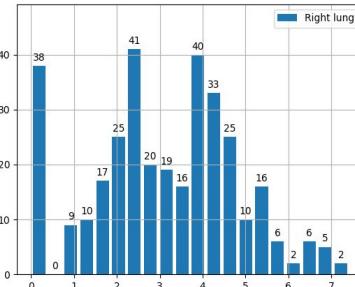
## Experimentation & results

- Experiment 2: Statistical Analysis of adding 30% DO on first 5 layers

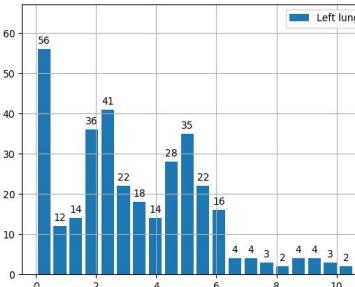
Hausdorff - Disease



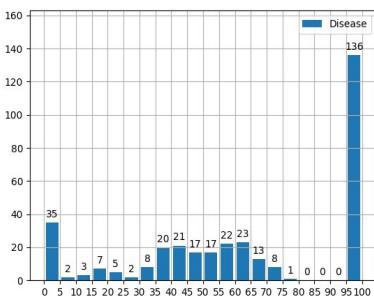
Hausdorff - RL



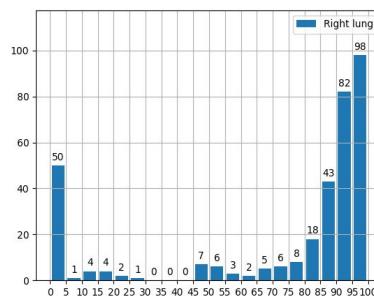
Hausdorff - LL



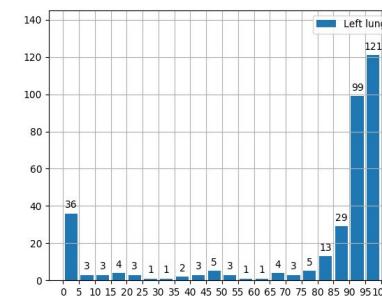
Dice coefficient - Disease



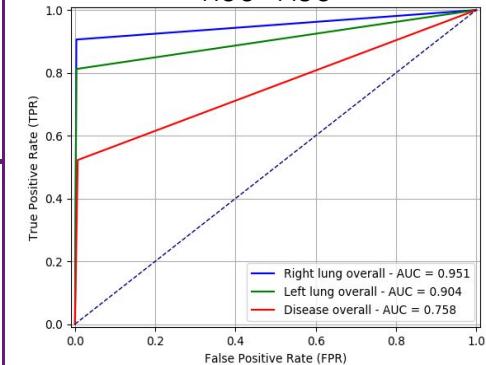
Dice coefficient - RL



Dice coefficient - LL



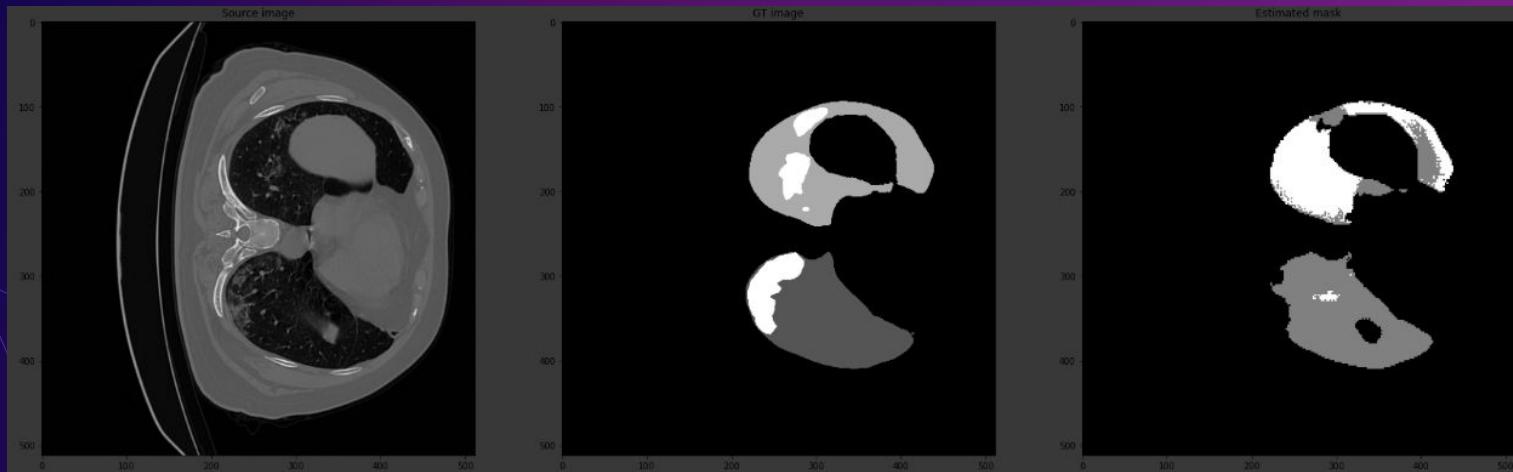
ROC - AUC



# 3

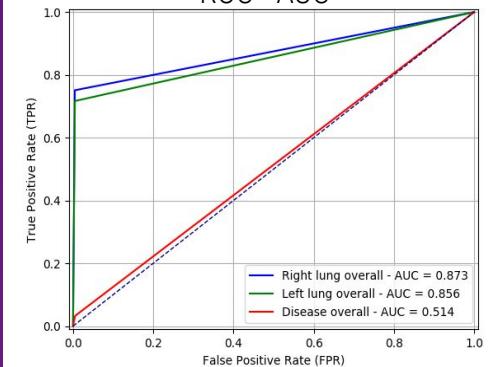
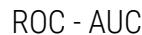
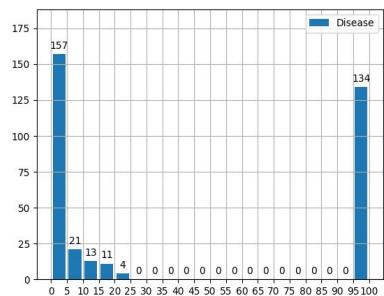
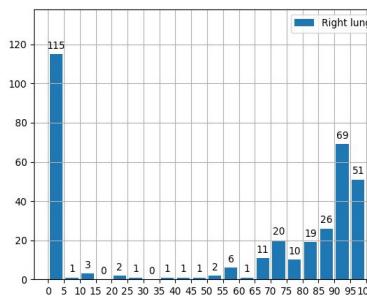
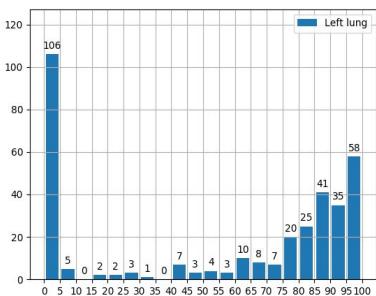
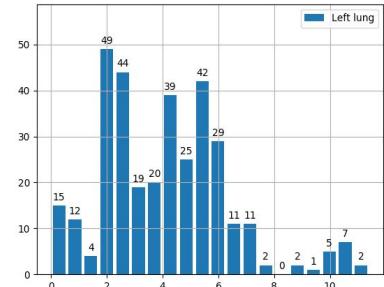
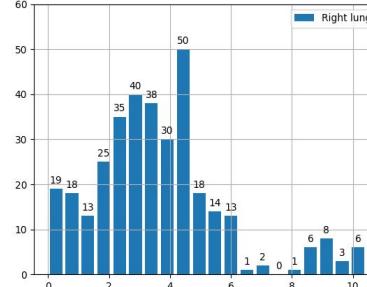
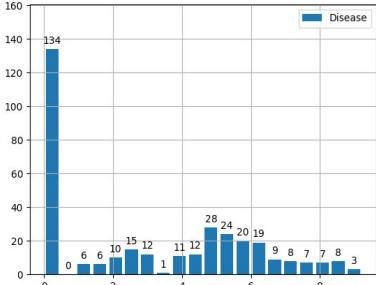
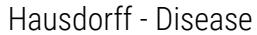
## Experimentation & results

- **Experiment 4: Adding DO (50%) on first 5 layers**



# 3 Experimentation & results

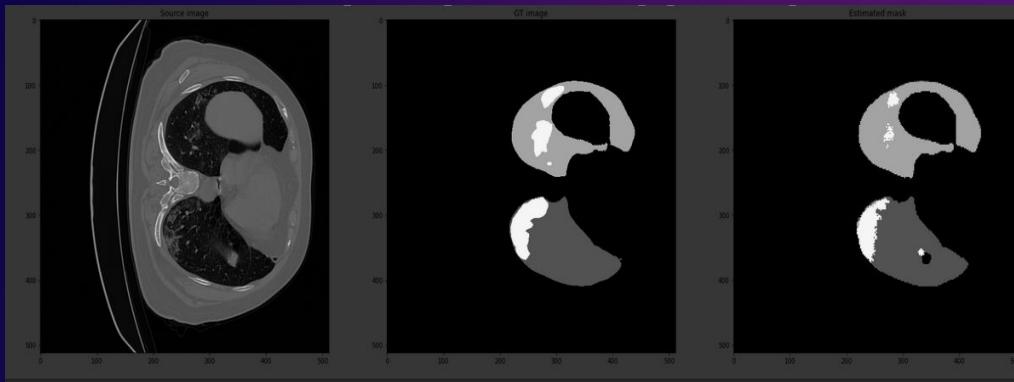
- Experiment 4: Statistical analysis Adding DO(50%) on first 5 layers



