

## SOURCE CODE:

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fruit.ipynb (due to limited page size the code vegetable.ipynb uploaded in
github)#!/usr/bin/env python

# coding:
utf-8# In[1]:

pwd

# In[2]: cd E:/IBM_MY_COURSE/Project/Dataset
PlantDisease/fruit-dataset/fruit-dataset

# # Apply ImageDataGenerator functionality to Train and Test set

# # Preprocessing # In[3]: from keras.preprocessing.image
import ImageDataGenerator train_datagen
=

ImageDataGenerator(rescale=1./255, shear_range=0.2, zoom_range=0.2, horizontal_
flip=True) test_datagen = ImageDataGenerator(rescale=1) #In[4]: pw
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# In[5]: x_train =
train_datagen.flow_from_directory('E:/IBM_MY_COURSE/Project/Dataset
Plant
Disease/fruit-dataset/fruitdataset/train',target_size=(128,128),batch_size=32,class_m
ode='categorical')

# In[6]:
x_test=test_datagen.flow_from_directory('E:/IBM_MY_COURSE/Project/ Dataset
et Plant Disease/fruit-dataset/fruit-
dataset/test',target_size=(128,128),
batch_size=32,class_mode='categorical') ## Import the models

# In[7]: from tensorflow.keras.models import Sequential from
tensorflow.keras.layers importDense,Convolution2D,MaxPool2D,Flatten

## Initializing the
models 10# In[8]:
model=Sequential() ##
Add CNN Layers

# In[9]:
model.add(Convolution2D(32,(3,3),input_shape=(128,128,3),activation= 'relu')) #

In[10]: x_train.class_indices

## Add Pooling layer#

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In[11]:model.add(MaxPool2D(pool_size=(2,2)

))# # Add Flatten layer # In[12]:

model.add(Flatten()) # # Add DenseLayer

#                               In[21]:                               model.add(Dense(40,
kernel_initializer='uniform',activation='relu'))                               model.add(Dense(20,
kernel_initializer='random_uniform',activation='relu'))

#                               #                               Add                               Output                               Layer                               #
In[24]:model.add(Dense(6,activation='softmax',kernel_initializer='random_uniform'))

#                               #                               Compile                               the                               model                               #
In[25]:model.compile(loss='categorical_crossentropy',optimizer='adam',
metrics=['accuracy' ]) # In[26]: len(x_train)

# In[27]:5384/32

# # Fit the Model

#                               In[28]:
model.fit_generator(x_train,steps_per_epoch=168,validation_data=x_test
,validation_steps=52,epochs=3)# #

Save the Model

# In[29]:

model.save("fruit.h5")#

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In[30]: ls
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# # Test the Model
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# In[32]: from keras.preprocessing import image from
tensorflow.keras.preprocessing.image import img_to_array from
tensorflow.keras.models import load_model import numpy as np
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# In[33]: model =
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load_model("fruit.h5") # # Test
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Apple_Healthy Class images
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# In[37]:img = image.load_img('E:/IBM_MY_COURSE/Project/Dataset
PlantDisease/fruitdataset/fruit-dataset/test/Apple_____
healthy/00fca0da-2db3-481b- b98a9b67bb7b105c
_____RS_H
L7708.JPG',target_size=(128,128)) 11
```

```
# In[39]: x=image.img_to_array(img) x=np.expand_dims(x,axis=0)#
```

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In[40]: pred =model.predict_classes(x)
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# In[41]: pred
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# In[45]: index
=['Apple__Black_rot','Apple__healthy','Corn_(maize)_____
Northern_Leaf_Blight','Corn_( maize)_____healthy','Peach_____
Bacterial_spot','Peach_____healthy']
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# In[46]: print('the given image belongs to=',index[pred[0]])
```

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# # Test Apple Black Rot class images # In[54]: img
=image.load_img('E:/IBM_MY_COURSE/Project/Dataset
PlantDisease/fruitdataset/fruit-dataset/test/Apple_____
Black_rot/0f3d45f4-e121-42cd- a5b6- be2f866a0574_____JR_FrgE.S
2870.JPG',target_size=(128,128))

# In[55]: x=image.img_to_array(img) x=np.expand_dims(x,axis=0)
pred =model.predict_classes(x) print('the given image
belogsto=',index[pred[0]])

# # Test Corn Northern leaf Blight classimages

# In[56]:img = image.load_img('E:/IBM_MY_COURSE/Project/Dataset
PlantDisease/fruitdataset/fruit-
dataset/test/Corn_(maize)_____Northern_Leaf_Blight/00a14441-7a62-
4034-bc40-b196aeab2785_____RS_NLB 3932.JPG',target_size=(128,128))

# In[57]: x=image.img_to_array(img) x=np.expand_dims(x,axis=0)
pred =model.predict_classes(x) print('the given image
belogsto=',index[pred[0]])

# # Test Corn Healthy class images # In[58]: img
=image.load_img('E:/IBM_MY_COURSE/Project/Dataset
Plant Disease/fruitdataset/fruit-dataset/test/Corn_(maize)_____
healthy/0a68ef5a-027c- 41ae-b227- 159dae77d3dd_____R.S_HL
7969 copy.jpg',target_size=(128,128))

# In[59]: x=image.img_to_array(img) x=np.expand_dims(x,axis=0)
pred =model.predict_classes(x) print('the given image belongs
to=',index[pred[0]]) # #Test Peach Bacterial spot class

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images # In[60]: img =
image.load_img('E:/IBM_MY_COURSE/Project/Dataset
PlantDisease/fruitdataset/fruit-dataset/test/Peach_____
Bacterial_spot/00ddc106-692e- 4c67-b2e8- 569c924caf49____
Rutg._Bact.S 1228.JPG',target_size=(128,128)) 12# In[61]:
x=image.img_to_array(img) x=np.expand_dims(x,axis=0) pred =
model.predict_classes(x) print('the given image
belongsto=',index[pred[0]])

# # Test Peach Healthy classimages

# In[62]:img = image.load_img('E:/IBM_MY_COURSE/Project/Dataset
PlantDisease/fruitdataset/fruit-dataset/test/Peach_____
healthy/1a07ce54-f4fd-41cf- b088- 144f6bf71859_____Rutg._HL
3543.JPG',target_size=(128,128))

# In[63]: x=image.img_to_array(img) x=np.expand_dims(x,axis=0)pred
=model.predict_classes(x) print('the given image
belongsto=',index[pred[0]])l

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