

## 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):
```

```
    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'),

```

        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
```

```
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>
```

```

.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>

```

```
<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>
```