# 3

## Experimentation & results

- Experiment 3: Trying different activation functions at the last layer



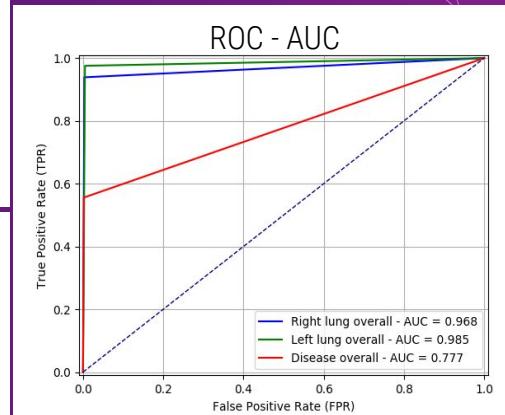
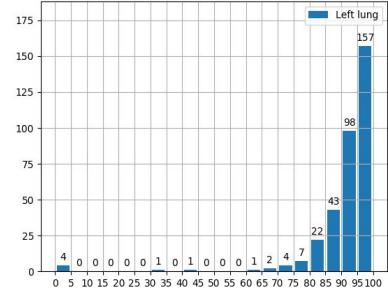
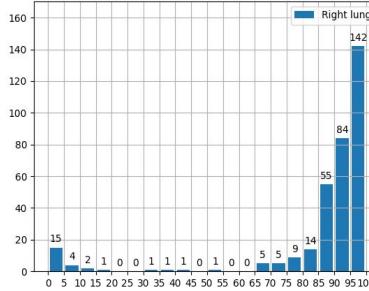
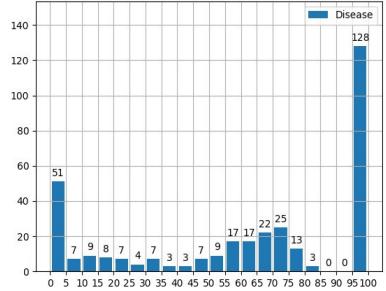
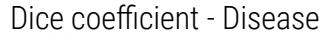
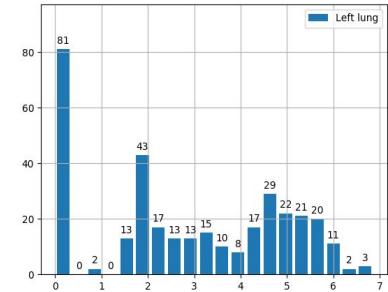
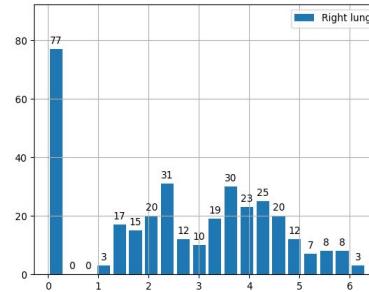
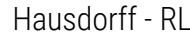
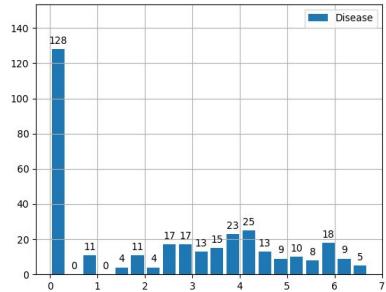
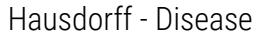
- Sigmoid



- Softmax

# 3 Experimentation & results

- **Experiment 3: Statistical Analysis of Sigmoid Activation function**

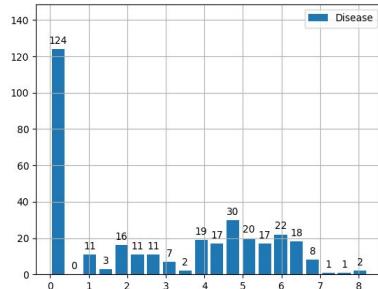


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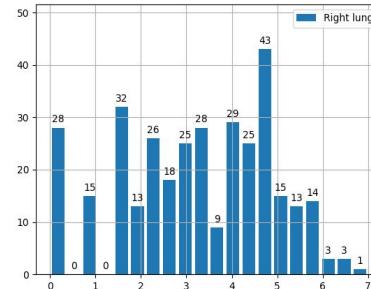
## Experimentation & results

- Experiment 3: Statistical Analysis of Softmax Activation function

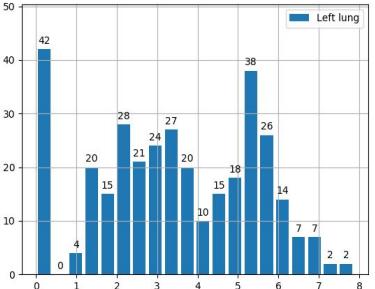
Hausdorff - Disease



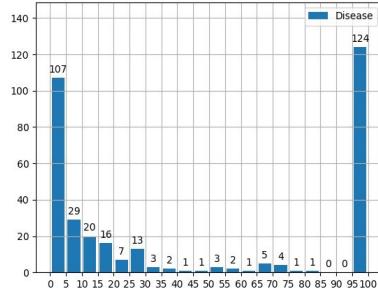
Hausdorff - RL



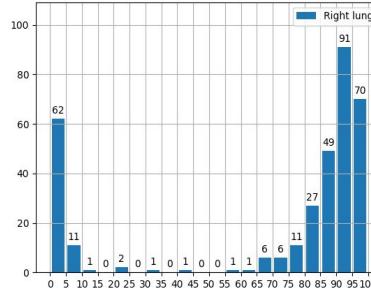
Hausdorff - LL



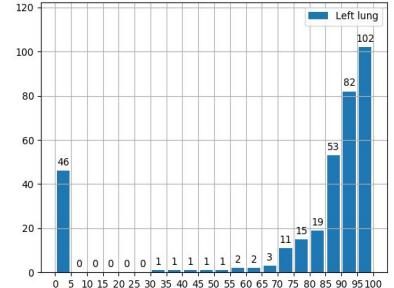
Dice coefficient - Disease



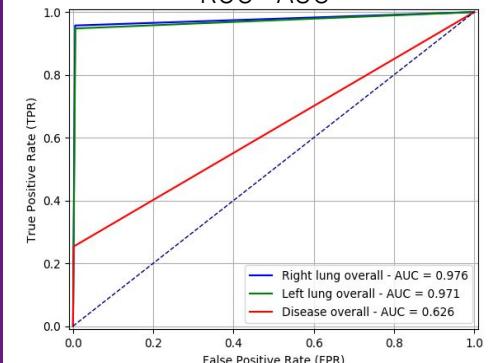
Dice coefficient - RL



Dice coefficient - LL



ROC - AUC



# 3

## Experimentation & results

- Experiment 4: Trying different loss functions

Using different loss functions on original U-Net model:

- Probabilistic Loss functions
  - Categorical Cross entropy
  - Poisson
  - KL - Divergence

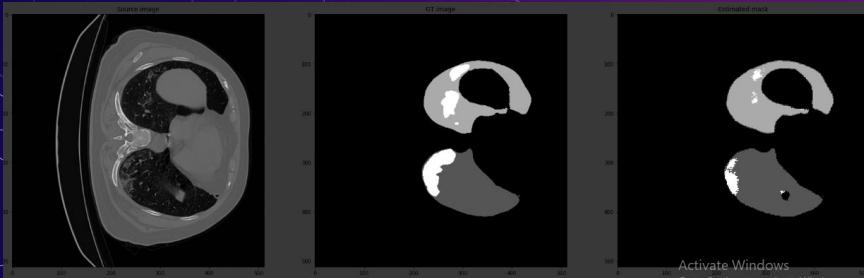
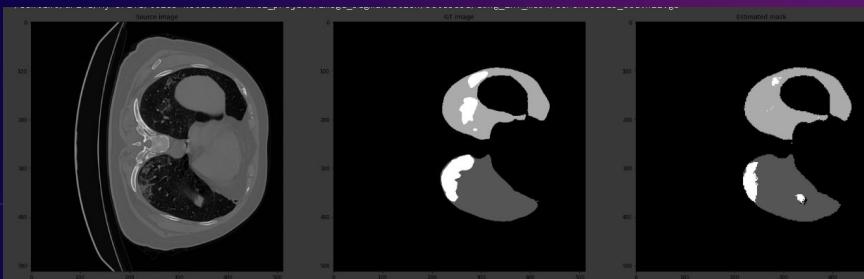
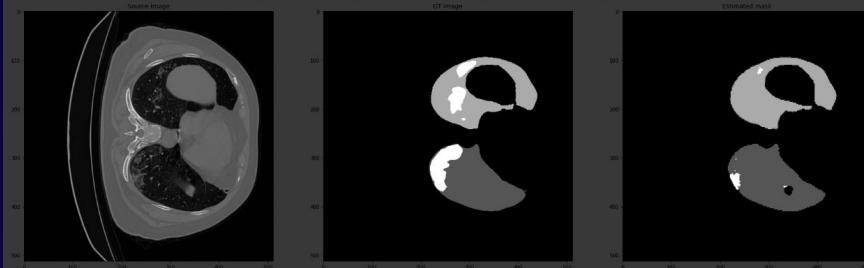
Regression loss functions

