

```
In [2]: from tensorflow.keras.preprocessing.image import ImageDataGenerator

In [3]: train_datagen=ImageDataGenerator(rescale=1./255,zoom_range=0.2,horizontal_flip=True,vertical_flip=False)

In [4]: test_datagen=ImageDataGenerator(rescale=1./255)

In [5]: x_train=train_datagen.flow_from_directory(r"C:\Users\maris_q3mm6nk\Desktop\FILES\data_for_ibm\Fertilizers_Recommendation_System_For_Disease_Predicti
        class_mode='categorical',batch_size=24)

Found 11385 images belonging to 9 classes.

In [6]: x_test=test_datagen.flow_from_directory(r"C:\Users\maris_q3mm6nk\Desktop\FILES\data_for_ibm\Fertilizers_Recommendation_System_For_Disease_Prediction
        class_mode='categorical',batch_size=24)

Found 3416 images belonging to 9 classes.

In [7]: from tensorflow.keras.models import Sequential
        from tensorflow.keras.layers import Dense,Convolution2D,MaxPooling2D,Flatten

In [8]: model=Sequential()

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

In [10]: model.add(MaxPooling2D(pool_size=(2,2)))

In [11]: model.add(Flatten())
```

In [12]:

```
model.summary()
```

Model: "sequential"

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 126, 126, 32)	896
max_pooling2d (MaxPooling2D)	(None, 63, 63, 32)	0
flatten (Flatten)	(None, 127008)	0
Total params: 896		
Trainable params: 896		
Non-trainable params: 0		

In [13]:

```
model.add(Dense(300,activation='relu'))  
model.add(Dense(150,activation='relu'))
```

In [20]:

```
model.add(Dense(9,activation='softmax'))
```

In [21]:

```
model.compile(loss='categorical_crossentropy',optimizer='adam',metrics=['accuracy'])
```

In [22]:

```
len(x_train)
```

Out[22]: 475

In [23]:

```
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```

```
Out[23]: 51.563333333333336
```

```
In [24]: model.fit(x_train, steps_per_epoch=len(x_train), validation_data=x_test, validation_steps=len(x_test), epochs=10)
```

Epoch 1/10

475/475 [=====] - 237s 498ms/step - loss: 2.1787 - accuracy: 0.1331 - val_loss: 2.1362 - val_accuracy: 0.1953

Epoch 2/10

475/475 [=====] - 224s 470ms/step - loss: 2.1077 - accuracy: 0.1868 - val_loss: 2.1022 - val_accuracy: 0.1953

Epoch 3/10

475/475 [=====] - 242s 509ms/step - loss: 2.0872 - accuracy: 0.1868 - val_loss: 2.0911 - val_accuracy: 0.1953

Epoch 4/10

475/475 [=====] - 244s 514ms/step - loss: 2.0795 - accuracy: 0.1868 - val_loss: 2.0859 - val_accuracy: 0.1953

Epoch 5/10

475/475 [=====] - 249s 525ms/step - loss: 2.0761 - accuracy: 0.1868 - val_loss: 2.0846 - val_accuracy: 0.1953

Epoch 6/10

475/475 [=====] - 249s 525ms/step - loss: 2.0745 - accuracy: 0.1868 - val_loss: 2.0837 - val_accuracy: 0.1953

Epoch 7/10

475/475 [=====] - 250s 526ms/step - loss: 2.0738 - accuracy: 0.1868 - val_loss: 2.0830 - val_accuracy: 0.1953

Epoch 8/10

475/475 [=====] - 248s 521ms/step - loss: 2.0735 - accuracy: 0.1868 - val_loss: 2.0842 - val_accuracy: 0.1953

Epoch 9/10

475/475 [=====] - 221s 466ms/step - loss: 2.0734 - accuracy: 0.1868 - val_loss: 2.0844 - val_accuracy: 0.1953

Epoch 10/10

475/475 [=====] - 221s 465ms/step - loss: 2.0734 - accuracy: 0.1868 - val_loss: 2.0836 - val_accuracy: 0.1953

```
Out[24]:
```

```
In [25]: model.save('vegetabledata.h5')
```

```
In [26]: import numpy as np
from tensorflow.keras.models import load_model
from tensorflow.keras.preprocessing import image
```

```
In [27]: model=load_model('vegetabledata.h5')
```

```

[203., 201., 214.],
[221., 219., 232.],
[207., 205., 218.]],

[[148., 146., 160.],
[165., 163., 177.],
[152., 150., 164.],
...,
[176., 174., 187.],
[192., 190., 203.],
[189., 187., 200.]],

