# **COVID 19 SEGMENTATION API**

#### **GUIDE**

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#### **Team Members**

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# INTRODUCTION

- COVID-19 is an infectious disease caused by the SARS-CoV-2
- Common method detecting COVID-19 RT-PCR testing,
- CT scan imaging fast and cost-effective alternative
- Deep learning techniques U-Net models
- This promising research field significant step towards rapid and reliable

COVID-19 detection

# **MOTIVATION**

• Detecting COVID-19 from CT-scan image using CNN models is a highly relevant and timely project, given the ongoing global pandemic.

- Creating a reliable CNN model to detect COVID-19 from CT-scans can assist healthcare professionals in early diagnosis and treatment, utilizing a common diagnostic tool.
- Developing a CNN model for CT-scan-based COVID-19 detection can aid in patient triage and improve outcomes in the fight against the pandemic.
   Ultimately, this project has the potential to contribute to the fight against COVID-19 and improve patient outcomes.

# PROBLEM STATEMENT

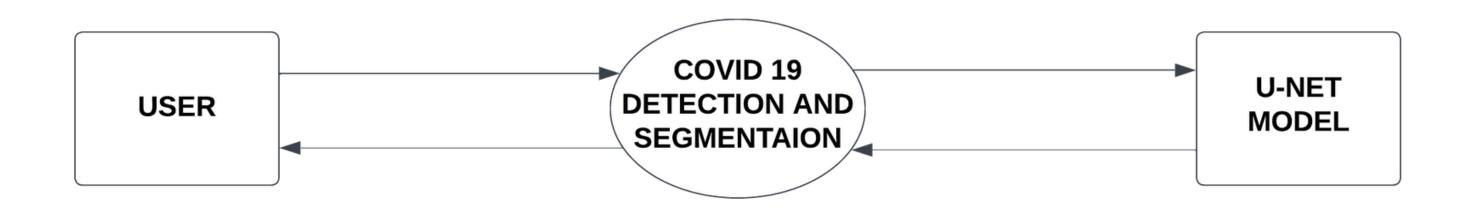
The accurate segmentation of COVID-19 infected regions in medical images is crucial for diagnosing and monitoring the disease. However, the manual segmentation process is time-consuming, subjective, and prone to human error. Therefore, there is a need to develop a reliable and efficient COVID-19 Segmentation API using the UNet deep learning model to automate the segmentation process and provide accurate and timely results.

# **OBJECTIVES**

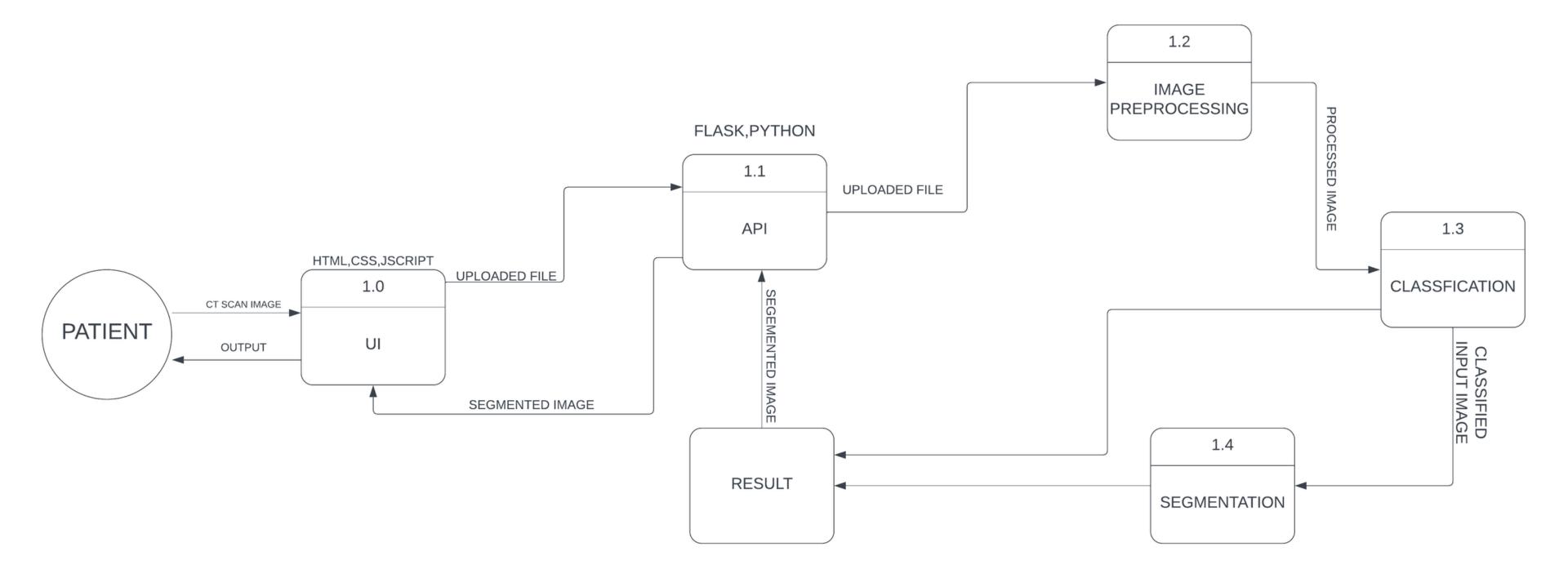
- Develop an API that allows users to perform automated segmentation of COVID-19 infected regions in medical images.
- Implement the UNet architecture, a deep learning model known for its effectiveness in image segmentation tasks, as the core algorithm for the API
- Optimize the model for efficient inference, ensuring that the API provides segmentation results in a timely manner
- Implement a user friendly interface.
- Provide comprehensive testing to validate the accuracy and reliability of the segmentation results across different types of COVID-19 lung images

