

face_recognize

April 22, 2025

1

```
[143]: import os
import numpy as np
from PIL import Image, ImageOps, ImageEnhance, ImageFilter
from sklearn.decomposition import PCA
from sklearn.svm import SVC
from sklearn.metrics import (
    confusion_matrix, accuracy_score, roc_curve, auc, f1_score,
    precision_score, recall_score, cohen_kappa_score
)
from sklearn.model_selection import GridSearchCV
import matplotlib.pyplot as plt
import seaborn as sns
```

1.1

```
[144]: #
train_dir = "split_data/train"
test_dir = "split_data/test"
test_out_dir = "split_data/test_out"
```

1.2

```
[145]: #
def preprocess_image(image_path, target_size=(64, 64), is_train=False):
    img = Image.open(image_path).convert("L") #
    width, height = img.size

    if is_train:
        # 1.
        scale = np.random.uniform(0.8, 1.2) #
        new_size = (int(target_size[0] * scale), int(target_size[1] * scale))
        img = img.resize(new_size)

        # 2.
        x = np.random.randint(0, new_size[0] - target_size[0] + 1)
        y = np.random.randint(0, new_size[1] - target_size[1] + 1)
```

```

img = img.crop((x, y, x + target_size[0], y + target_size[1]))

# 3.
if np.random.random() > 0.5:
    img = img.rotate(np.random.randint(-15, 15)) #
if np.random.random() > 0.5:
    img = img.transpose(Image.FLIP_LEFT_RIGHT)
enhancer = ImageEnhance.Brightness(img)
img = enhancer.enhance(np.random.uniform(0.7, 1.3)) #
enhancer = ImageEnhance.Contrast(img)
img = enhancer.enhance(np.random.uniform(0.8, 1.2)) #
if np.random.random() > 0.3: #
    img = img.filter(ImageFilter.GaussianBlur(radius=np.random.
↪uniform(0, 1)))

else:
    #
    x = (width - target_size[0]) // 2
    y = (height - target_size[1]) // 2
    img = img.crop((x, y, x + target_size[0], y + target_size[1]))

#
img = ImageOps.equalize(img)
img = np.array(img) / 255.0
return img.flatten() #

```

1.3

```

[146]: #
#
def load_dataset(data_dir, target_size=(64, 64), max_images_per_person=None):
    # print(f"    {data_dir}    ...")
    data = []
    labels = []
    filenames = [] #
    for subject_id in os.listdir(data_dir):
        subject_path = os.path.join(data_dir, subject_id)
        if os.path.isdir(subject_path):
            images = os.listdir(subject_path)
            if max_images_per_person:
                images = images[:max_images_per_person]
            for img_name in images:
                img_path = os.path.join(subject_path, img_name)
                img = preprocess_image(img_path, target_size)
                data.append(img)
                labels.append(subject_id)
                filenames.append(f"{subject_id}/{img_name}") #

```

```

print(f"{data_dir}      ")
return np.array(data), np.array(labels), filenames

```

1.4 PCA

```

[147]: # PCA
def pca_reduction(train_data, test_data, test_out_data, n_components):
    print(f"    PCA    {n_components}    ...")
    train_data_flat = train_data.reshape(train_data.shape[0], -1)
    test_data_flat = test_data.reshape(test_data.shape[0], -1)
    test_out_data_flat = test_out_data.reshape(test_out_data.shape[0], -1)

    mean_face = np.mean(train_data_flat, axis=0)

    train_data_centered = train_data_flat - mean_face
    test_data_centered = test_data_flat - mean_face
    test_out_data_centered = test_out_data_flat - mean_face

    #    n_components
    max_components = min(train_data_centered.shape[0], train_data_centered.
↪shape[1])
    if n_components > max_components:
        print(f" : n_components={n_components}    {max_components}")
        n_components = max_components

    pca = PCA(n_components=n_components)
    train_data_pca = pca.fit_transform(train_data_centered)
    test_data_pca = pca.transform(test_data_centered)
    test_out_data_pca = pca.transform(test_out_data_centered)
    print("PCA    ")
    return train_data_pca, test_data_pca, test_out_data_pca