- Mean squared error
- Mean absolute error

# 3

## Experimentation & results

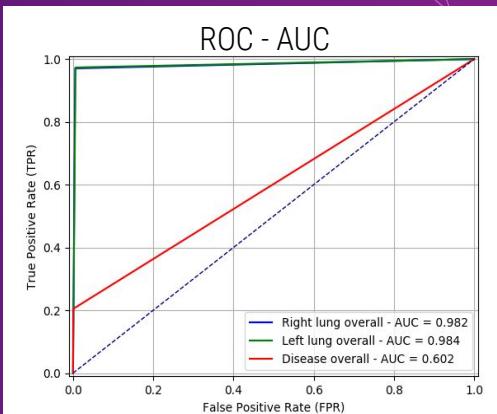
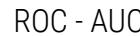
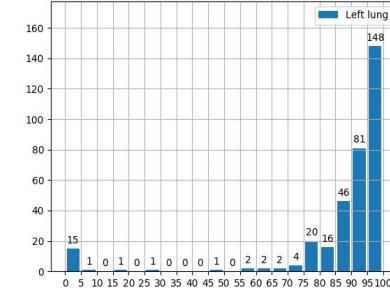
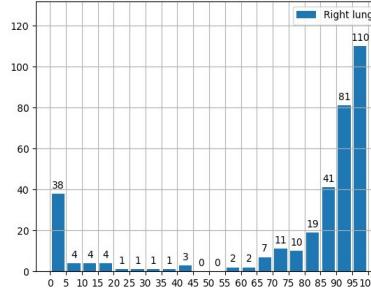
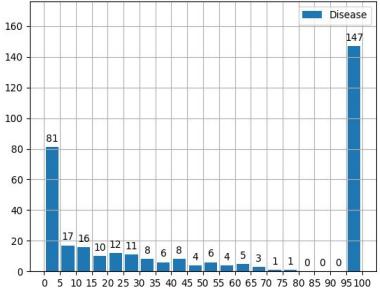
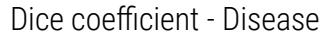
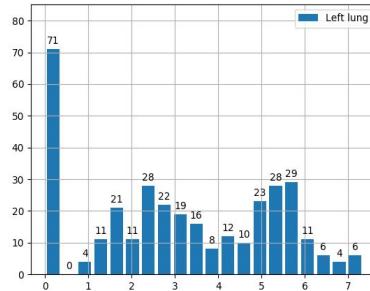
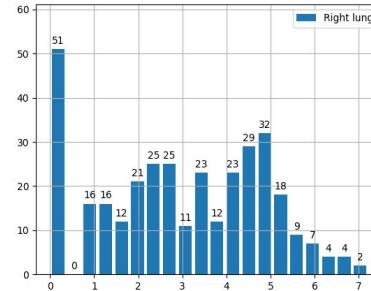
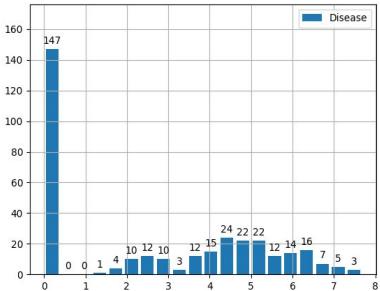
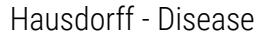
- Experiment 4 - A: Probabilistic loss functions



- Categorical cross entropy
- Poisson
- Kullback-Leibler divergence

# 3 Experimentation & results

- Experiment 4 - A: Statistical Analysis of Categorical Cross Entropy LF

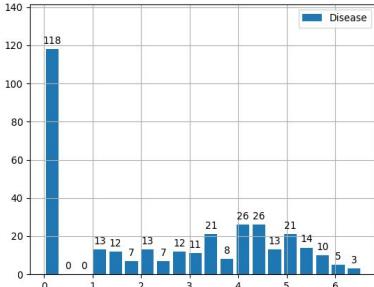


# 3

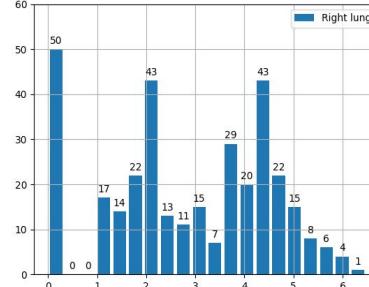
## Experimentation & results

- Experiment 4 - A: Statistical Analysis of Poisson LF

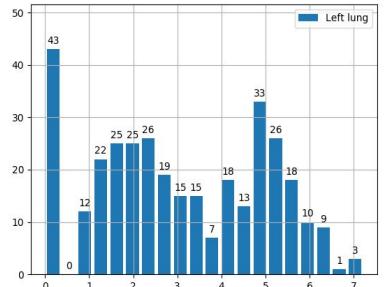
Hausdorff - Disease



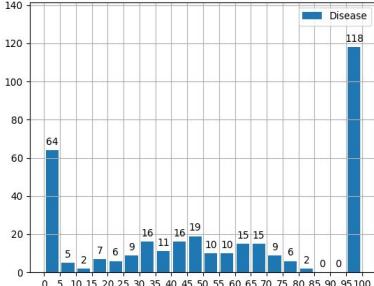
Hausdorff - RL



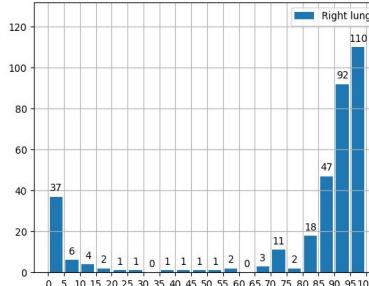
Hausdorff - LL



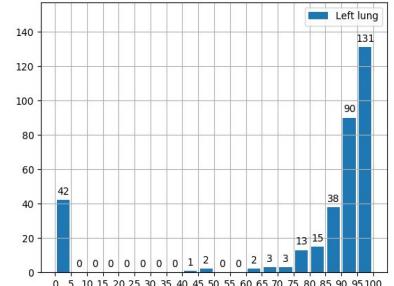
Dice coefficient - Disease



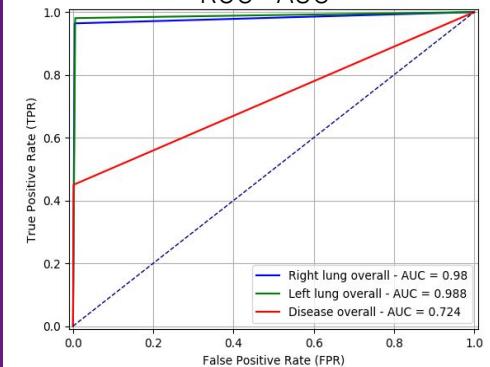
Dice coefficient - RL



Dice coefficient - LL



ROC - AUC

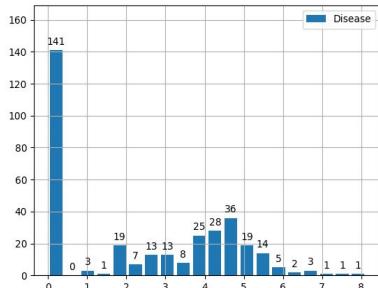


# 3

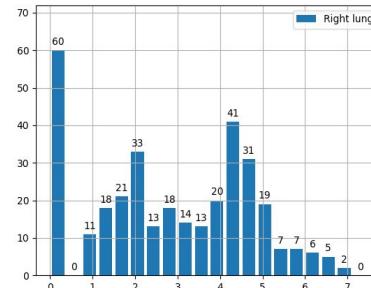
## Experimentation & results

### • Experiment 4 - A: Statistical Analysis of Kullback-Leibler divergence LF

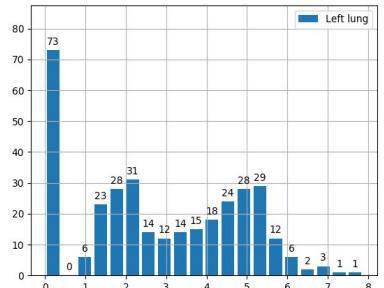
Hausdorff - Disease



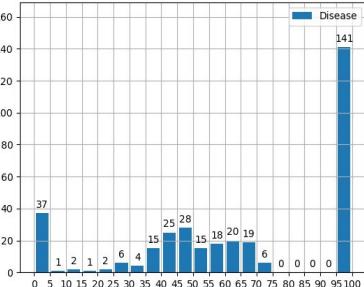
Hausdorff - RL



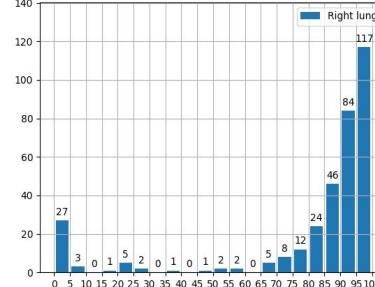
Hausdorff - LL



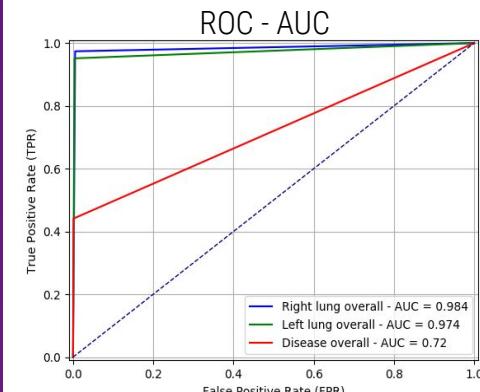
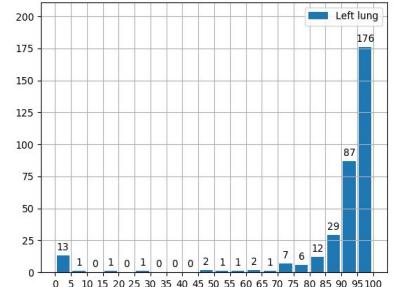
Dice coefficient - Disease