# **DATA FLOW DIAGRAM**

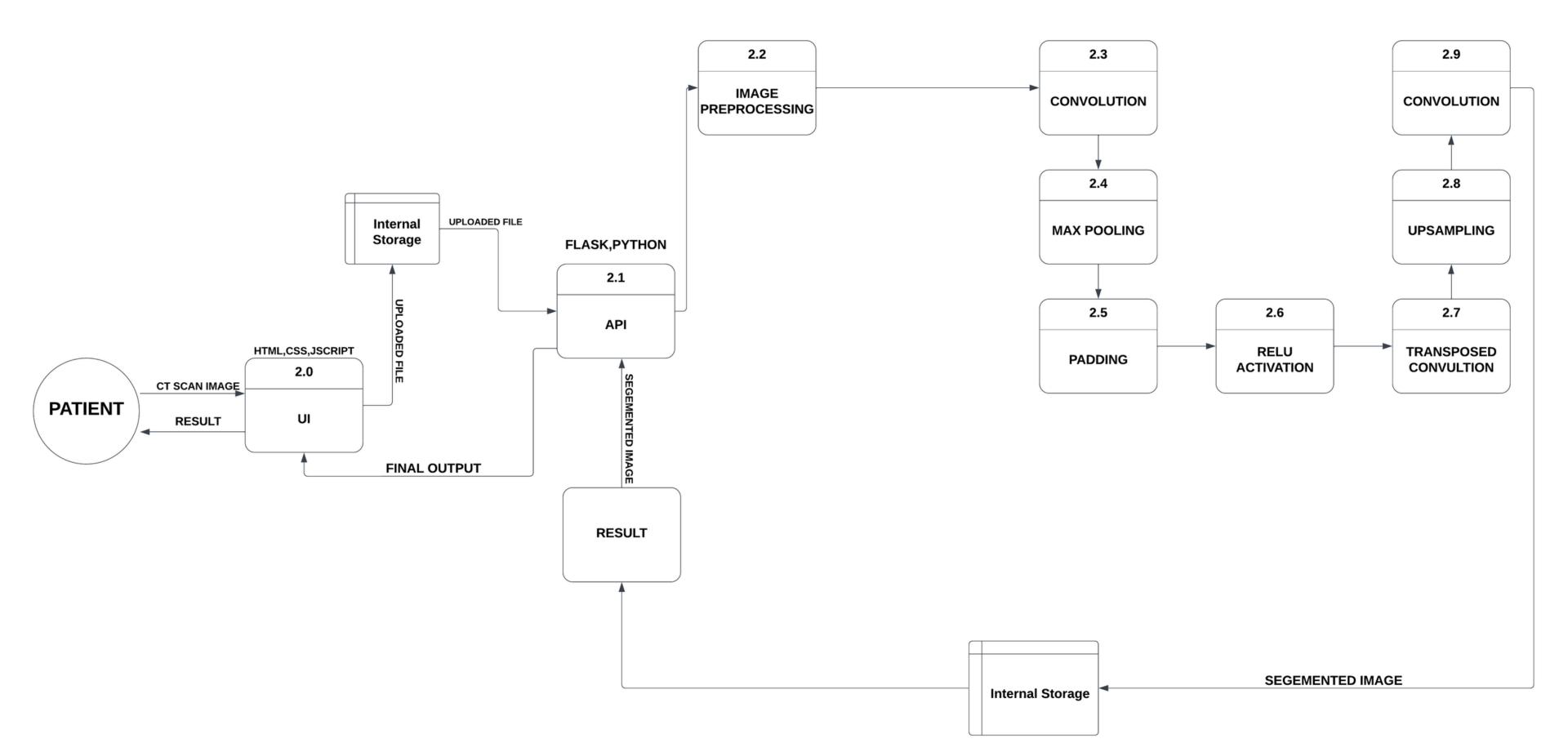
# **LEVEL 0**

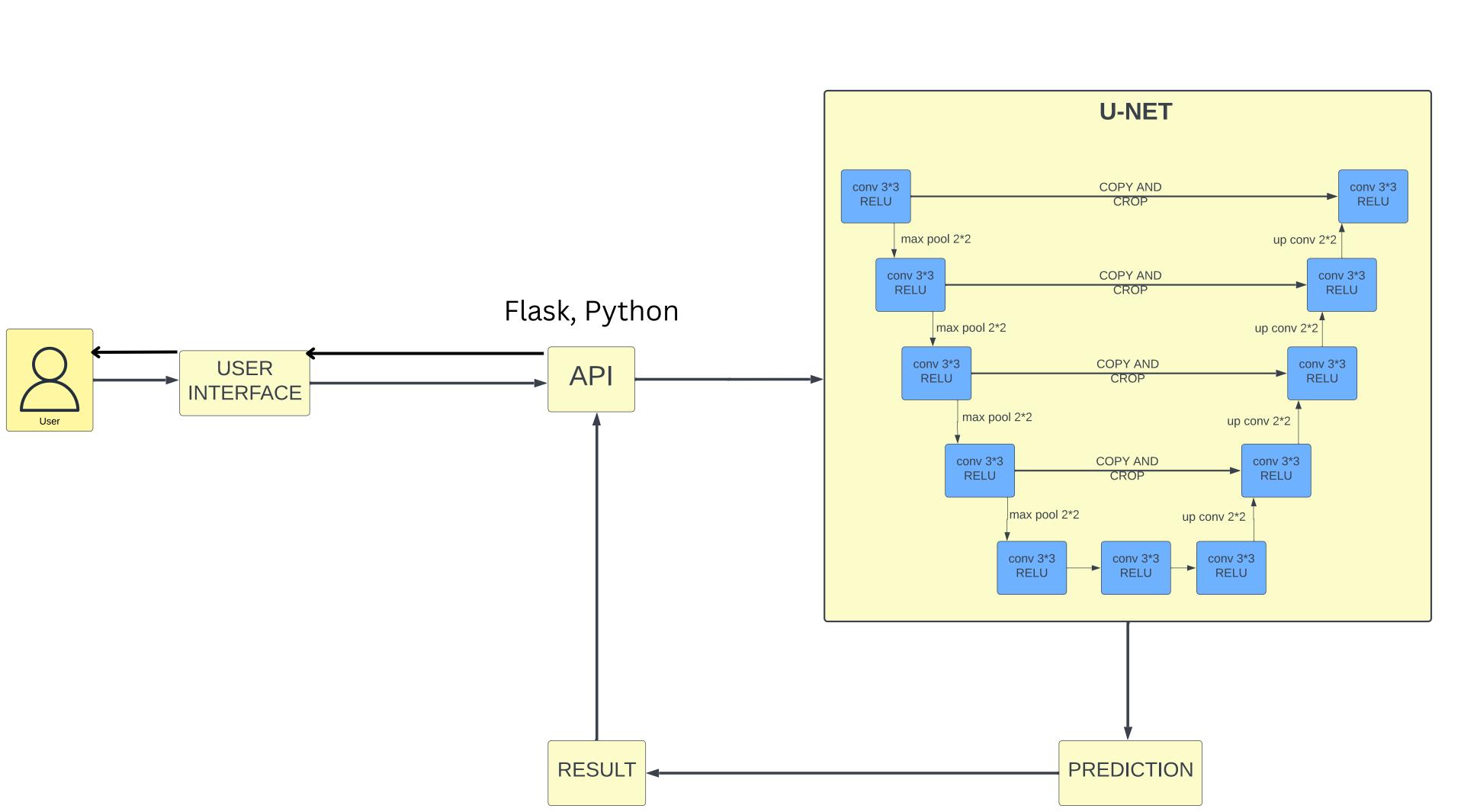


# LEVEL 1

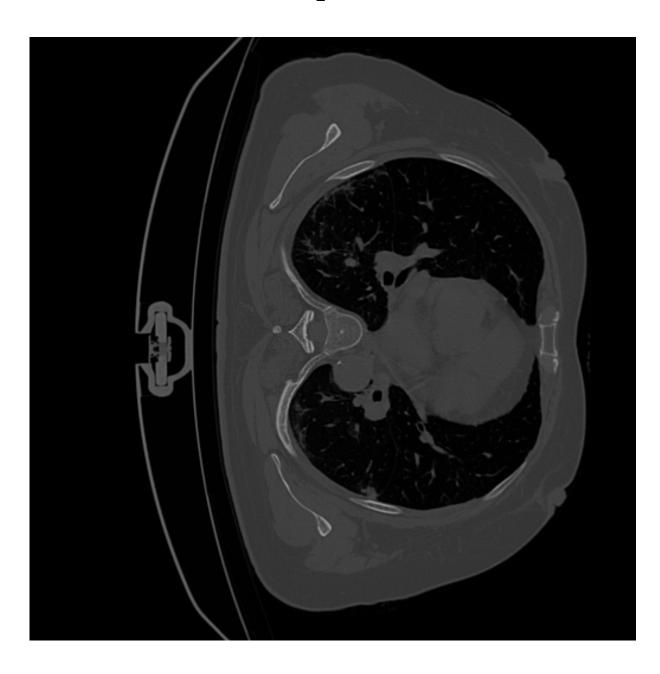


# LEVEL 2

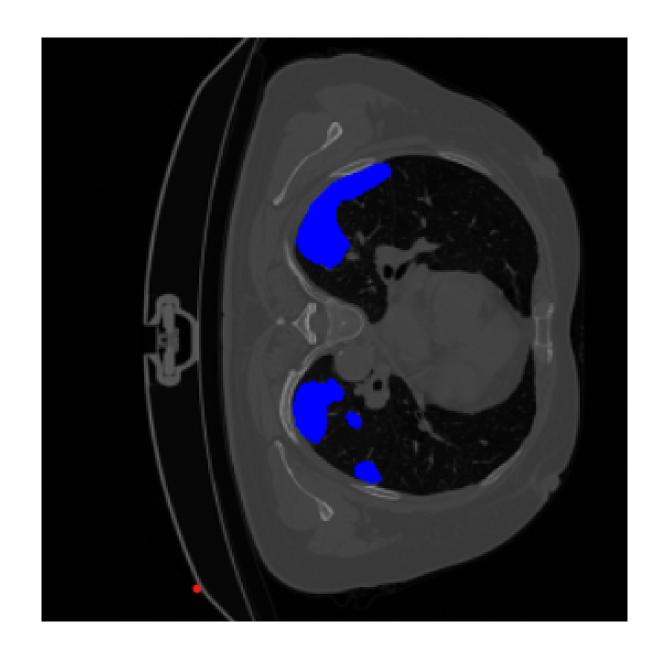




# Input



# Output



# **MODULE DETAILS**

- 1. Design the user interface
- 2.Build API
- 3. Collection of the dataset
- 4.Implement UNET Model
- 5. Prediction

# **DESIGNING USER INTERFACE**

- 1. Design an intuitive and user-friendly interface.
- 2.It consists of a simple UI that is connected through an API to the model.
- 3. An option to add the CT Scan will be provided.
- 4.The result is then fetched from the API and is displayed to the user.

