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!pip install google-api-python-client
!pip install oauth2client
!pip install oauth2client gspread google-api-python-client
import tensorflow as tf
print("TensorFlow version:", tf. version )
import gspread
from googleapiclient.discovery import build
from googleapiclient.http import MediaFileUpload
import os
import cv2
import numpy as np
from sklearn.model selection import train test split
from sklearn.preprocessing import LabelEncoder
import tensorflow as tf
from oauth2client.service account import ServiceAccountCredentials
# Path to the dataset
data dir =
'C:/Users/MALAYAPPAN/OneDrive/Desktop/all/aiot/computervision/plantDisease monitoring
/PlantVillage/PlantVillage'
# Parameters
img size = 128 # You can adjust this size
# Function to load and preprocess images
def load data(data dir, img size):
  images = []
  labels = []
  categories = os.listdir(data dir)
  for category in categories:
    category path = os.path.join(data dir, category)
    if not os.path.isdir(category path):
       continue
    for img in os.listdir(category_path):
       img_path = os.path.join(category_path, img)
       try:
          img_array = cv2.imread(img_path)
          if img array is None:
            print(f"Warning: Could not read image {img path}")
          resized_img = cv2.resize(img_array, (img_size, img_size))
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images.append(resized img)

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labels.append(category)
       except Exception as e:
          print(f"Error processing image {img_path}: {e}")
          continue
  return np.array(images), np.array(labels)
# Load data
images, labels = load_data(data_dir, img_size)
# Encode labels
le = LabelEncoder()
labels encoded = le.fit transform(labels)
# Split data
X_train, X_test, y_train, y_test = train_test_split(images, labels_encoded, test_size=0.2,
random state=42)
# Normalize images
X train = X train / 255.0
X_{test} = X_{test} / 255.0
print("Data loaded and preprocessed successfully.")
print(f"Number of training samples: {len(X train)}")
print(f"Number of testing samples: {len(X test)}")
import tensorflow as tf
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Conv2D, MaxPooling2D, Flatten, Dense, Dropout
from tensorflow.keras.optimizers import Adam
from tensorflow.keras.callbacks import EarlyStopping
# Build the model
model = Sequential([
  Conv2D(32, (3, 3), activation='relu', input shape=(img size, img size, 3)),
  MaxPooling2D((2, 2)),
  Conv2D(64, (3, 3), activation='relu'),
  MaxPooling2D((2, 2)),
  Conv2D(128, (3, 3), activation='relu'),
  MaxPooling2D((2, 2)),
  Flatten(),
  Dense(128, activation='relu'),
  Dropout(0.5),
  Dense(len(np.unique(labels encoded)), activation='softmax')
])
# Compile the model
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model.compile(optimizer=Adam(), loss='sparse categorical crossentropy',
metrics=['accuracy'])
# Print the model summary
model.summary()
# Early stopping to prevent overfitting
early stopping = EarlyStopping(monitor='val loss', patience=5, restore best weights=True)
# Train the model
history = model.fit(
  X_train, y_train,
  epochs=25,
  validation_data=(X_test, y_test),
  callbacks=[early stopping]
)
# Evaluate the model
test loss, test_acc = model.evaluate(X_test, y_test, verbose=2)
print(f"Test accuracy: {test_acc}")
# Save the model
model.save('plant disease model.h5')
def predict disease(image path, model):
  img = cv2.imread(image path)
  img = cv2.resize(img, (img size, img size))
  img = img / 255.0
  img = np.expand_dims(img, axis=0)
  prediction = model.predict(img)
  predicted class = np.argmax(prediction, axis=1)
  return le.inverse transform(predicted class)[0]
# Load the model
model = tf.keras.models.load model('plant disease model.h5')
# Predict the disease
image path =
r'C:\Users\MALAYAPPAN\OneDrive\Desktop\all\aiot\computervision\'Cercospora capsici.jpg'
predicted disease = predict disease(image path, model)
print(f"The predicted disease is: {predicted_disease}")
# Load the model
model = tf.keras.models.load model('plant disease model.h5')
# Predict the disease
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```
image path =
r'C:\Users\MALAYAPPAN\OneDrive\Desktop\all\aiot\computervision\WhatsApp Image
predicted disease = predict disease(image path, model)
print(f"The predicted disease is: {predicted disease}")
# Load the model
model = tf.keras.models.load model('plant disease model.h5')
# Predict the disease
image path = r'C:\Users\MALAYAPPAN\OneDrive\Pictures\banyan.jpg'
predicted disease = predict disease(image path, model)
print(f"The predicted disease is: {predicted disease}")
# Load the model
model = tf.keras.models.load model('plant disease model.h5')
# Predict the disease
image path = r'C:\Users\MALAYAPPAN\OneDrive\Pictures\bilwa.jpg'
predicted disease = predict disease(image path, model)
print(f"The predicted disease is: {predicted disease}")
def update google sheet(predicted disease, sheet name):
  # Define the scope and create credentials
  scope = ["https://spreadsheets.google.com/feeds",
"https://www.googleapis.com/auth/drive"]
  creds =
ServiceAccountCredentials.from_json_keyfile_name(r"C:\Users\MALAYAPPAN\plant-426909
-8c6454e8fafb.json", scope)
  client = gspread.authorize(creds)
  # Open the Google Sheet and select the first worksheet
  sheet = client.open(sheet name).sheet1
  # Append the predicted disease to the sheet
  sheet.append row([predicted disease])
# Update the Google Sheet with the predicted disease
sheet name = "live plant detection"
update google sheet(predicted disease, sheet name)
print("Google Sheet updated with the predicted disease.")
```