Dice coefficient - RL



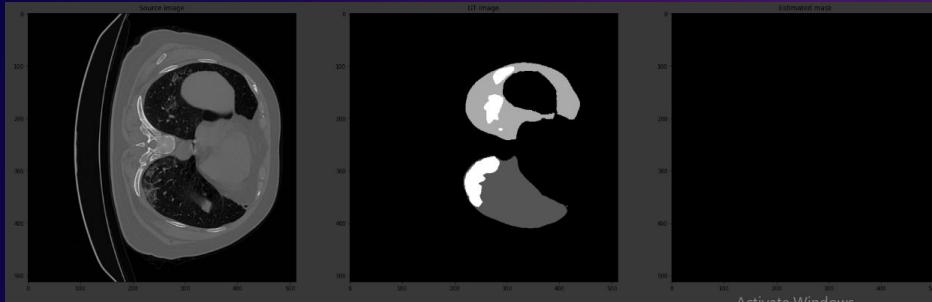
Dice coefficient - LL



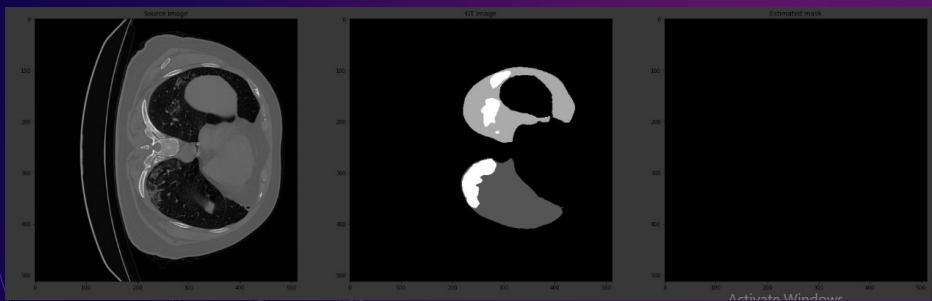
# 3

## Experimentation & results

- **Experiment 4 - B: Regression models LF**



- **Mean square error**



- **Mean absolute error**

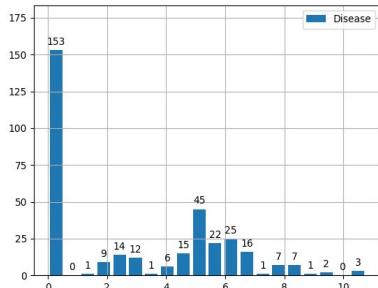
We can see that, even though the PLF work for the training, the absolutely fail into segmenting the image (It succeeded in segmenting only background)

# 3

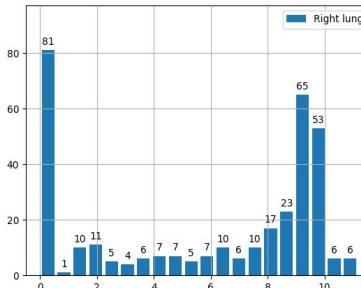
## Experimentation & results

### • Experiment 5 - B: Statistical Analysis of Absolute / Mean Squared error LF.

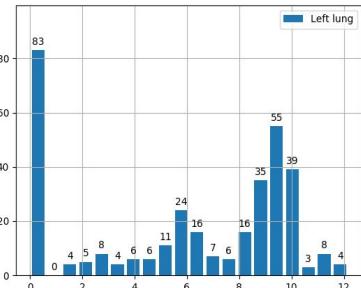
Hausdorff - Disease



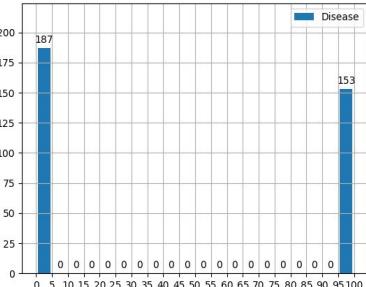
Hausdorff - RL



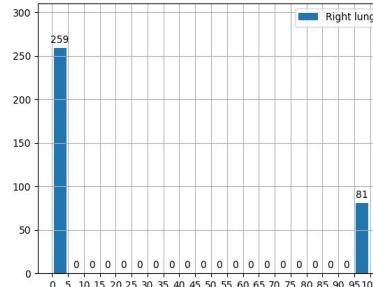
Hausdorff - LL



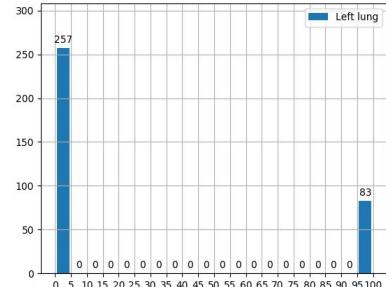
Dice coefficient - Disease



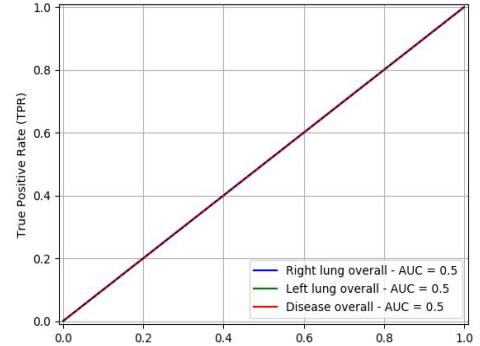
Dice coefficient - RL



Dice coefficient - LL



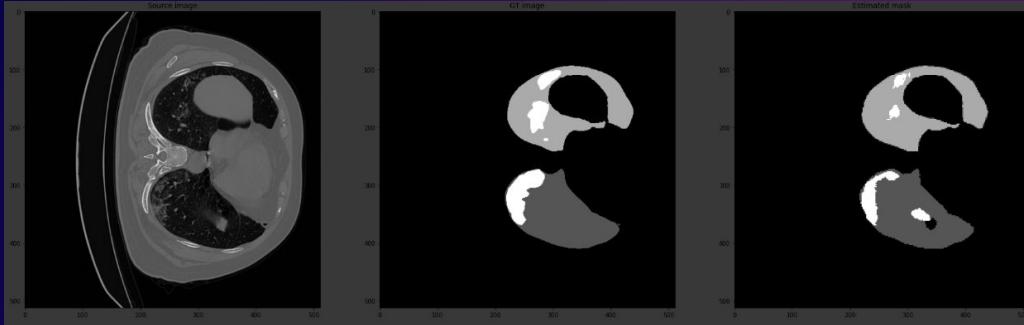
ROC - AUC



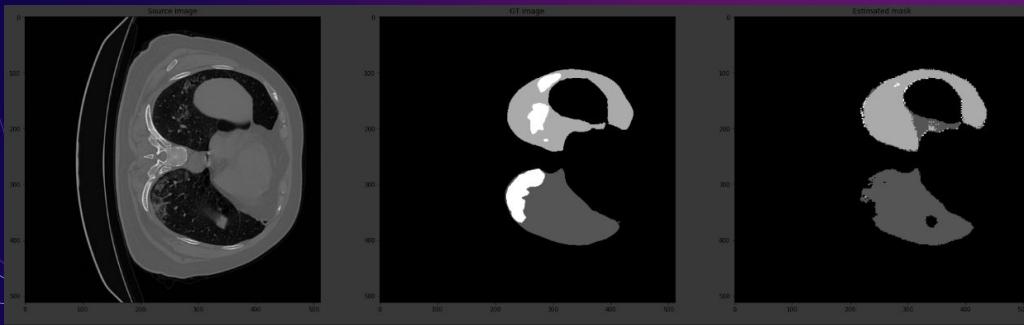
# 3

## Experimentation & results

- Experiment 5: Using Different Learning rates (Adam optimizer)