# **BUILDING API**

- The back-end API will receive the CT-scan image from the front-end and process it using UNET model.
- To build an API a web framework such as Flask (Python), Express (Node.js), or Ruby on Rails (Ruby) is required.

• When the API receives the image, it will need to preprocess the image to prepare it for input to the UNET model.

# **COLLECTION & TRAINING**

• Public dataset from **Ma et al.** which consists of 700 annotated COVID-19 chest CT volumes.

• 3D volume set with annotated COVID-19 infection segmentation.

• It is used by the model to provide accurate and helpful responses to users.

# IMPLEMENTING U-NET MODEL

- Segmentation: It provides fine-grained information about the image as well as the shapes and boundaries of the objects.
- Encoding layers: Extracting features from the input image using convolution, padding, maxpooling operations.
- Decoding Layers: precise localization using transposed convolutions

# **SEGMENTATION & RESULT**

Receive a CT scanned image of a lung through an API to be Segmented by a Convolutional neural network.

# **SYSTEM REQUIREMENTS**

# Software Requirement

- Language: Python
- Supporting libraries like KERAS
- API framework: FLASK
- IDE: VS Code

# Hardware Requirement

- Processor : Intel(R) Xeon w-2255
   CPU(3.7Ghz)
- RAM: 64 GB
- GPU: RTX 3080 10 GB

# **COMPARISON WITH EXISTING SYSTEM**

- Current Existing systems are
  - Reverse-Transcription Polymerase Chain Reaction (RT-PCR)
  - Rapid Antigen Test
  - Chest X-Ray/ CT Scan

- Compared to these systems the model will have:
  - Lower False Positives / False Negatives
  - Easily Available
  - Lesser Turnaround Time
  - Reduced Cost

# WORKPLAN

# Week 1-3: Planning and research

- Define project requirements and objectives
- Research on CNN U-NET
- Determine feasibility of project within the given timeline and resources

# Week 4-5:Development Process

- Collect required training datas
- Training the U-NET model with the datas and tuning hyperparameters
- Developing API for the system
- Making user interface

# Week 6-7: Testing and refinement

- Test the model with sample data and check for accuracy
- Gather feedback from users and identify areas for improvement.

### **IMPLEMENTATION**

#### **IMPORTS**

```
import os
os.environ["TF_CPP_MIN_LOG_LEVEL"] = "2"
import numpy as np
import cv2
from glob import glob
from sklearn.model_selection import train_test_split
import tensorflow as tf
from tensorflow.keras.callbacks import ModelCheckpoint, CSVLogger, ReduceLROnPlateau
from tensorflow.keras.optimizers import Adam
from tensorflow.keras.metrics import Recall, Precision
from model import build_unet
from metrics import dice_loss, dice_coef, iou
```

All the necessary libraries & packages required to run the program are installed.

#### **CONVULTION BLOCK**

```
def conv_block(input, num_filters):
    x = Conv2D(num_filters, 3, padding="same")(input)
    x = BatchNormalization()(x)
    x = Activation("relu")(x)

x = Conv2D(num_filters, 3, padding="same")(x)
    x = BatchNormalization()(x)
    x = Activation("relu")(x)

return x
```

- The convolution operation, when performed on an image, extracts features from it.
- The number of filters in the Conv2D block is defined with num\_filters
- Normalization ensures that all the values are normalized between 0 and 1.
- The activation function chosen is the Relu function.
- A single convolution block contains two Conv2D layers, 2 Normalization layers followed by an activation function

#### **ENCODER BLOCK & DECODER BLOCK**

```
def encoder_block(input, num_filters):
    x = conv_block(input, num_filters)
    p = MaxPool2D((2, 2))(x)
    return x, p

def decoder_block(input, skip_features, num_filters):
    x = Conv2DTranspose(num_filters, (2, 2), strides=2, padding="same")(input)
    x = Concatenate()([x, skip_features])
    x = conv_block(x, num_filters)
    return x
```

The above block contains the code for U-Net Architecture.It has 2 parts:

- Encoder Block
  - Responsible for extracting patterns in the image, such as infected regions.
  - Uses convolutions and max-pooling layers.
  - Convolutions make patterns more obscure and reduce image size.
  - Max-pooling ensures important features are retained.
- Decoder Block
  - Opposite of Encoder Block
  - Localizes the extracted features

#### **U-NET MODEL**

```
def build_unet(input_shape):
    inputs = Input(input_shape)
   s1, p1 = encoder_block(inputs, 64)
   s2, p2 = encoder_block(p1, 128)
    s3, p3 = encoder_block(p2, 256)
   s4, p4 = encoder_block(p3, 512)
   b1 = conv_block(p4, 1024)
   d1 = decoder_block(b1, s4, 512)
   d2 = decoder_block(d1, s3, 256)
   d3 = decoder_block(d2, s2, 128)
   d4 = decoder_block(d3, s1, 64)
   outputs = Conv2D(1, 1, padding="same", activation="sigmoid")(d4)
   model = Model(inputs, outputs, name="U-Net")
    return model
```

#### The U-Net model consists of:

- 4 Encoder blocks
- Bottle neck layer
- 4 Decoder Blocks
- Activation Function : Sigmoid Function

#### **MODEL SUMMARY**

The summary of the model that we have constructed to perform segmentation of covid-19 infected CT scans.

