

```
In [1]: from tensorflow.keras.preprocessing.image import ImageDataGenerator
```

```
In [2]: train_datagen=ImageDataGenerator(rescale=1./255, zoom_range=0.2, horizontal_flip=True, vertical_flip=False)
```

```
In [3]: test_datagen=ImageDataGenerator(rescale=1./255)
```

```
In [6]: x_train=train_datagen.flow_from_directory('/content/drive/MyDrive/Classroom/Dataset Plant Disease/Veg-dataset/Veg-dataset/train_set',  
                                                class_mode='categorical', batch_size=24)
```

Found 10410 images belonging to 9 classes.

```
In [7]: x_test=test_datagen.flow_from_directory('/content/drive/MyDrive/Classroom/Dataset Plant Disease/fruit-dataset/fruit-dataset/train', class_mode='catego
```

Found 56 images belonging to 6 classes.

```
In [8]: from tensorflow.keras.models import Sequential  
from tensorflow.keras.layers import Dense, Convolution2D, MaxPooling2D, Flatten
```

```
In [9]: model=Sequential()
```

```
In [10]: model.add(Convolution2D(32,(3,3),input_shape=(128,128,3),activation='relu'))
```

```
In [11]: model.add(MaxPooling2D(pool_size=(2,2)))
```

```
In [12]: model.add(Flatten())
```

```
In [13]: 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 [14]: model.add(Dense(300,activation='relu'))
         model.add(Dense(150,activation='relu'))
```

```
In [15]: model.add(Dense(9,activation='softmax'))
```

```
In [16]: model.compile(loss='categorical_crossentropy',optimizer='adam',metrics=['accuracy'])
```

```
In [17]: len(x_train)
```

```
Out[17]: 434
```

```
In [18]: 1238/24
```

```
Out[18]: 51.583333333333336
```

```
In [21]: model.save('vegetabledata.h5')
```

```
In [22]: import numpy as np  
from tensorflow.keras.models import load_model  
from tensorflow.keras.preprocessing import image
```

```
In [23]: model=load_model('vegetabledata.h5')
```

```
In [25]: img=image.load_img('/content/drive/MyDrive/Classroom/Dataset Plant Disease/Veg-dataset/Veg-dataset/train_set/Pepper,_bell___healthy/00208a93-7687-4e8c')
```

```
In [26]: img
```

Out[26]:



```
In [27]: x=image.img_to_array(img)
```

```
In [27]: x=image.img_to_array(img)
```

```
In [29]: img=image.load_img('/content/drive/MyDrive/Classroom/Dataset Plant Disease/Veg-dataset/Veg-dataset/train_set/Pepper,_bell___healthy/01468dda-44f3-4de8  
img
```

Out[29]:



```
In [30]: x=image.img_to_array(img)
```

```
In [31]: x
```

```
Out[31]: array([[197., 190., 198.],  
               [195., 188., 196.],  
               [193., 186., 194.],  
               ...,  
               [208., 197., 203.],  
               [211., 200., 206.],  
               [212., 201., 207.]],  
              [[195., 188., 196.],  
               [191., 184., 192.],  
               [187., 180., 188.]])
```

```

[107., 100., 100.],
...,
[208., 197., 203.],
[208., 197., 203.],
[208., 197., 203.]],

[[199., 192., 200.],
[195., 188., 196.],
[190., 183., 191.],
...,
[212., 201., 207.],
[209., 198., 204.],
[209., 198., 204.]],

...,

[[183., 172., 178.],
[184., 173., 179.],
[186., 175., 181.],
...,
[139., 124., 129.],
[202., 187., 192.],
[146., 131., 136.]],

[[182., 171., 177.],
[183., 172., 178.],
[184., 173., 179.],
...,
[145., 130., 135.],
[167., 152., 157.],
[187., 172., 177.]],

[[191., 180., 186.],
[191., 180., 186.],
[191., 180., 186.],
...,
[211., 196., 201.],
[173., 158., 163.],
[160., 145., 150.]]], dtype=float32)

```

```
In [32]: x=np.expand_dims(x,axis=0)
```

```
In [33]: x
```

```

[[195., 188., 196.],
 [191., 184., 192.],
 [187., 180., 188.],
 ...,
 [208., 197., 203.],
 [208., 197., 203.],
 [208., 197., 203.]],

[[199., 192., 200.],
 [195., 188., 196.],
 [190., 183., 191.],
 ...,
 [212., 201., 207.],
 [209., 198., 204.],
 [209., 198., 204.]],

...,

[[183., 172., 178.],
 [184., 173., 179.],
 [186., 175., 181.],
 ...,
 [139., 124., 129.],
 [202., 187., 192.],
 [146., 131., 136.]],

[[182., 171., 177.],
 [183., 172., 178.],
 [184., 173., 179.],
 ...,
 [145., 130., 135.],
 [167., 152., 157.],
 [187., 172., 177.]],

[[191., 180., 186.],
 [191., 180., 186.],
 [191., 180., 186.],
 ...,
 [211., 196., 201.],
 [173., 158., 163.],
 [160., 145., 150.]]], dtype=float32)

```

```
In [41]: y=np.expand_dims(x,axis=0)
```

```
In [36]: x_train.class_indices
```

```
Out[36]: {'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 [37]: index=['Pepper_bell_Bacterial_spot','Pepper_bell_healthy','Potato_Early_blight','Potato_Late_blight']
```

```
In [42]: index[y[0]]
```

```
-----
TypeError                                Traceback (most recent call last)
in
----> 1 index[y[0]]

TypeError: only integer scalar arrays can be converted to a scalar index
```

```
In [40]: img=image.load_img('/content/drive/MyDrive/Classroom/Dataset Plant Disease/Veg-dataset/Veg-dataset/train_set/Pepper_bell_healthy/0119205b-cfac-4322
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_spot','Tomat
index[y[0]]
```