- Learning rate = 0.001



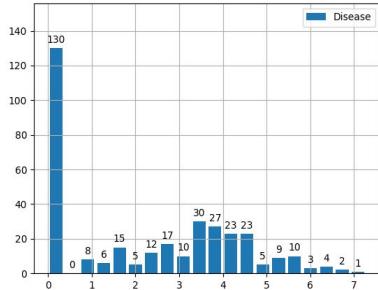
- Learning rate = 0.00001

# 3

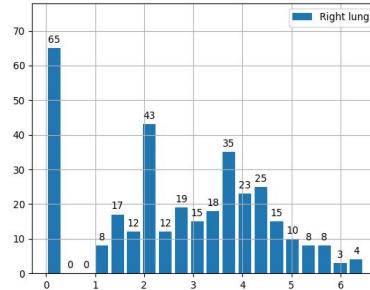
## Experimentation & results

- Experiment 5: Statistical Analysis of 0.001 learning rate

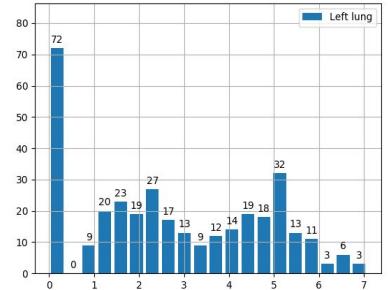
Hausdorff - Disease



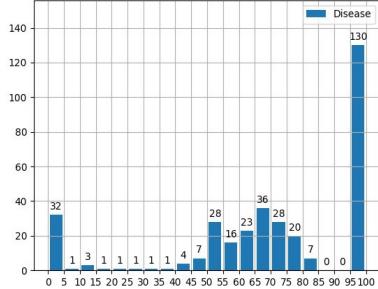
Hausdorff - RL



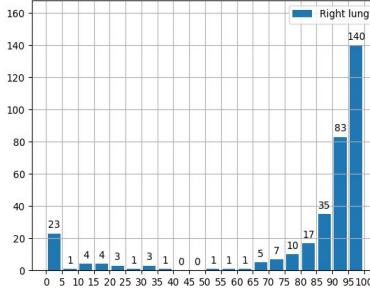
Hausdorff - LL



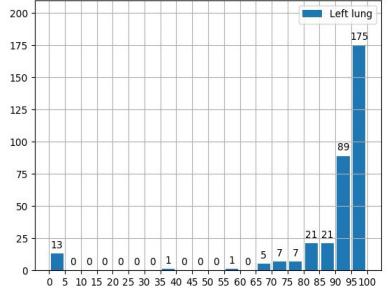
Dice coefficient - Disease



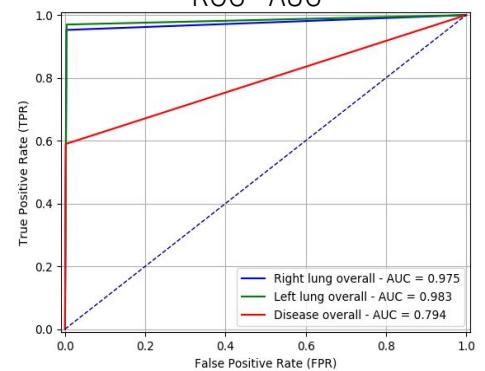
Dice coefficient - RL



Dice coefficient - LL



ROC - AUC

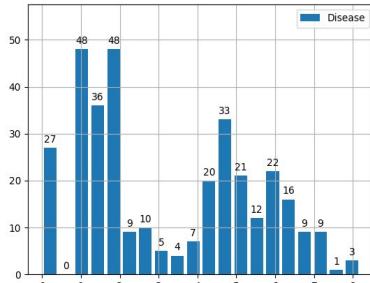


# 3

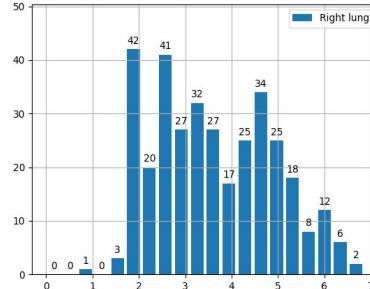
## Experimentation & results

- Experiment 5: Statistical Analysis of 0.00001 learning rate

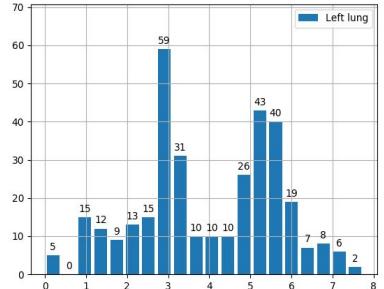
Hausdorff - Disease



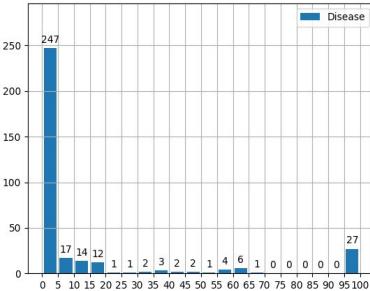
Hausdorff - RL



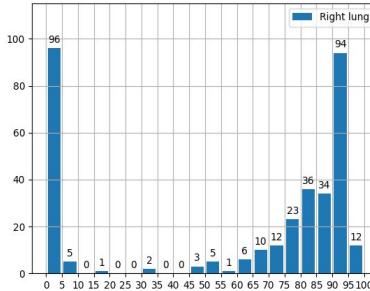
Hausdorff - LL



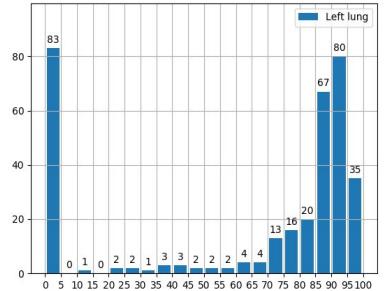
Dice coefficient - Disease



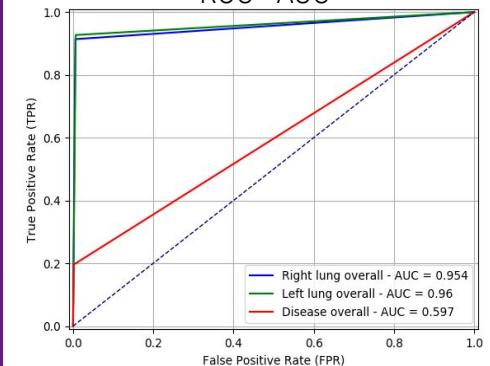
Dice coefficient - RL



Dice coefficient - LL



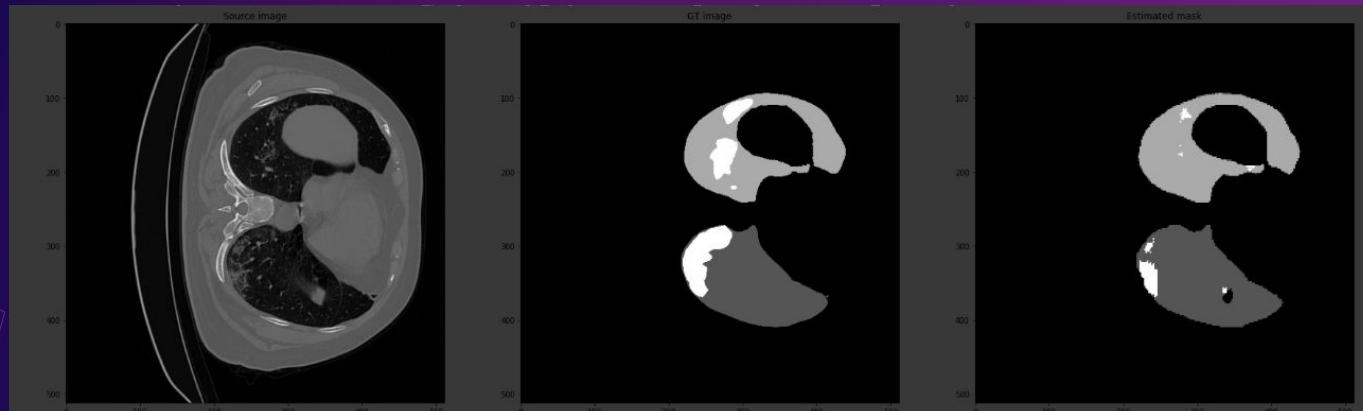
ROC - AUC



# 3

## Experimentation & results

- Experiment 6: Introducing inception module/layer to the U-Net model

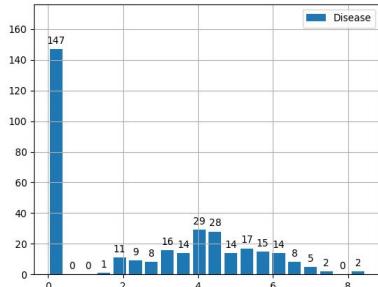