Model: "U-Net"			
Layer (type)	Output Shape	Param #	Connected to
input_3 (InputLayer)	[(None, 512, 512, 3 )]	Ø	D .
conv2d_16 (Conv2D)	(None, 512, 512, 64 )	1792	['input_3[0][0]']
<pre>batch_normalization_14 (BatchN ormalization)</pre>	(None, 512, 512, 64 )	256	['conv2d_16[0][0]']
activation_14 (Activation)	(None, 512, 512, 64 )	0	['batch_normalization_14[0][0]']
<pre>max_pooling2d_6 (MaxPooling2D)</pre>	(None, 256, 256, 64 )	0	['activation_14[0][0]']
conv2d_17 (Conv2D)	(None, 256, 256, 12 8)	73856	['max_pooling2d_6[0][0]']
<pre>batch_normalization_15 (BatchN ormalization)</pre>	(None, 256, 256, 12 8)	512	['conv2d_17[0][0]']
activation_15 (Activation)	(None, 256, 256, 12 8)	0	['batch_normalization_15[0][0]']
<pre>max_pooling2d_7 (MaxPooling2D)</pre>	(None, 128, 128, 12	0	['activation_15[0][0]']

```
activation_18 (Activation)
                                                                ['batch_normalization_18[0][0]']
                               (None, 128, 128, 25 0
                               6)
 conv2d_transpose_7 (Conv2DTran (None, 256, 256, 12 131200
                                                                ['activation_18[0][0]']
 spose)
                                                                ['conv2d_transpose_7[0][0]',
 concatenate 7 (Concatenate)
                               (None, 256, 256, 25 0
                                                                  'activation_15[0][0]']
 conv2d_21 (Conv2D)
                               (None, 256, 256, 12 295040
                                                                ['concatenate_7[0][0]']
                               8)
batch_normalization_19 (BatchN (None, 256, 256, 12 512
                                                                ['conv2d_21[0][0]']
ormalization)
 activation_19 (Activation)
                                                                ['batch_normalization_19[0][0]']
                                (None, 256, 256, 12 0
 conv2d transpose 8 (Conv2DTran (None, 512, 512, 64 32832
                                                                ['activation_19[0][0]']
 spose)
                                                                ['conv2d_transpose_8[0][0]',
 concatenate_8 (Concatenate)
                               (None, 512, 512, 12 0
                                                                  'activation_14[0][0]']
                                                                ['concatenate_8[0][0]']
 conv2d 22 (Conv2D)
                               (None, 512, 512, 64 73792
                                                                ['conv2d_22[0][0]']
batch normalization 20 (BatchN (None, 512, 512, 64 256
ormalization)
                                                                 ['batch_normalization_20[0][0]']
 activation_20 (Activation)
                                (None, 512, 512, 64 0
 conv2d_23 (Conv2D)
                                                                 ['activation_20[0][0]']
                                (None, 512, 512, 1) 65
Total params: 3,793,985
Trainable params: 3,791,169
Non-trainable params: 2,816
```

The input shape of the image is (512,512) The output shape of the image is (512,512)

### TRAINING & TESTING

```
""" Global parameters """
H = 512
W = 512
def create_dir(path):
    """ Create a directory. """
   if not os.path.exists(path):
        os.makedirs(path)
def load_data(path, split=0.1):
    images = sorted(glob(os.path.join(path, "images", "*.png")))
   masks = sorted(glob(os.path.join(path, "masks", "*.png")))
    split_size = int(len(images) * split)
    train_x, valid_x = train_test_split(images, test_size=split_size, random_state=42)
    train_y, valid_y = train_test_split(masks, test_size=split_size, random_state=42)
    train_x, test_x = train_test_split(train_x, test_size=split_size, random_state=42)
    train_y, test_y = train_test_split(train_y, test_size=split_size, random_state=42)
    return (train_x, train_y), (valid_x, valid_y), (test_x, test_y)
```

Total dataset: 700 images

Training Data: 400 images

Validation : 200 images

Testing : 100 images

#### MODEL DEFINITION

```
model = build_unet((H, W, 3))
metrics = [dice_coef, iou, Recall(), Precision()]
model.compile(loss=dice_loss, optimizer=Adam(lr), metrics=metrics)

callbacks = [
    ModelCheckpoint(model_path, verbose=1, save_best_only=True),
    ReduceLROnPlateau(monitor='val_loss', factor=0.1, patience=5, min_lr=1e-7, verbose=1),
    CSVLogger(csv_path)
]

model.fit(
    train_dataset,
    epochs=num_epochs,
    validation_data=valid_dataset,
    callbacks=callbacks
)
```

#### The model is build by:

- Defining it's structure
- Setting metrics
- Implementing callbacks when desired performance is achieved
- model.fit() is used to train the model with pre-defined epochs & datasets

### **METRICS**

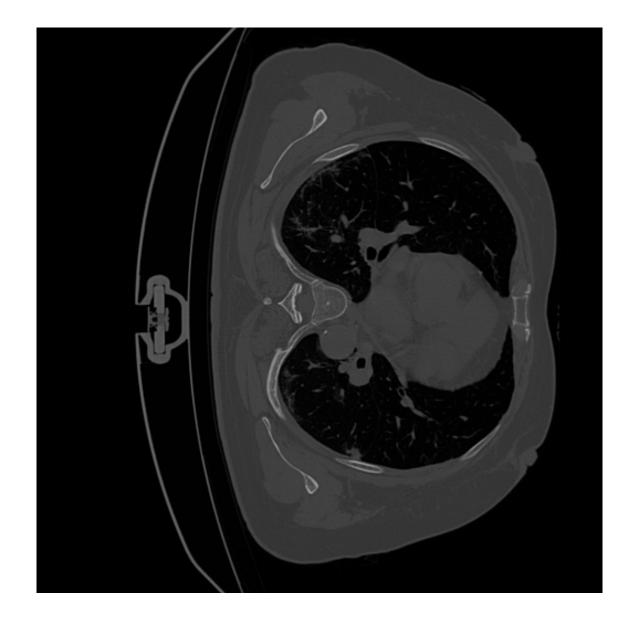
```
def iou(y_true, y_pred):
    def f(y_true, y_pred):
       intersection = (y_true * y_pred).sum()
       union = y_true.sum() + y_pred.sum() - intersection
       x = (intersection + 1e-15) / (union + 1e-15)
       x = x.astype(np.float32)
        return x
   return tf.numpy_function(f, [y_true, y_pred], tf.float32)
smooth = 1e-15
def dice_coef(y_true, y_pred):
   y_true = tf.keras.layers.Flatten()(y_true)
   y_pred = tf.keras.layers.Flatten()(y_pred)
   intersection = tf.reduce sum(y true * y pred)
    return (2. * intersection + smooth) / (tf.reduce_sum(y_true) + tf.reduce_sum(y_pred) + smooth)
def dice_loss(y_true, y_pred):
   return 1.0 - dice_coef(y_true, y_pred)
```

