

Plant_Pathology Dataset



- Given a photo of an apple leaf, can you accurately assess its health? This competition will challenge you to distinguish between leaves which are healthy, those which are infected with apple rust, those that have apple scab, and those with more than one disease.
- A folder containing the train and test images, in jpg format.

1-PreProcessing

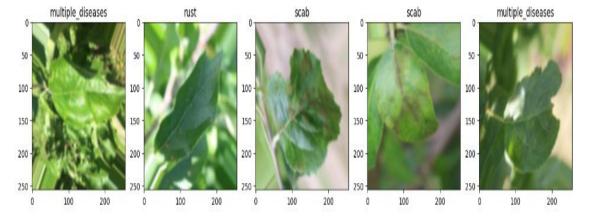
Read Data

Determine Directions

```
In [ ]: from shutil import copyfile
        # deLete temp dir
        if os.path.exists('/kaggle/temp/'):
            shutil.rmtree('/kaggle/temp/')
        os.mkdir('/kaggle/temp/')
        # train directory
        os.mkdir('/kaggle/temp/train')
        os.mkdir('/kaggle/temp/train/healthy')
        os.mkdir('/kaggle/temp/train/multiple_diseases')
        os.mkdir('/kaggle/temp/train/rust')
        os.mkdir('/kaggle/temp/train/scab')
        # validation directory
        os.mkdir('/kaggle/temp/valid')
        os.mkdir('/kaggle/temp/valid/healthy')
        os.mkdir('/kaggle/temp/valid/multiple_diseases')
        os.mkdir('/kaggle/temp/valid/rust')
        os.mkdir('/kaggle/temp/valid/scab')
In [ ]: SOURCE = '/kaggle/input/plant-pathology-2020-fgvc7/images/'
        TRAIN_DIR = '/kaggle/temp/train/'
        # copy images to train directory
        for index, data in train_set.iterrows():
            label = df.columns[np.argmax(data)]
            filepath = os.path.join(SOURCE, index + ".jpg")
            destination = os.path.join(TRAIN_DIR, label, index + ".jpg")
            copyfile(filepath, destination)
        for subdir in os.listdir(TRAIN_DIR):
            print(subdir, len(os.listdir(os.path.join(TRAIN_DIR, subdir))))
        healthy 416
        scab 465
        multiple_diseases 73
        rust 502
In [ ]: VALID_DIR = '/kaggle/temp/valid/'
        # copy images to valid directory
        for index, data in valid_set.iterrows():
            label = df.columns[np.argmax(data)]
            filepath = os.path.join(SOURCE, index + ".jpg")
            destination = os.path.join(VALID_DIR, label, index + ".jpg")
            copyfile(filepath, destination)
        for subdir in os.listdir(VALID_DIR):
            print(subdir, len(os.listdir(os.path.join(VALID_DIR, subdir))))
        healthy 100
        scab 127
        multiple_diseases 18
        rust 120
In [ ]: healthy_dir = os.path.join(TRAIN_DIR, 'healthy')
        mdiseases_dir = os.path.join(TRAIN_DIR, 'multiple_diseases')
        scab_dir = os.path.join(TRAIN_DIR, 'scab')
        rust_dir = os.path.join(TRAIN_DIR, 'rust')
        healthy_files = os.listdir(healthy_dir)
        mdiseases_files = os.listdir(mdiseases_dir)
        scab_files = os.listdir(scab_dir)
        rust files = os.listdir(rust dir)
```

Data Visualization

```
In [ ]: %matplotlib inline
         import matplotlib.pyplot as plt
        import matplotlib.image as mpimg
        pic index = 2
         next_healthy = [os.path.join(healthy_dir, fname) for fname in healthy_files[pic_index-2:pic_index]]
         next_mdiseases = [os.path.join(mdiseases_dir, fname) for fname in mdiseases_files[pic_index-2:pic_index]]
         next_scab = [os.path.join(scab_dir, fname) for fname in scab_files[pic_index-2:pic_index]]
         next_rust = [os.path.join(rust_dir, fname) for fname in rust_files[pic_index-2:pic_index]]
         nrows = 4
         ncols = 4
        fig = plt.gcf()
        fig.set_size_inches(ncols*4, nrows*4)
        for i, img_path in enumerate(next_healthy+next_mdiseases+next_scab+next_rust):
            sp = plt.subplot(nrows, ncols, i + 1)
            sp.axis('Off') # Don't show axes (or gridlines)
            img = mpimg.imread(img_path)
            plt.title(img_path.split('/')[-2])
            plt.imshow(img)
        plt.show()
                    healthy
                                                                                 multiple diseases
                                                                                                                  multiple_diseases
```