# 3

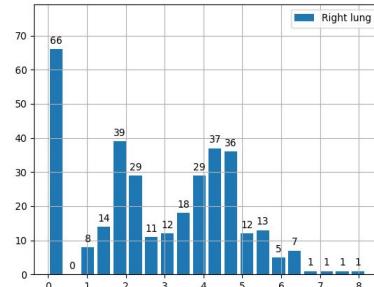
## Experimentation & results

- Experiment 6: Statistical Analysis of Inception Layer 1

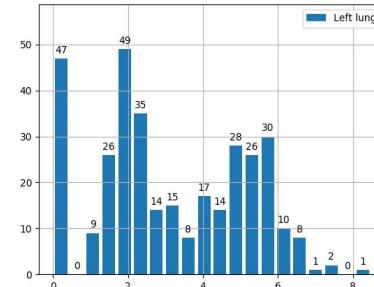
Hausdorff - Disease



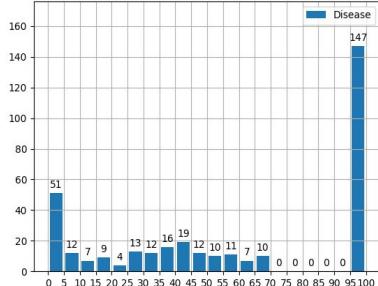
Hausdorff - RL



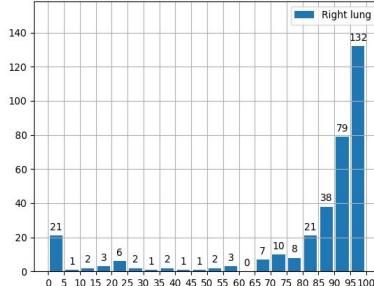
Hausdorff - LL



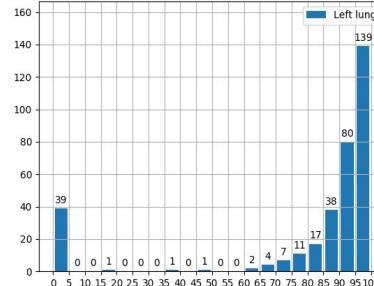
Dice coefficient - Disease



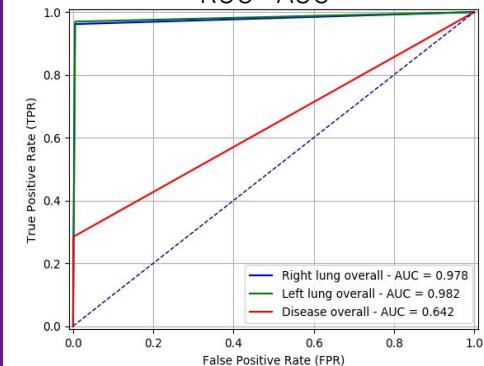
Dice coefficient - RL



Dice coefficient - LL



ROC - AUC



# 3

## Experimentation & results

- So far, we saw that the inception module works well with U-Net model. The biggest challenge here is to decide the size of the filters for this module
- Also, the value of 0.001 for the learning rate increases the performance of the model
- Some overfitting might be happening
- Further experiments made for testing the performance of the model with different batch size and epochs value

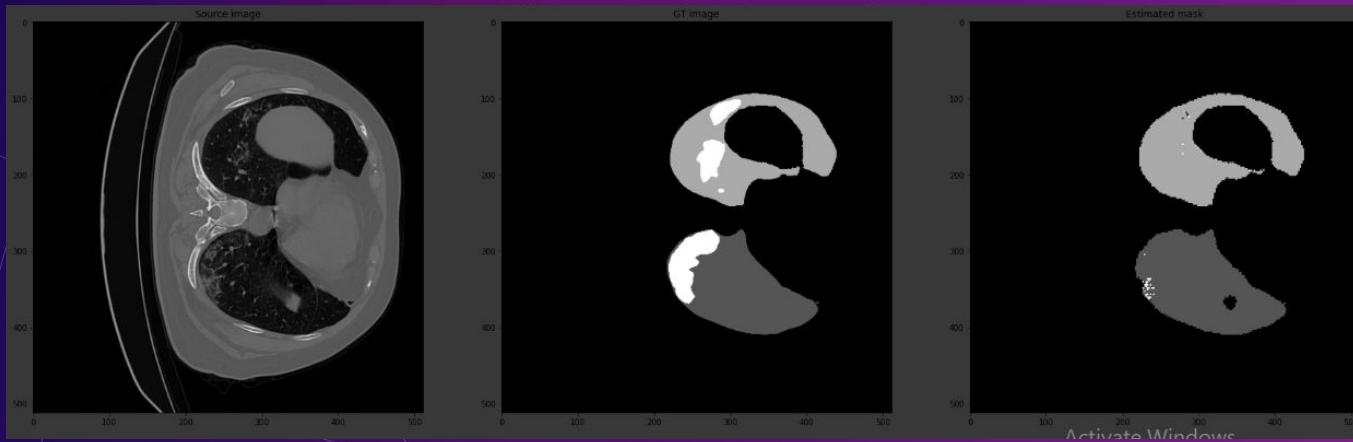
# 3

## Experimentation & results

- Experiment 7: Reducing Epoch and increasing Batch size

### Experiment details

- Epoch = 4
- Batch size = 12

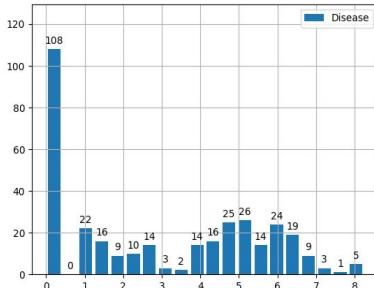


# 3

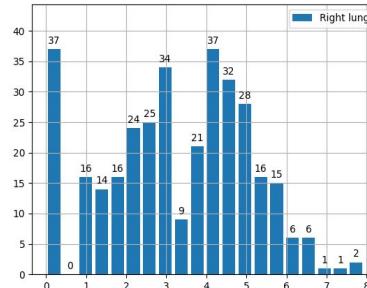
## Experimentation & results

- **Experiment 7: Statistical Analysis of Reducing Epoch and increasing Batch size**