```

1.5 SVM

```

[148]: from sklearn.model_selection import StratifiedKFold

def train_svm_classifiers(train_data, train_labels):
    print("    SVM    ...")
    unique_labels = np.unique(train_labels)
    classifiers = {}
    # for label in unique_labels:
    #     binary_labels = (train_labels == label).astype(int)
    #     param_grid = {'C': [0.1, 1, 10], 'gamma': [0.001, 0.01], 'kernel': '
↪[rbf]'}
    #     #
    #     cv = StratifiedKFold(n_splits=3)
    #     grid_search = GridSearchCV(SVC(probability=True), param_grid, cv=cv)

```

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#     grid_search.fit(train_data, binary_labels)
#     best_sum = grid_search.best_estimator_
#     classifiers[label] = best_sum
for label in unique_labels:
    binary_labels = (train_labels == label).astype(int)
    #
    svm = SVC(probability=True, C=0.01, gamma=0.001, kernel='rbf')
    svm.fit(train_data, binary_labels)
    classifiers[label] = svm
print("SVM      ")
return classifiers

```

1.6

```

[149]: #
def vote_predict(classifiers, data, threshold):
    # print("      ...")
    predictions = []
    all_scores = []
    for sample in data:
        scores = []
        for label, classifier in classifiers.items():
            score = classifier.predict_proba([sample])[0][1]
            scores.append(score)
        max_score = np.max(scores)
        all_scores.append(max_score)
        if max_score < threshold:
            predictions.append("OUT")
        else:
            predictions.append("IN")
    # print("      ")
    return np.array(predictions), np.array(all_scores)

```

1.7

```

[150]: #
def predict_person(classifiers, data):
    predictions = []
    for sample in data:
        scores = []
        labels = []
        for label, classifier in classifiers.items():
            score = classifier.predict_proba([sample])[0][1]
            scores.append(score)
            labels.append(label)
        predicted_label = labels[np.argmax(scores)]
        predictions.append(predicted_label)

```

```
return np.array(predictions)
```

1.8

```
[151]: # 10
def plot_mean_and_top_eigenfaces(train_data, n_components=50, top_n=10):
    print(" 10 ...")
    train_data_flat = train_data.reshape(train_data.shape[0], -1)
    mean_face = np.mean(train_data_flat, axis=0)

    # PCA
    pca = PCA(n_components=n_components)
    pca.fit(train_data_flat - mean_face)
    eigenfaces = pca.components_.reshape((n_components, 64, 64))
    explained_variance_ratio = pca.explained_variance_ratio_

    #
    plt.figure(figsize=(12, 6))
    plt.subplot(1, top_n + 1, 1)
    plt.imshow(mean_face.reshape(64, 64), cmap='gray')
    plt.title("Mean Face")
    plt.axis("off")

    # 10
    for i in range(top_n):
        plt.subplot(1, top_n + 1, i + 2)
        plt.imshow(eigenfaces[i], cmap='gray')
        plt.title(f"Eigenface {i + 1}\n({explained_variance_ratio[i]:.2%})")
        plt.axis("off")
    plt.tight_layout()
    plt.show()
    print(" ")

# Excel =50
import pandas as pd

def save_pca_to_excel_with_filenames(train_data, filenames, n_components=50,
    output_file="pca_results.xlsx"):
    print(" PCA Excel ...")
    train_data_flat = train_data.reshape(train_data.shape[0], -1)
    mean_face = np.mean(train_data_flat, axis=0)

    # PCA
    pca = PCA(n_components=n_components)
    train_data_pca = pca.fit_transform(train_data_flat - mean_face)
    explained_variance_ratio = pca.explained_variance_ratio_

```

```

#
df_pca = pd.DataFrame(train_data_pca, columns=[f"PC{i + 1}" for i in
↪range(n_components)])
df_pca.insert(0, "Filename", filenames) #
df_pca.to_excel(output_file, index=False, sheet_name="PCA Results")