Data Balance (Over Sampling)

```
: target_multi_cols = ['healthy', 'multiple_diseases', 'rust', 'scab']
  print("Multi Classification Targets")
 print(train_df[target_multi_cols].sum())
  Multi Classification Targets
  healthy
                      516
  multiple diseases
                       91
                      622
  scab
 dtype: int64
  Maka data balance (over sampling)
: def balance set(df, x cols, y cols):
      ros = RandomOverSampler(random state=42)
     x_multi, y_multi = ros.fit_resample(df[x_cols], df[y_cols].values)
      data = pd.concat([x_multi, pd.DataFrame(y_multi, columns= y_cols)], axis=1)
      return data
  train_multi = balance_set(train_df,
                           x_cols = ["image_id", "img_name"],
                           y_cols = target_multi_cols)
  labels = train_multi[target_multi_cols]
  label_names = labels[labels==1].stack().reset_index()['level_1']
  label_names.index = train_multi.index
  train multi['label names'] = label names
  print("Multi Classification Labels")
  print(train_multi[target_multi_cols].sum())
  Multi Classification Labels
  healthy
                      622
  multiple_diseases
                      622
  rust
                      622
  scab
                       622
  dtype: int64
```

Data Split (Train&Validation)

```
from sklearn.model_selection import train_test_split

train_set, valid_set = train_test_split(df, test_size=0.2, random_state=42)

print(train_set.shape)
print(valid_set.shape)

(1456, 4)
(365, 4)
```

Image Generator

Before Balance

After Balance

```
import tensorflow as tf
from tensorflow import keras
import tensorflow datasets as tfds
import os
import cv2
from sklearn.model selection import train test split
import numpy as no
import matplotlib.pyplot as plt
import tensorflow.keras.layers as tfl
import pandas as pd
from tensorflow.keras.models import Model
from tensorflow.keras.preprocessing import image_dataset_from_directory
from tensorflow.keras.layers.experimental.preprocessing import RandomFlip, RandomRotation
from PIL import Image
import csv
from tensorflow.keras.preprocessing import image
from tensorflow.keras.preprocessing.image import ImageDataGenerator
training datagen = ImageDataGenerator(rescale = 1./255,
                                     rotation range=40,
                                     width shift range=0.2.
                                     height shift range=0.2,
                                      shear range=0.2,
                                     zoom_range=0.2,
                                     horizontal_flip=True,
                                     fill mode='nearest')
validation_datagen = ImageDataGenerator(rescale = 1./255)
#test_datagen = ImageDataGenerator( rescale = 1.0/255. )
train generator = training datagen.flow from directory(TRAIN DIR, target size=(150,150), class mode='categorical', batch size=32
validation generator = validation datagen.flow from directory(VALID DIR, target size=(150,150), class mode='categorical', batch
```

```
def blur preprocessing(img):
    return cv2.blur(img, (5, 5))
tf.random.set seed(99)
img data generator = ImageDataGenerator(rescale=1/255,
                                        validation split=0.2,
                                        rotation range = 180,
                                        horizontal flip = True,
                                        vertical flip = True,
                                        preprocessing function=blur preprocessing
train data multi = img data generator.flow from dataframe(dataframe=train multi,
                                                    directory="/kaggle/input/plant-pathology-2020-fgvc7/images/",
                                                    x col="img name",
                                                   y col= "label names",
                                                    target_size=(256, 256),
                                                    class_mode='categorical',
                                                    batch size=32,
                                                    subset='training',
                                                    shuffle=True,
                                                    seed=42)
val data multi = img_data_generator.flow_from_dataframe(dataframe=train_multi,
                                                    directory="/kaggle/input/plant-pathology-2020-fgvc7/images/",
                                                    x col="img name",
                                                    y_col="label_names",
                                                    target size=(256, 256),
                                                    class mode='categorical',
                                                    batch size=32,
                                                    subset='validation',
                                                    shuffle=True,
                                                    seed=42)
```

Found 1456 images belonging to 4 classes. Found 365 images belonging to 4 classes.