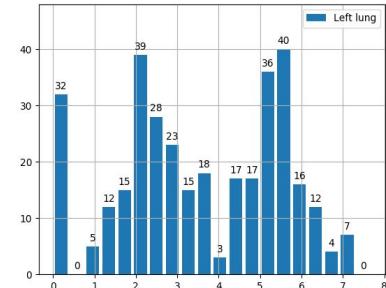
Hausdorff - Disease



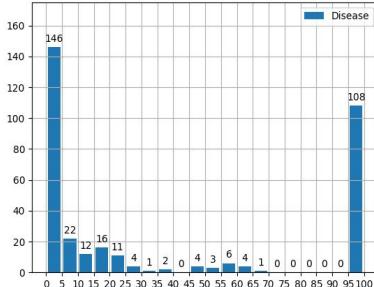
Hausdorff - RL



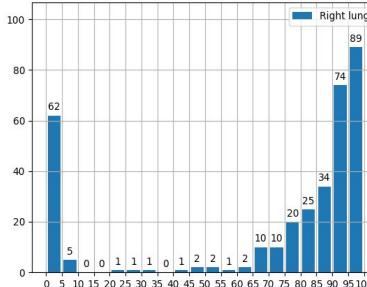
Hausdorff - LL



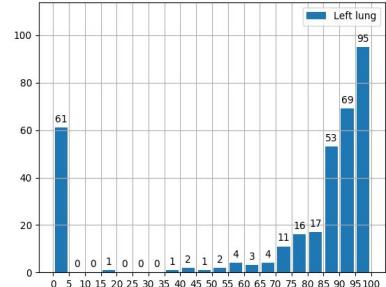
Dice coefficient - Disease



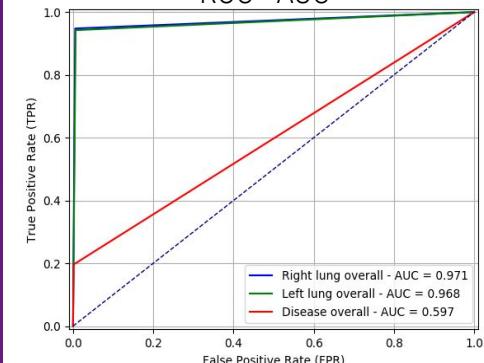
Dice coefficient - RL



Dice coefficient - LL



ROC - AUC



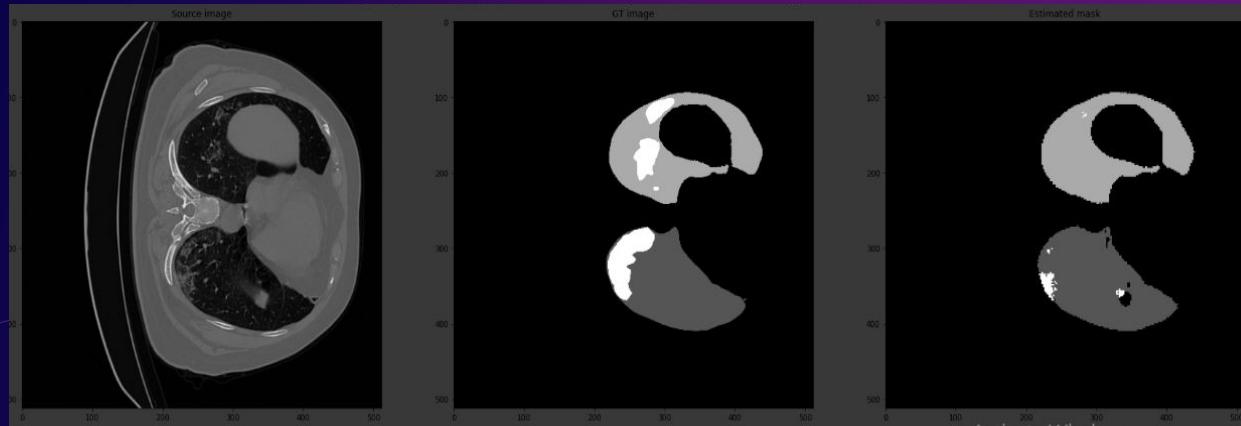
# 3

## Experimentation & results

- Experiment 8: Using categorical cross entropy for inception U-net model

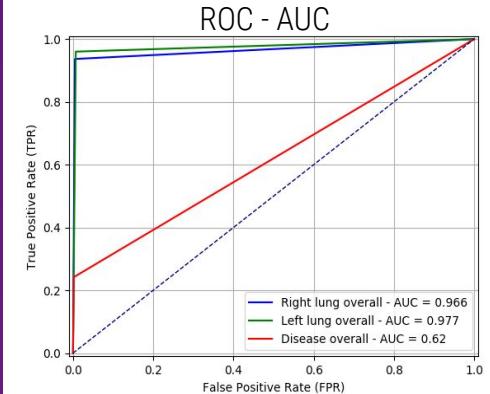
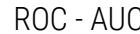
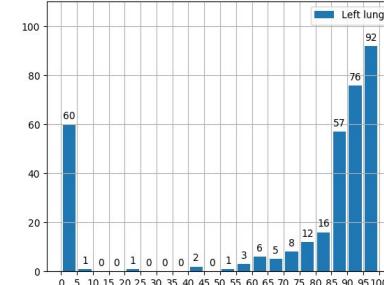
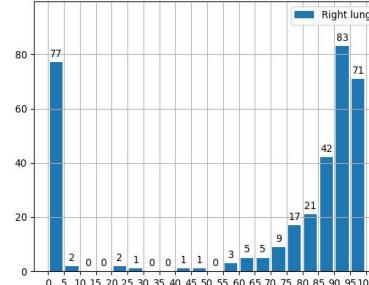
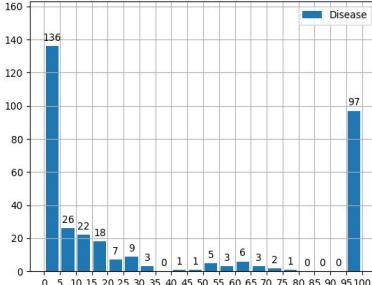
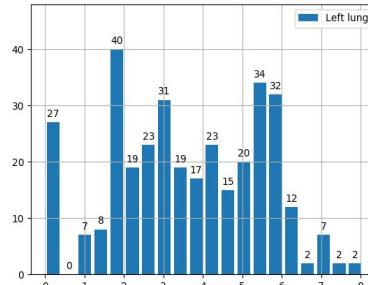
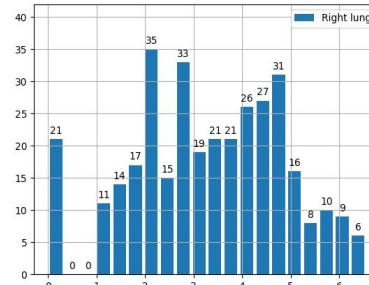
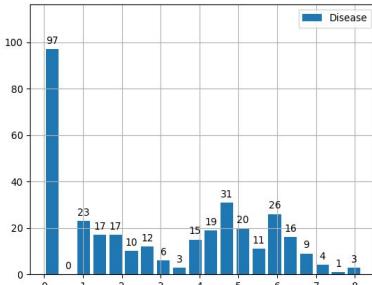
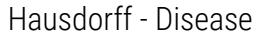
### Experimental Details

- Using Categorical cross entropy instead of binary cross entropy as a loss function
- Epochs = 5
- Batch size = 16



# 3 Experimentation & results

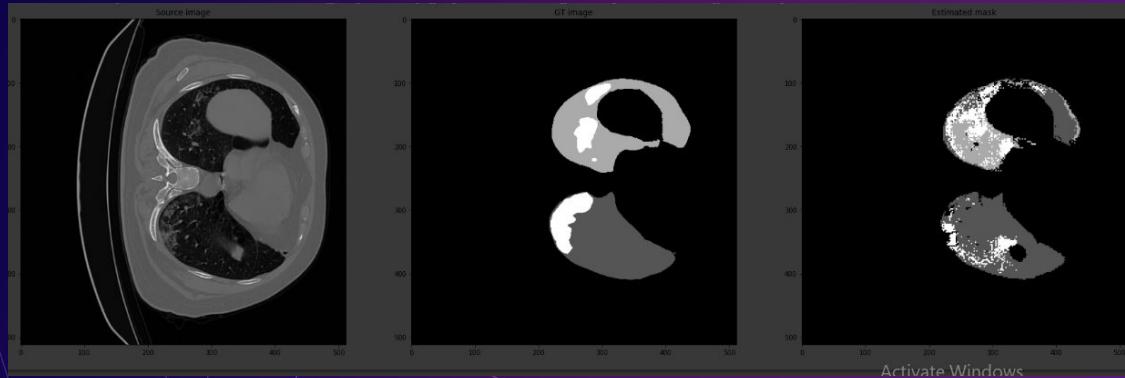
- **Experiment 8: Statistical Analysis of using categorical cross entropy for inception U-net model**



# 3

## Experimentation & results

- Experiment 9: Introducing second layer of Inception in U-net



### Experimental Details

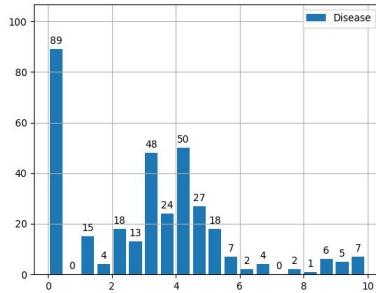
- **Total Params:** **8,990,692**
- **2 layers of inception**
- **2 drop out layers**
- **Epochs = 5**
- **Batch size = 12**

# 3

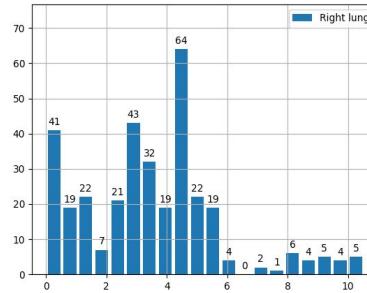
## Experimentation & results

- Experiment 9: Statistical Analysis of Introducing second layer of Inception in U-net

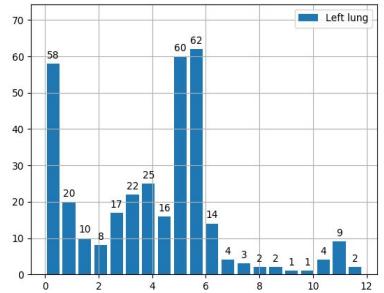
Hausdorff - Disease



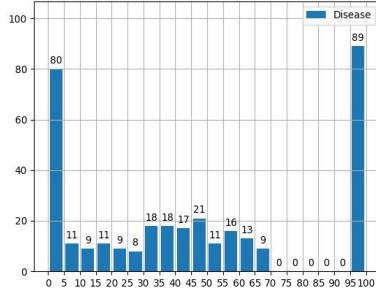
Hausdorff - RL



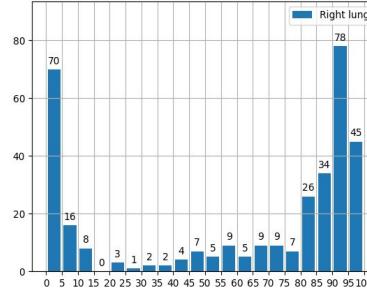
Hausdorff - LL



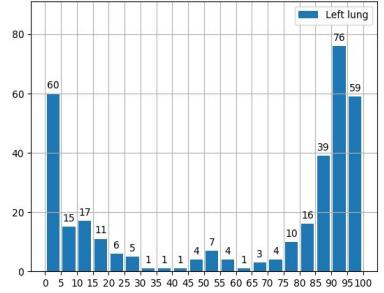
Dice coefficient - Disease



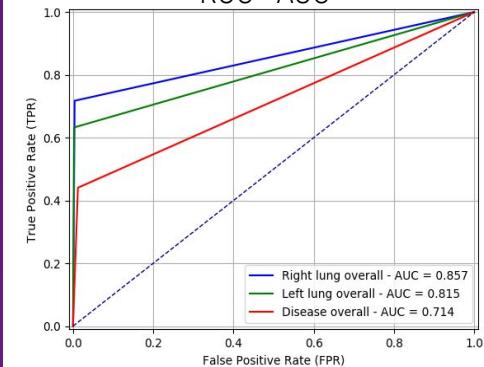
Dice coefficient - RL



Dice coefficient - LL



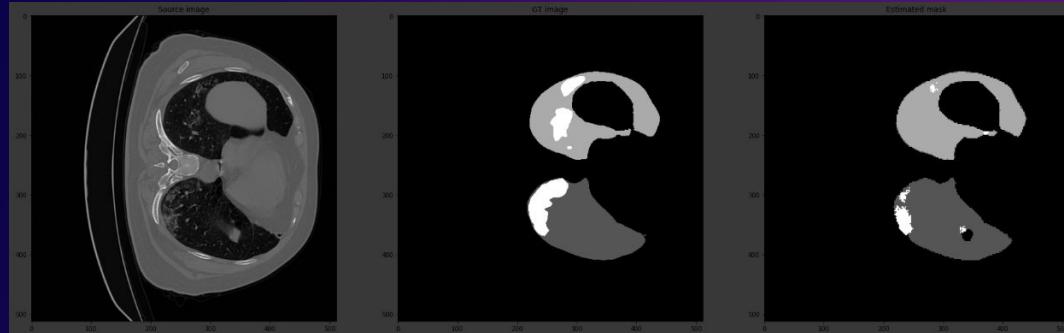
ROC - AUC



# 3

## Experimentation & results

- Experiment 10: Adding One inception layer and one DO (30%) in initial U-net



### Experimental details

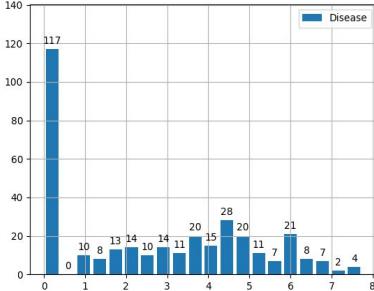
- 1 Drop out layer
- 1 Inception layer in unet
- Categorical cross entropy
- epochs = 8
- Batch size = 8
- Adam optimizer. LR 0.0001

