

SAMPLE CODE

Trained_Model.py

```
import os
import cv2
import numpy as np
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Conv2D, MaxPooling2D, Flatten, Dense, Dropout, Input
from tensorflow.keras.utils import to_categorical
from sklearn.model_selection import train_test_split

# Suppress TensorFlow logs and warnings
os.environ["TF_CPP_MIN_LOG_LEVEL"] = "3" # Suppress TensorFlow logs
os.environ["TF_ENABLE_ONEDNN_OPTS"] = "0" # Disable oneDNN warnings

def load_dataset(folder_path, image_size=(128, 128)):
    """
    Load images from folders where each folder represents a blood group.

    Args:
        folder_path (str): Path to the root dataset folder.
        image_size (tuple): Size to resize the images (width, height).

    Returns:
        images (numpy.ndarray): Array of preprocessed images.
        labels (numpy.ndarray): Corresponding labels for the images.
        label_map (dict): Mapping of blood group names to numeric labels.
    """
    images = []
    labels = []
    label_map = {}
    current_label = 0

    for folder_name in sorted(os.listdir(folder_path)):
        folder_full_path = os.path.join(folder_path, folder_name)
        if not os.path.isdir(folder_full_path):
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        continue
    label_map[folder_name] = current_label
    for file_name in os.listdir(folder_full_path):
        file_path = os.path.join(folder_full_path, file_name)
        img = cv2.imread(file_path, cv2.IMREAD_GRAYSCALE)
        if img is None:
            print(f"Warning: Skipping invalid image {file_path}")
            continue
        img = cv2.resize(img, image_size)
        images.append(img)
        labels.append(current_label)
    current_label += 1

if not images:
    raise ValueError("No images found in the dataset. Please check the folder structure.")

images = np.array(images, dtype=np.float32) / 255.0
labels = np.array(labels, dtype=np.int32)
return images, labels, label_map

def build_cnn_model(input_shape, num_classes):
    """
    Build and compile a CNN model.

    Args:
        input_shape (tuple): Shape of the input images (height, width, channels).
        num_classes (int): Number of output classes.

    Returns:
        model: Compiled CNN model.
    """
    model = Sequential([
        Input(shape=input_shape),
        Conv2D(32, (3, 3), activation='relu'),
        MaxPooling2D((2, 2)),
        Conv2D(64, (3, 3), activation='relu'),
        MaxPooling2D((2, 2)),
        Conv2D(128, (3, 3), activation='relu'),

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    MaxPooling2D((2, 2)),
    Flatten(),
    Dense(128, activation='relu'),
    Dropout(0.5),
    Dense(num_classes, activation='softmax')
])
model.compile(optimizer='adam', loss='categorical_crossentropy', metrics=['accuracy'])
return model

if __name__ == "__main__":
    # Path to your dataset folder
    dataset_path = "dataset" # Ensure this folder exists in the same directory as this script
    image_size = (128, 128)

    # Load the dataset
    print("Loading dataset...")
    images, labels, label_map = load_dataset(dataset_path, image_size)
    print(f'Loaded {len(images)} images.')
    print(f'Label Map: {label_map}')

    # Preprocess data
    images = images.reshape(-1, image_size[0], image_size[1], 1) # Reshape for grayscale
    labels = to_categorical(labels, num_classes=len(label_map))

    # Split data into training and testing sets
    X_train, X_test, y_train, y_test = train_test_split(images, labels, test_size=0.2, random_state=42)

    # Build and train the model
    print("Building and training the model...")
    model = build_cnn_model(input_shape=(image_size[0], image_size[1], 1), num_classes=len(label_map))
    history = model.fit(X_train, y_train, epochs=20, batch_size=32, validation_data=(X_test, y_test))

    # Save the trained model
    model.save("blood_group_model.h5")
    print("Model saved as blood_group_model.h5")