Found 1991 validated image filenames belonging to 4 classes. Found 497 validated image filenames belonging to 4 classes.

Blur preprocessing

Learning rate scheduler

```
def blur_preprocessing(img):
    return cv2.blur(img, (5, 5))
```

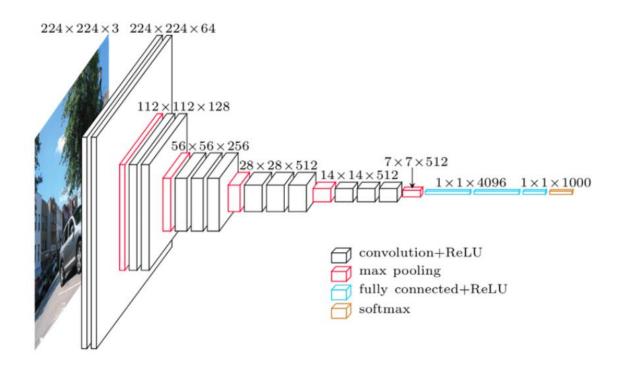
2-Image Classification Models

1- VGG16 (Pretrained Model)

What Is VGG16?

- ➤ VGG16 refers to the VGG model, also called VGGNet. It is a convolution neural network (CNN) model supporting 16 layers.
- ➤ The VGG16 model can achieve a test accuracy of 92.7% in ImageNet, a dataset containing more than 14 million training images across 1000 object classes. It is one of the top models from the ILSVRC-2014 competition.

VGG16 Architecture



VGG16 modeling

(Transfer learning – compile - fit – load model - evaluation)

```
from tensorflow.keras.applications.vgg16 import VGG16
base_model = VGG16(input_shape = (150, 150, 3), # Shape of our images
include_top = False, # Leave out the last fully connected layer
weights = 'imagenet')
```

```
import tensorflow as tf
import keras
from keras import layers
for layer in base_model.layers:
    layer.trainable = False
    tf.keras.layers.BatchNormalization()
```

```
# Flatten the output layer to 1 dimension
x = layers.Flatten()(base_model.output)
# Add a fully connected layer with 512 hidden units and ReLU activation
x = layers.Dense(512, activation='relu')(x)
# Add a dropout rate of 0.5
x = layers.Dropout(0.5)(x)
# Add a final sigmoid layer with 1 node for classification output
x = layers.Dense(4, activation='softmax')(x)
model = tf.keras.models.Model(base_model.input, x)
model.compile(optimizer='Adam', loss='categorical_crossentropy',metrics = ['acc'])
early_stopping_cb = tf.keras.callbacks.EarlyStopping(patience=5)
checkpoint_cb = tf.keras.callbacks.ModelCheckpoint("vgg16.h5", save_best_only=True)
```

```
vgghist = model.fit(train_generator, validation_data = validation_generator, steps_per_epoch = 10, epochs = 15, validation_s
```

Visualize train and validation accuracy

```
import matplotlib.pyplot as plt
acc = vgghist.history['acc']
val_acc = vgghist.history['val_acc']
loss = vgghist.history['loss']
val_loss = vgghist.history['val_loss']
epochs = range(len(acc))
plt.plot(epochs, acc, 'r', label='Training accuracy')
plt.plot(epochs, val_acc, 'b', label='Validation accuracy')
plt.title('Training and validation accuracy')
plt.legend(loc=0)
plt.show()
```

Evatuation

```
model = tf.keras.models.load_model("vgg16.h5") # rollback to best model
model.evaluate(validation_generator)
```

VGG16 modeling

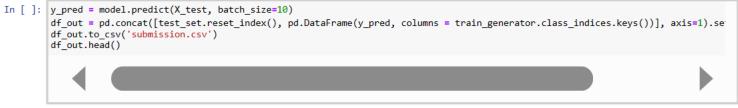
(read test data - make predictions)

```
In [ ]: test_set = pd.read_csv("/kaggle/input/plant-pathology-2020-fgvc7/test.csv", index_col=0)

X_test = []
    for index, data in test_set.iterrows():
            filepath = os.path.join(SOURCE, index + ".jpg")
            img = image.load_img(filepath, target_size=(150, 150))
            x = image.img_to_array(img)
            x = np.expand_dims(x, axis=0)
            X_test.append(x)

X_test = np.vstack(X_test) / 255 # rescale images
```