The metrics that we have chosen to evaluate our model are:

- Dice coefficient: (2 \* intersection) / (sum of sizes).
- Intersection over Union (IoU): IoU = intersection / union
- Precision : Precision = True Positives / (True Positives + False Positives)
- Recall : Recall = True Positives / (True Positives + False Negatives)

# **SAMPLE TRAINING DATA**

# CT SCAN



# MASK

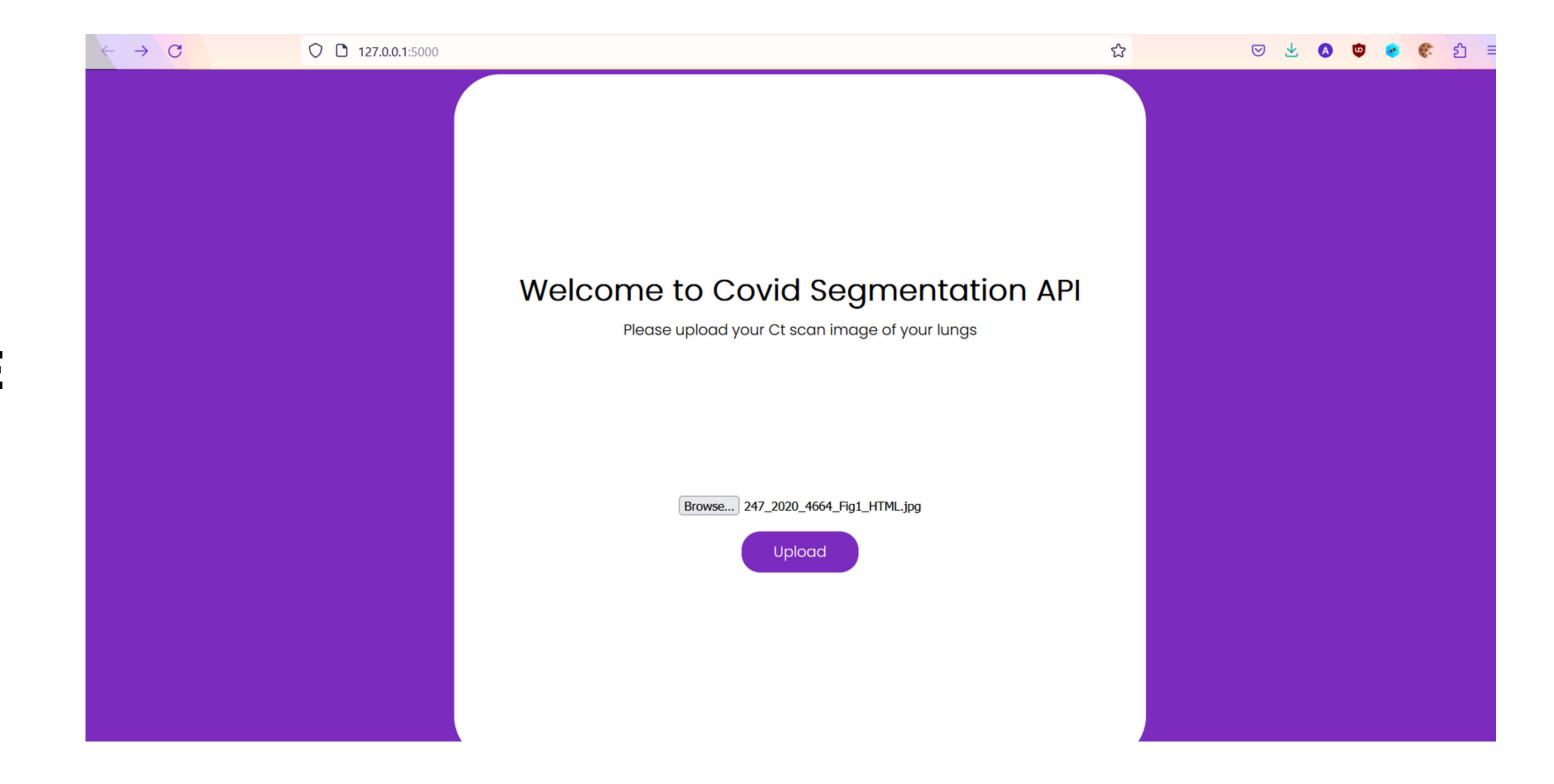


#### **SEGMENTATION**

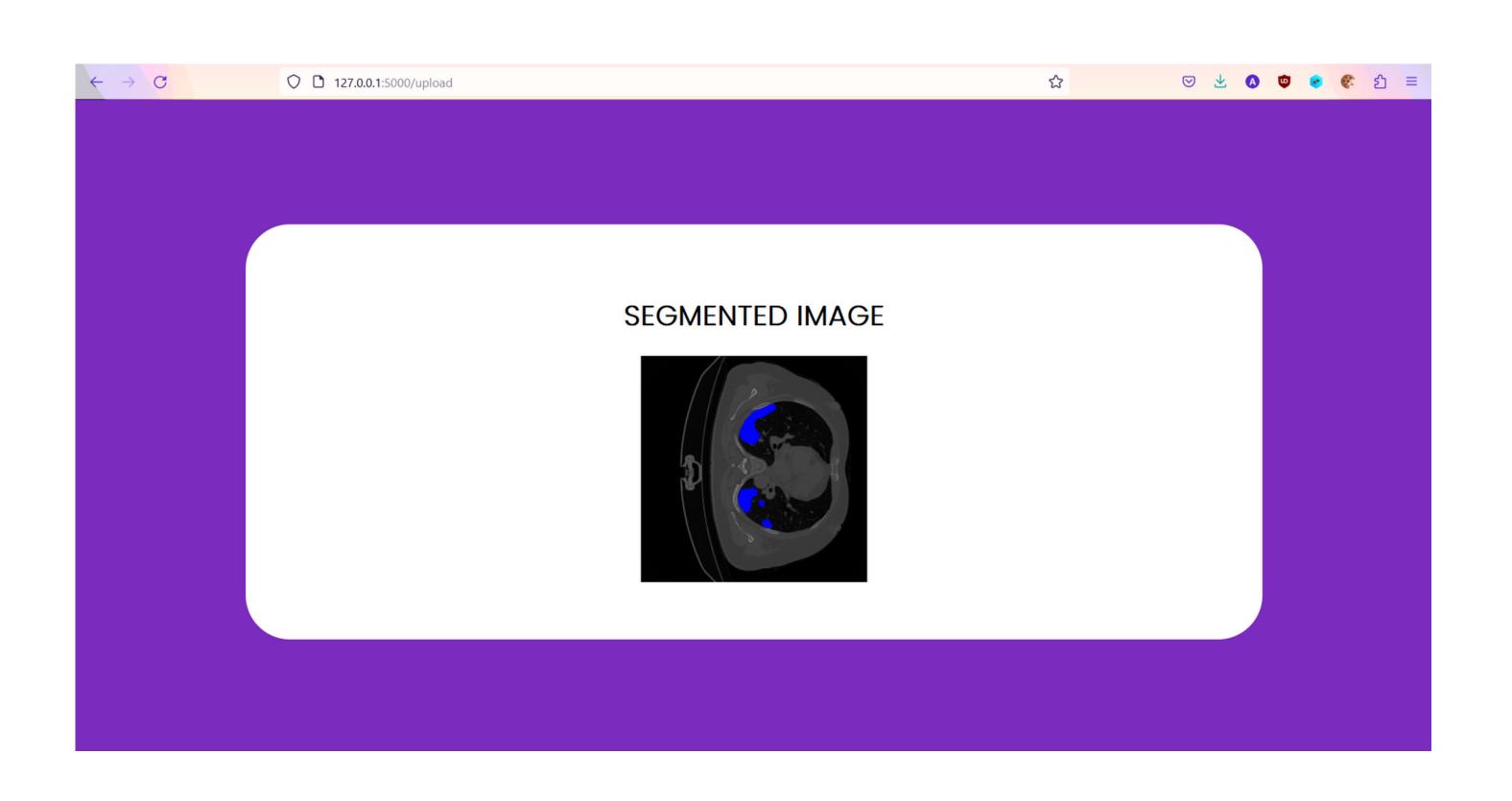
```
"" Predicting the mask """
for x, y in tqdm(zip(test_x, test_y), total=len(test_x)):
    """ Extracing the image name. """
   image_name = x.split("/")[-1]
   """ Reading the image """
   ori_x = cv2.imread(x, cv2.IMREAD_COLOR)
   ori_x = cv2.resize(ori_x, (W, H))
   x = ori_x/255.0
   x = x.astype(np.float32)
   x = np.expand_dims(x, axis=0)
    """ Reading the mask """
   ori_y = cv2.imread(y1, cv2.IMREAD_GRAYSCALE)
   # ori_y2 = cv2.imread(y2, cv2.IMREAD_GRAYSCALE)
   # ori_y = ori_y1 + ori_y2
   ori_y = cv2.resize(ori_y, (W, H))
   ori_y = np.expand_dims(ori_y, axis=-1) ## (512, 512, 1)
   ori_y = np.concatenate([ori_y, ori_y, ori_y], axis=-1) ## (512, 512, 3)
   """ Predicting the mask. """
   y_pred = model.predict(x)[0] > 0.5
   y_pred = y_pred.astype(np.int32)
   """ Saving the predicted mask along with the image and GT """
   save_image_path = f"results/{image_name}"
   y_pred = np.concatenate([y_pred, y_pred, y_pred], axis=-1)
   sep_line = np.ones((H, 10, 3)) * 255
   cat_image = np.concatenate([ori_x, sep_line, ori_y, sep_line, y_pred*255], axis=1)
    cv2.imwrite(save_image_path, cat_image)
```