```

App.py

```
from flask import Flask, request, render_template
import tensorflow as tf
import cv2
import numpy as np
import os

app = Flask(__name__)
model = tf.keras.models.load_model("blood_group_model.h5") # Ensure this file is in the same folder
label_map = {0: "A+", 1: "A-", 2: "AB+", 3: "AB-", 4: "B+", 5: "B-", 6: "O+", 7: "O-"} # Adjust based on your dataset


UPLOAD_FOLDER = "static/uploads"
if not os.path.exists(UPLOAD_FOLDER):
    os.makedirs(UPLOAD_FOLDER)
app.config["UPLOAD_FOLDER"] = UPLOAD_FOLDER


@app.route("/")
def index():
    return render_template("index.html")


@app.route("/predict", methods=["POST"])
def predict():
    if "file" not in request.files:
        return "No file part", 400

    file = request.files["file"]
    if file.filename == "":
        return "No selected file", 400

    # Save the uploaded file
    file_path = os.path.join(app.config["UPLOAD_FOLDER"], file.filename)
    file.save(file_path)

    # Preprocess the image
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```
img = cv2.imread(file_path, cv2.IMREAD_GRAYSCALE)
img = cv2.resize(img, (128, 128)) / 255.0 # Normalize pixel values
img = img.reshape(1, 128, 128, 1)

# Predict blood group
predictions = model.predict(img)
predicted_label = label_map[np.argmax(predictions)]

return render_template("index.html", prediction=predicted_label, image_path=file_path)

if __name__ == "__main__":
    app.run(debug=True)
```

Index.html

```
<!DOCTYPE html>
<html lang="en">
<head>
  <meta charset="UTF-8">
  <meta name="viewport" content="width=device-width, initial-scale=1.0">
  <title>Blood Group Detection</title>
  <link rel="stylesheet" href="https://cdn.jsdelivr.net/npm/bootstrap@5.3.0-alpha1/dist/css/bootstrap.min.css">
  <style>
    body {
      background: linear-gradient(to bottom right, #003366, #ff6600); /* Dark blue to orange gradient */
      color: #fff;
      font-family: 'Arial', sans-serif;
      height: 100vh;
      display: flex;
      justify-content: center;
      align-items: center;
      margin: 0;
    }
    .container {
      width: 90%;
      max-width: 600px;
      padding: 20px;
      background-color: rgba(255, 255, 255, 0.15); /* Transparent white effect */
      border-radius: 10px;
      box-shadow: 0 4px 8px rgba(0, 0, 0, 0.3);
      text-align: center;
    }
    h1 {
      color: #ffcc00; /* Bright yellow-orange for the title */
    }
    p {
      color: #ffffff; /* Keep text readable */
    }
  </style>
</head>
<body>
  <div class="container">
    <h1>Blood Group Detection</h1>
    <p>Welcome to the Blood Group Detection System. Please enter your details to proceed.</p>
  </div>
</body>
</html>
```

```
.btn-primary {
  background-color: #ff6600; /* Vibrant orange button */
  border-color: #ff6600;
}

.btn-primary:hover {
  background-color: #cc5200; /* Darker orange on hover */
  border-color: #cc5200;
}

.form-label {
  color: #ffcc00; /* Bright yellow-orange for labels */
}

.result-box {
  background-color: rgba(0, 51, 102, 0.8); /* Semi-transparent dark blue */
  padding: 20px;
  border-radius: 10px;
  color: #ffcc00; /* Yellow text in result box */
  margin-top: 20px;
}

.result-text {
  font-size: 1.5rem;
  font-weight: bold;
  color: #ffcc00; /* Yellow for the result text */
}

.thumbnail {
  margin-top: 20px;
  border: 2px solid #ffcc00; /* Border matches yellow-orange theme */
  border-radius: 10px;
}
```

</style>

</head>

<body>

<div class="container">

<h1>Blood Group Detection</h1>

<p>Upload a fingerprint image to predict the blood group.</p>

<form action="/predict" method="POST" enctype="multipart/form-data" class="mt-4">

<div class="mb-3">

<label for="file" class="form-label">Choose Fingerprint Image:</label>

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        <input type="file" class="form-control" id="file" name="file" required>
    </div>

    <button type="submit" class="btn btn-primary">Predict</button>
</form>

{% if prediction %}
<div class="result-box">
    <h2>Prediction Result</h2>

    <p class="result-text">Blood Group: {{ prediction }}</p>

    
</div>

{% endif %}

</div>
</body>
</html>
```