predict the test data



183/183 [========] - 7s 30ms/step

Out[19]:

mage_id				
Test_0	0.106944	0.044960	0.239254	0.608842
Test_1	0.199255	0.047042	0.460560	0.293144
Test_2	0.014253	0.024624	0.089598	0.871525
Test_3	0.552810	0.033190	0.152912	0.261087
Test_4	0.187862	0.061119	0.353388	0.397631

healthy multiple diseases

VGG16 modeling

(Predictions)

```
ncols = 4
fig = plt.gcf()
fig.set_size_inches(ncols*4, nrows*4)
for i, (idx, row) in enumerate(df_out.sample(nrows*ncols).iterrows()):
    filepath = filepath = os.path.join(SOURCE, idx + ".jpg")
    sp = plt.subplot(nrows, ncols, i + 1)
sp.axis('Off') # Don't show axes (or gridLines)
    img = mpimg.imread(filepath)--
    plt.title(df_out.columns[np.argmax(row)])
    plt.imshow(img)
```

2- Second Model (Build the model)

Second model (Created model)

```
model = tf.keras.models.Sequential([
    tf.keras.layers.Conv2D(64, (3,3), activation='relu', input_shape=(150, 150, 3)),
    tf.keras.layers.MaxPooling2D(2, 2),
    tf.keras.layers.MaxPooling2D(2,2),
    tf.keras.layers.Conv2D(128, (3,3), activation='relu'),
    tf.keras.layers.Conv2D(128, (3,3), activation='relu'),
    tf.keras.layers.MaxPooling2D(2,2),
    tf.keras.layers.Conv2D(128, (3,3), activation='relu'),
    tf.keras.layers.MaxPooling2D(2,2),
    tf.keras.layers.Flatten(),
    tf.keras.layers.Flatten(),
    tf.keras.layers.Dense(512, activation='relu'),
    tf.keras.layers.Dense(4, activation='relu')]
```

Model: "sequential 1"

1792 0 36928 0 73856
36928 0 73856
0 73856
73856
0
147584
0
0
0
3211776
2052

Second Model (compile - fit)

```
: model.compile(loss = 'categorical_crossentropy', optimizer='rmsprop', metrics=['accuracy'])
early stopping cb = tf.keras.callbacks.EarlyStopping(patience=5)
checkpoint_cb = tf.keras.callbacks.ModelCheckpoint("cm.h5", save_best_only=True)
history = model.fit(train generator, epochs=50, steps per epoch=46,
         validation_data = validation_generator, validation_steps=12, callbacks=[early_stopping_cb, checkpoint_cb]
Epoch 1/50
Epoch 2/50
8712
46/46 [=============] - 28s 606ms/step - loss: 0.5331 - accuracy: 0.8015 - val_loss: 0.5373 - val_accuracy: 0.
Epoch 4/50
7479
Epoch 5/50
46/46 [===========] - 27s 597ms/step - loss: 0.4785 - accuracy: 0.8310 - val_loss: 0.3099 - val_accuracy: 0.
7863
Enoch 7/50
8822
Epoch 8/50
8192
Epoch 9/50
8548
9288
Epoch 11/50
8795
46/46 [============] - 27s 592ms/step - loss: 0.4077 - accuracy: 0.8516 - val_loss: 0.2552 - val_accuracy: 0.
9151
```

46/46 [===========] - 27s 591ms/step - loss: 0.4016 - accuracy: 0.8654 - val_loss: 0.4763 - val_accuracy: 0.

46/46 [============] - 27s 587ms/step - loss: 0.3788 - accuracy: 0.8757 - val loss: 0.2590 - val accuracy: 0.

46/46 [===========] - 27s 595ms/step - loss: 0.3446 - accuracy: 0.8839 - val_loss: 0.2467 - val_accuracy: 0.