# 3

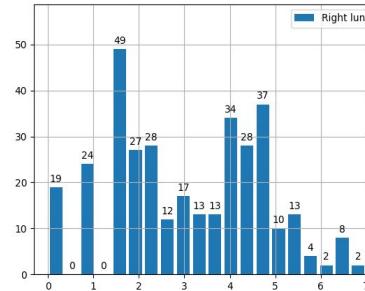
## Experimentation & results

- Experiment 10: Statistical Analysis of Adding One inception layer and one DO (30%) in initial U-net

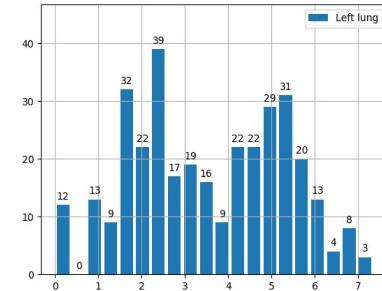
Hausdorff - Disease



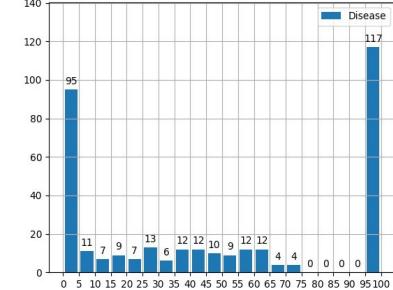
Hausdorff - RL



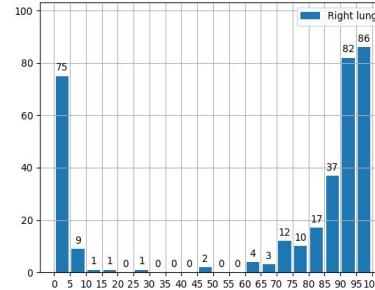
Hausdorff - LL



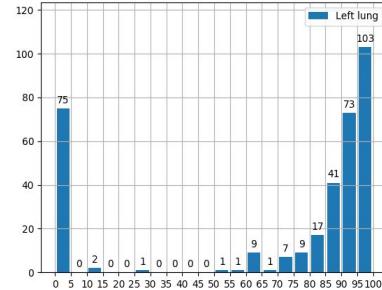
Dice coefficient - Disease



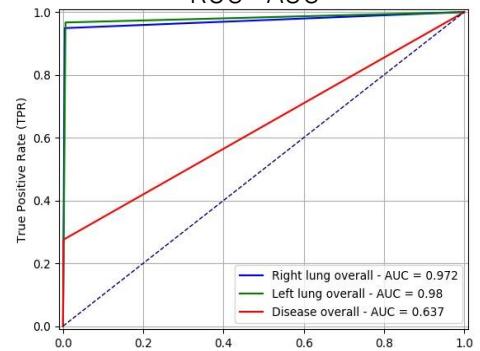
Dice coefficient - RL



Dice coefficient - LL



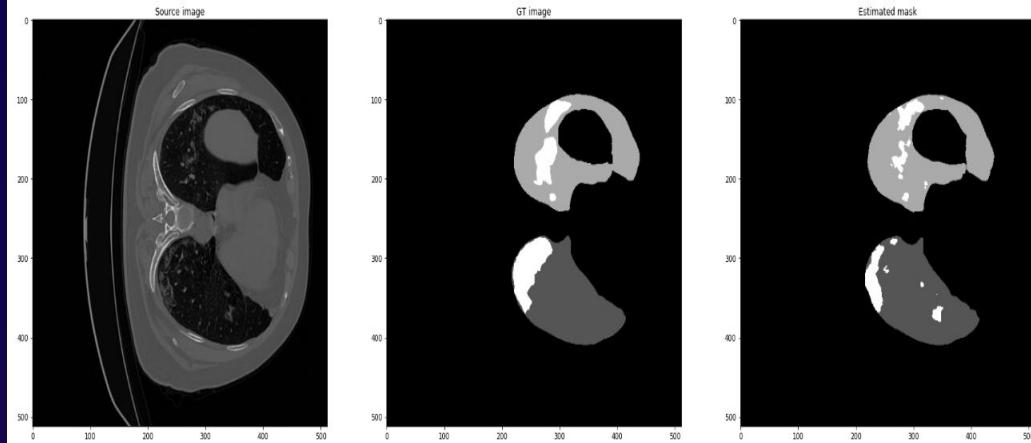
ROC - AUC



# 3

## Experimentation & results

- Experiment 11: Base U-Net model using EfficientNet pre-trained weights



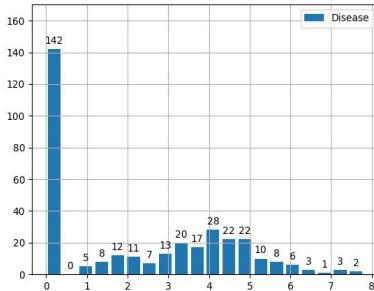
- Optimizer: Adam - LR = 0.0001
- Activation Func.: Sigmoid
- LF: Categorical Cross Entropy
- Epochs: 10
- Batch size: 8

# 3

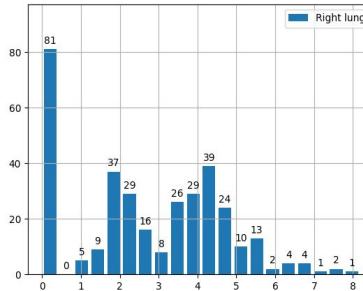
## Experimentation & results

- Experiment 11: Base U-Net model using EfficientNet pre-trained weights

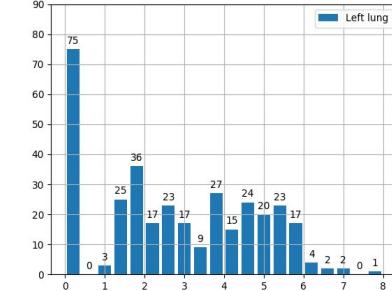
Hausdorff - Disease



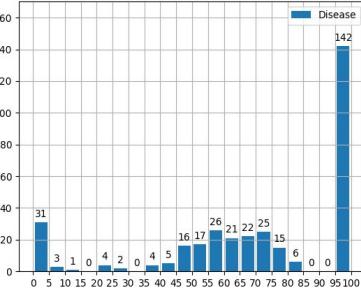
Hausdorff - RL



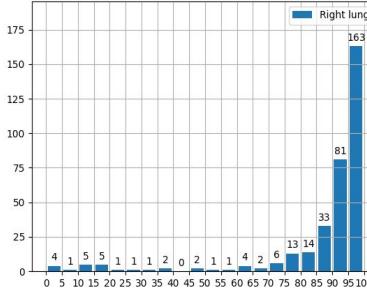
Hausdorff - LL



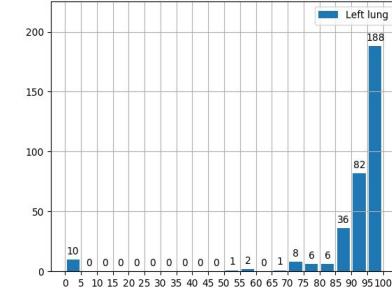
Dice coefficient - Disease



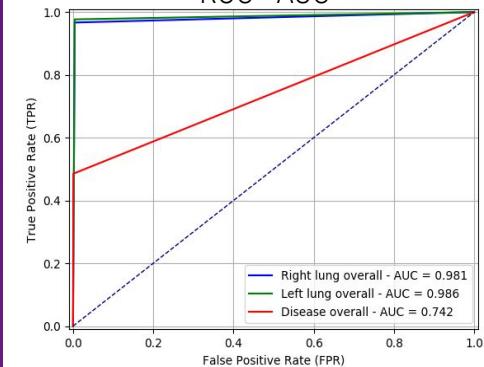
Dice coefficient - RL



Dice coefficient - LL



ROC - AUC



## 4 Conclusions & Further improvements

- The EfficientNet model performs as good as our custom model
- Since there are more pixels about the lungs than for the disease, the model performs better on them than in the disease area
- Even only 20 CT scans are provided, it is easy to get 99% accuracy for the lungs using deep learning
- Inception modules showed to improve the segmentation quality
- Further work should include:
  - Further studies of dropout layers are required. Interdependent learning
  - Data augmentation is strongly suggested
  - Batch normalization when using sigmoid activation function
  - Reduction of the background area at each image

# Questions?

# Thank you!