# API

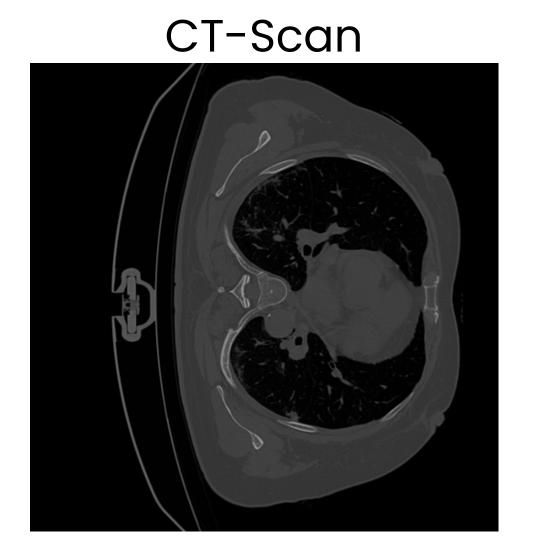
```
@app.route('/')
def index():
    return render_template('h.html')
@app.route('/upload', methods=['GET', 'POST'])
def upload file():
    if request.method == 'POST':
        uploaded_file = request.files['file']
        temp_file_path='D:\\downloads\\flsk\\static\\input.png'
        uploaded file.save(temp file path)
        output= process_image(temp_file_path)
        return render_template('result.html',output='output.png')
app.run()
```

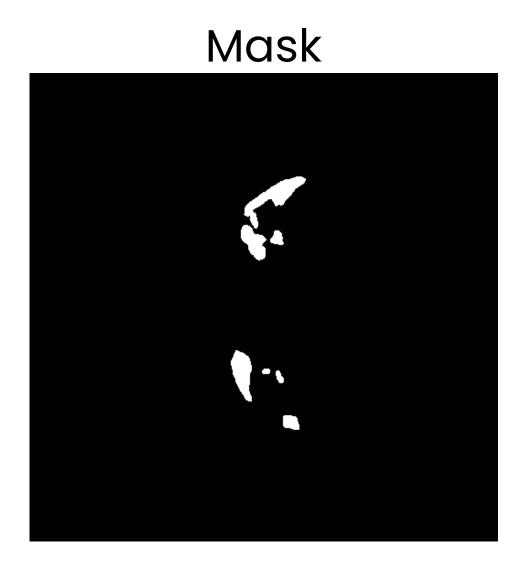


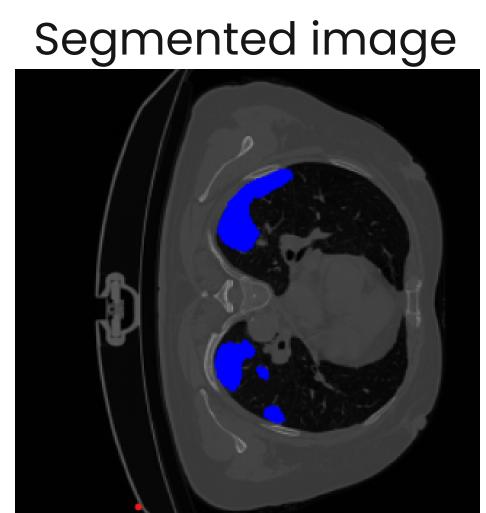
# **INTERFACE**



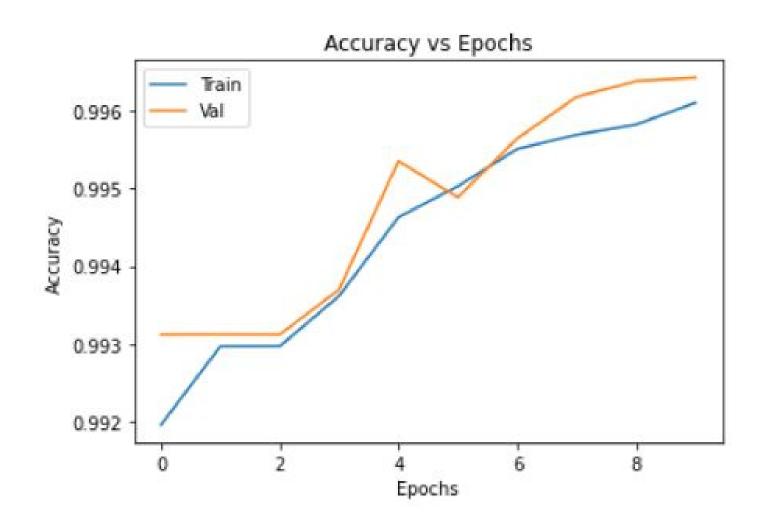
# Sample Test Data

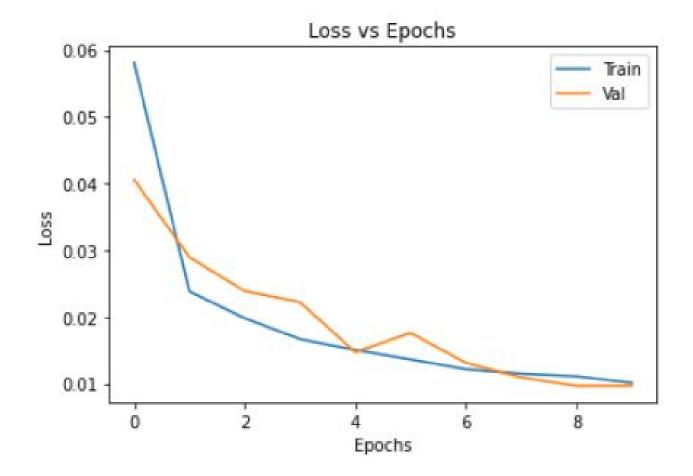






# **PERFORMANCE**





- No overfitting is observed as the graphs align with each other and rise together.
- The training accuracy curve converges at 97% at 10 epochs.
- The validation accuracy reaches 95% at 10 epochs.

# CONCLUSION

- The 'Covid-19 Segmentation API' utilizes the U-net model to accurately segment COVID-19 infections.
- It provides a user-friendly interface for researchers and medical professionals to input COVID-19 images and obtain segmented results efficiently.
- It has the potential to be extended for segmenting other types of infections, making it valuable for medical research and diagnosis