Epoch 13/50

8904 Epoch 14/50

8438 Epoch 15/50

9260 Epoch 16/50

9041 Epoch 17/50

8986

9178

8822 Epoch 20/50

9151

Second Model

(train and validation accuracy visualization – load model - evaluation)

```
In []: import matplotlib.pyplot as plt
    acc = history.history['accuracy']
    val_acc = history.history['val_accuracy']
    loss = history.history['loss']
    val_loss = history.history['val_loss']

epochs = range(len(acc))

plt.plot(epochs, acc, 'r', label='Training accuracy')
    plt.plot(epochs, val_acc, 'b', label='Validation accuracy')
    plt.title('Training and validation accuracy')
    plt.legend(loc=0)
```

Training and validation accuracy



```
In [ ]: model = tf.keras.models.load_model("cm.h5") # rollback to best model
model.evaluate(validation_generator)
```

12/12 [=======] - 4s 337ms/step - loss: 0.2368 - accuracy: 0.9260

Out[25]: [0.2368302196264267, 0.9260274171829224]

Second Model

(read test data - make predictions)

```
In [ ]: | test_set = pd.read_csv("/kaggle/input/plant-pathology-2020-fgvc7/test.csv", index_col=0)
         X_test = []
         for index, data in test_set.iterrows():
             filepath = os.path.join(SOURCE, index + ".jpg")
             img = image.load img(filepath, target size=(150, 150))
             x = image.img_to_array(img)
             x = np.expand_dims(x, axis=0)
             X_test.append(x)
         X_test = np.vstack(X_test) / 255 # rescale images
In [ ]: y_pred = model.predict(X_test, batch_size=10)
         df_out = pd.concat([test_set.reset_index(), pd.DataFrame(y_pred, columns = train_generator.class_indices.keys())], axis=1).se
         df out.to csv('submission.csv')
         df_out.head()
         183/183 [======] - 1s 4ms/step
Out[27]:
                       healthy multiple_diseases
                                                rust
                                                            scab
          image_id
            Test 0 6.722297e-12
                                     0.000572 0.999428 7.045307e-13
            Test_1 1.999887e-08
                                     0.015762 0.984238 1.652383e-08
            Test_2 4.754157e-02
                                     0.017981 0.000729 9.337487e-01
            Test_3 8.150271e-01
                                     0.023522 0.002990 1.584612e-01
            Test 4 1.529118e-18
                                     0.001206 0.998794 3.132368e-18
```

Second Model

(Predictions)

```
In [ ]: nrows = 4 ncols = 4
              fig = plt.gcf()
fig.set_size_inches(ncols*4, nrows*4)
             for i, (idx, row) in enumerate(df_out.sample(nrows*ncols).iterrows()):
    filepath = filepath = os.path.join(SOURCE, idx + ".jpg")
    sp = plt.subplot(nrows, ncols, i + 1)
                    sp.axis('Off') # Don't show axes (or gridlines)
                    img = mpimg.imread(filepath)
plt.title(df_out.columns[np.argmax(row)])
                    plt.imshow(img)
              plt.show()
```

A DenseNet is a type of convolutional neural network that utilises dense connections between layers, through Dense Blocks, where we connect all layers (with matching feature-map sizes) directly with each other.

3- Dense Net Pretrained Model

3- Dense Net

Pretrained Model

(Import dense model -Transfer learning – compile - fit)

```
from tensorflow.keras.applications import DenseNet201
def dense net model(trainable weights=False, weights path=None):
    tf.keras.backend.clear session()
   dense net = DenseNet201(input shape=(256, 256, 3), weights="imagenet", include top=False)
   for layer in dense net.layers:
       layer.trainable=trainable_weights
   model = tf.keras.models.Sequential([dense_net,
                                        tf.keras.layers.GlobalAveragePooling2D(),
                                        tf.keras.layers.Dense(128, activation='relu'),
                                        tf.keras.layers.Dropout(0.3),
                                        tf.keras.layers.Dense(4, activation='softmax')])
   if weights path:
       model.load_weights(weights_path)
    optimizer = tf.keras.optimizers.Adam(learning rate=learning rate scheduler)
   model.compile(loss="categorical_crossentropy", optimizer=optimizer, metrics=['accuracy'])
   return model
early_stopping cb = tf.keras.callbacks.EarlyStopping(patience=3)
checkpoint cb = tf.keras.callbacks.ModelCheckpoint("dense.h5", save best only=True)
dense net transfer = dense net model(trainable weights=True)
dense net transfer history = dense net transfer.fit(train data multi, validation data=val data multi, epochs=25, steps per epo
```