[[162., 160., 174.],
[155., 153., 167.],
[141., 139., 153.],
...,
[180., 178., 191.],
[190., 188., 201.],
[191., 189., 202.]]], dtype=float32)

```

```
In [47]: x=np.expand_dims(x,axis=0)
```

```
In [48]: x
```

```
Out[48]: array([[[[135., 131., 145.],
[134., 130., 144.],
[133., 129., 143.],
...,
[166., 164., 178.],
[188., 186., 200.],
[213., 211., 225.]],

[[141., 137., 151.],
[139., 135., 149.],
[128., 124., 138.],
...,

```

```
In [45]: x=image.img_to_array(img)
```

```
In [46]: x
```

```
Out[46]: array([[135., 131., 145.],
               [134., 130., 144.],
               [133., 129., 143.],
               ...,
               [166., 164., 178.],
               [188., 186., 200.],
               [213., 211., 225.]],

              [[141., 137., 151.],
               [139., 135., 149.],
               [128., 124., 138.],
               ...,
               [201., 199., 213.],
               [157., 155., 169.],
               [172., 170., 184.]],

              [[136., 132., 146.],
               [135., 131., 145.],
               [141., 137., 151.],
               ...,
               [166., 164., 178.],
               [169., 167., 181.],
               [165., 164., 178.]],

              ...,

              [[163., 161., 175.],
               [154., 152., 166.],
               [160., 158., 172.],
               ...,
               [203., 201., 214.],
               [221., 219., 232.]])
```

```

...
[201., 199., 213.],
[157., 155., 169.],
[172., 170., 184.]],

[[136., 132., 146.],
 [135., 131., 145.],
 [141., 137., 151.],
 ...,
 [166., 164., 178.],
 [169., 167., 181.],
 [166., 164., 178.]],

...,

[[163., 161., 175.],
 [154., 152., 166.],
 [160., 158., 172.],
 ...,
 [203., 201., 214.],
 [221., 219., 232.],
 [207., 205., 218.]],

[[148., 146., 160.],
 [165., 163., 177.],
 [152., 150., 164.],
 ...,
 [176., 174., 187.],
 [192., 190., 203.],
 [189., 187., 200.]],

[[162., 160., 174.],
 [155., 153., 167.],
 [141., 139., 153.],
 ...,
 [180., 178., 191.],
 [190., 188., 201.],
 [191., 189., 202.]]], dtype=float32)

```

```
In [33]: img=image.load_img(r"C:\Users\maris_q3mm6nk\Desktop\FILES\data_for_ibm\Fertilizers_Recommendation_System_For_Disease_Prediction\Dataset Plant Diseases
```

```
In [34]: img
```

```
Out[34]:
```



```
In [35]: x=image.img_to_array(img)
```

```
In [44]: img=image.load_img(r"C:\Users\maris_q3mm6nk\Desktop\FILES\data_for_ibm\Fertilizers_Recommendation_System_For_Disease_Prediction\Dataset Plant Diseases  
img
```

```
Out[44]:
```




```
In [49]: y=np.argmax(model.predict(x),axis=1)
```

```
1/1 [=====] - 0s 89ms/step
```

```
In [50]: x_train.class_indices
```

```
Out[50]: {'Pepper__bell__Bacterial_spot': 0,  
          'Pepper__bell__healthy': 1,  
          'Potato__Early_blight': 2,  
          'Potato__Late_blight': 3,  
          'Potato__healthy': 4,  
          'Tomato__Bacterial_spot': 5,  
          'Tomato__Late_blight': 6,  
          'Tomato__Leaf_Mold': 7,  
          'Tomato__Septoria_leaf_spot': 8}
```

```
In [51]: index=['Pepper__bell__Bacterial_spot','Pepper__bell__healthy','Potato__Early_blight','Potato__Late_blight','Potato__healthy','Tomato__Bacterial_
```

```
In [52]: index[y[0]]
```

```
Out[52]: 'Tomato__Bacterial_spot'
```

```
In [53]: img=image.load_img(r"C:\Users\maris_q3mm6nk\Desktop\FILES\data_for_ibm\Fertilizers_Recommendation_System_For_Disease_Prediction\Dataset Plant Diseases  
x=image.img_to_array(img)  
x=np.expand_dims(x,axis=0)  
y=np.argmax(model.predict(x),axis=1)  
index=['Pepper__bell__Bacterial_spot','Pepper__bell__healthy','Potato__Early_blight','Potato__Late_blight','Potato__healthy','Tomato__Bacterial_  
index[y[0]]
```

```
1/1 [=====] - 0s 38ms/step
```

```
Out[53]: 'Tomato__Bacterial_spot'
```