Pretrained Model

(Import dense model -Transfer learning – compile - fit)

```
]: def dense net model(trainable weights=False, weights path=None):
  tf.keras.backend.clear_session()
  dense_net = DenseNet281(input_shape=(256, 256, 3), weights="imagenet", include_top=False)
  for layer in dense net.layers:
   layer.trainable=trainable_weights
  model = tf.keras.models.Sequential([dense net.
             tf.keras.layers.GlobalAveragePooling2D(),
             tf.keras.layers.Dense(128, activation='relu'),
             tf.keras.layers.Dropout(0.3),
             tf.keras.layers.Dense(4, activation='softmax')])
   model.load_weights(weights_path)
  optimizer = tf.keras.optimizers.Adam(learning_rate=learning_rate_scheduler)
  model.compile(loss="categorical_crossentropy", optimizer=optimizer, metrics=['accuracy'])
 early stopping cb = tf.keras.callbacks.EarlyStopping(patience=3)
 checkpoint cb = tf.keras.callbacks.ModelCheckpoint("dense.h5", save_best_only=True)
 dense_net_transfer = dense_net_model(trainable_weights=True)
 dense net transfer history = dense net transfer.fit(train data multi, validation data=val data multi, epochs=25, steps per epo
 Epoch 1/25
 Epoch 2/25
     Epoch 3/25
 Epoch 5/25
 Epoch 8/25
 32/32 [----
     32/32 [----
 Enoch 11/25
      Epoch 12/25
      Epoch 14/25
 32/32 [-----
       Epoch 18/25
 Epoch 28/25
 Epoch 21/25
```

Pretrained Model

(train and validation accuracy visualization – load model - evaluation)

```
In []: import matplotlib.pyplot as plt
    acc = dense_net_transfer_history.history['accuracy']
    val_acc = dense_net_transfer_history.history['val_accuracy']
    loss = dense_net_transfer_history.history['loss']
    val_loss = dense_net_transfer_history.history['val_loss']
    epochs = range(len(acc))

plt.plot(epochs, acc, 'r', label='Training accuracy')
    plt.plot(epochs, val_acc, 'b', label='Validation accuracy')
    plt.title('Training and validation accuracy')
    plt.legend(loc=0)
```

Training and validation accuracy 1.0 0.9 0.8 0.7 0.6 0.5 0.4 0.0 2.5 5.0 7.5 10.0 12.5 15.0 17.5 20.0

```
In [ ]: #model = tf.keras.models.load_model("dense.h5") # rollback to best model
dense_net_transfer.evaluate(val_data_multi)
```

16/16 [======= 0.915 - accuracy: 0.9678

Out[18]: [0.09147188812494278, 0.9678068161010742]

Pretrained Model

(read test data – make predictions – Save submission file)

Found 1821 validated image filenames.

Out[20]:

 image_id
 healthy
 multiple_diseases
 rust
 scab

 0
 Test_0
 0.000111
 0.002933
 0.996945
 0.000011

 1
 Test_1
 0.000018
 0.001202
 0.998775
 0.000004

 2
 Test_2
 0.177084
 0.414771
 0.087757
 0.320388

 3
 Test_3
 0.999806
 0.000059
 0.00040
 0.000095

 4
 Test_4
 0.000008
 0.000232
 0.999758
 0.000002

```
In [ ]: submission.to_csv("submission_DenseNet.csv", index=False)
```

Pretrained Model

(read test data – make predictions – Save submission file)

Found 1821 validated image filenames.

```
In []: preds_df = pd.DataFrame(preds, columns=["healthy", "multiple_diseases", "rust", "scab"])
submission = pd.concat([test_df.image_id, preds_df], axis=1)
submission.head()
```

607/607 [=========] - 46s 67ms/step

Out[20]:

_	image_id	healthy	multiple_diseases	rust	scab
0	Test_0	0.000111	0.002933	0.996945	0.000011
1	Test_1	0.000018	0.001202	0.998775	0.000004
2	Test_2	0.177084	0.414771	0.087757	0.320388
3	Test_3	0.999806	0.000059	0.000040	0.000095
4	Test_4	0.000008	0.000232	0.999758	0.000002

```
In [ ]: submission.to_csv("submission_DenseNet.csv", index=False)
```

```
In [ ]: den_model = tf.keras.models.load_model("dense.h5") # rollback to best model
den_model.evaluate(val_data_multi)
```

16/16 [============] - 17s 826ms/step - loss: 0.0863 - accuracy: 0.9799

Out[23]: [0.08627106249332428, 0.9798792600631714]

Made by

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Thank You