#
df_variance = pd.DataFrame({
    "Principal Component": [f"PC{i + 1}" for i in range(n_components)],
    "Explained Variance (%)": explained_variance_ratio * 100
})
with pd.ExcelWriter(output_file, mode="a", engine="openpyxl") as writer:
    df_variance.to_excel(writer, index=False, sheet_name="Explained
↪Variance")

print(f"PCA      {output_file} ")
#
n = 6 #
train_data, train_labels, filenames = load_dataset(train_dir,
↪max_images_per_person=n)

#      10
plot_mean_and_top_eigenfaces(train_data, n_components=40, top_n=10)

#      PCA      Excel
save_pca_to_excel_with_filenames(train_data, filenames, n_components=40,
↪output_file="pca_results.xlsx")

```

split_data/train

10 ...



PCA Excel ...
PCA pca_results.xlsx

1.9

```
[155]: #
n = 6 #
train_data, train_labels, filenames = load_dataset(train_dir,
    ↪max_images_per_person=n)
test_data, test_labels, filenames = load_dataset(test_dir,
    ↪max_images_per_person=n)
test_out_data, test_out_labels, filenames = load_dataset(test_out_dir,
    ↪max_images_per_person=n)

print(" ")
print(f" : {train_data.shape}, : {test_data.shape}, : {test_out_data.
    ↪shape}")

# PCA
n_components = 50
train_data_pca, test_data_pca, test_out_data_pca = pca_reduction(train_data,
    ↪test_data, test_out_data, n_components)

#
classifiers = train_svm_classifiers(train_data_pca, train_labels)

#
# print(" ...")
all_predictions, all_scores = vote_predict(classifiers, np.
    ↪concatenate([test_data_pca, test_out_data_pca]), 0.5)
true_labels = np.concatenate([np.ones(len(test_labels)), np.
    ↪zeros(len(test_out_labels))])
fpr, tpr, thresholds = roc_curve(true_labels, all_scores)
roc_auc = auc(fpr, tpr)
optimal_threshold = thresholds[np.argmax(tpr - fpr)]
print(f" : {optimal_threshold:.3f}")

#
test_predictions, _ = vote_predict(classifiers, test_data_pca,
    ↪optimal_threshold)
test_out_predictions, _ = vote_predict(classifiers, test_out_data_pca,
    ↪optimal_threshold)
all_predictions = np.concatenate([test_predictions, test_out_predictions])
all_true_labels = np.concatenate([["IN"] * len(test_labels), ["OUT"] *
    ↪len(test_out_labels)])

# =====
#
cm = confusion_matrix(all_true_labels, all_predictions, labels=["IN", "OUT"])
tp, fp, fn, tn = cm[0,0], cm[1,0], cm[0,1], cm[1,1]
```

```

#
in_out_accuracy = accuracy_score(all_true_labels, all_predictions)
in_out_precision = precision_score(all_true_labels, all_predictions,
    pos_label="IN")
in_out_recall = recall_score(all_true_labels, all_predictions, pos_label="IN")
in_out_f1 = f1_score(all_true_labels, all_predictions, pos_label="IN")
in_out_specificity = tn / (tn + fp) if (tn + fp) != 0 else 0.0
in_out_kappa = cohen_kappa_score(all_true_labels, all_predictions)

#
print("\n====      =====")
print(f"      : {in_out_accuracy:.2f}")
print(f"      : {in_out_precision:.2f}")
print(f"      : {in_out_recall:.2f}")
print(f"      : {in_out_specificity:.2f}")
print(f"F1      : {in_out_f1:.2f}")
print(f"Kappa : {in_out_kappa:.2f}")
print(f"AUC : {roc_auc:.2f}")

#
plt.figure(figsize=(10, 8))
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues',
            xticklabels=["IN", "OUT"], yticklabels=["IN", "OUT"])
plt.xlabel('Predicted Labels')
plt.ylabel('True Labels')
plt.title('IN/OUT Confusion Matrix')
plt.show()

```

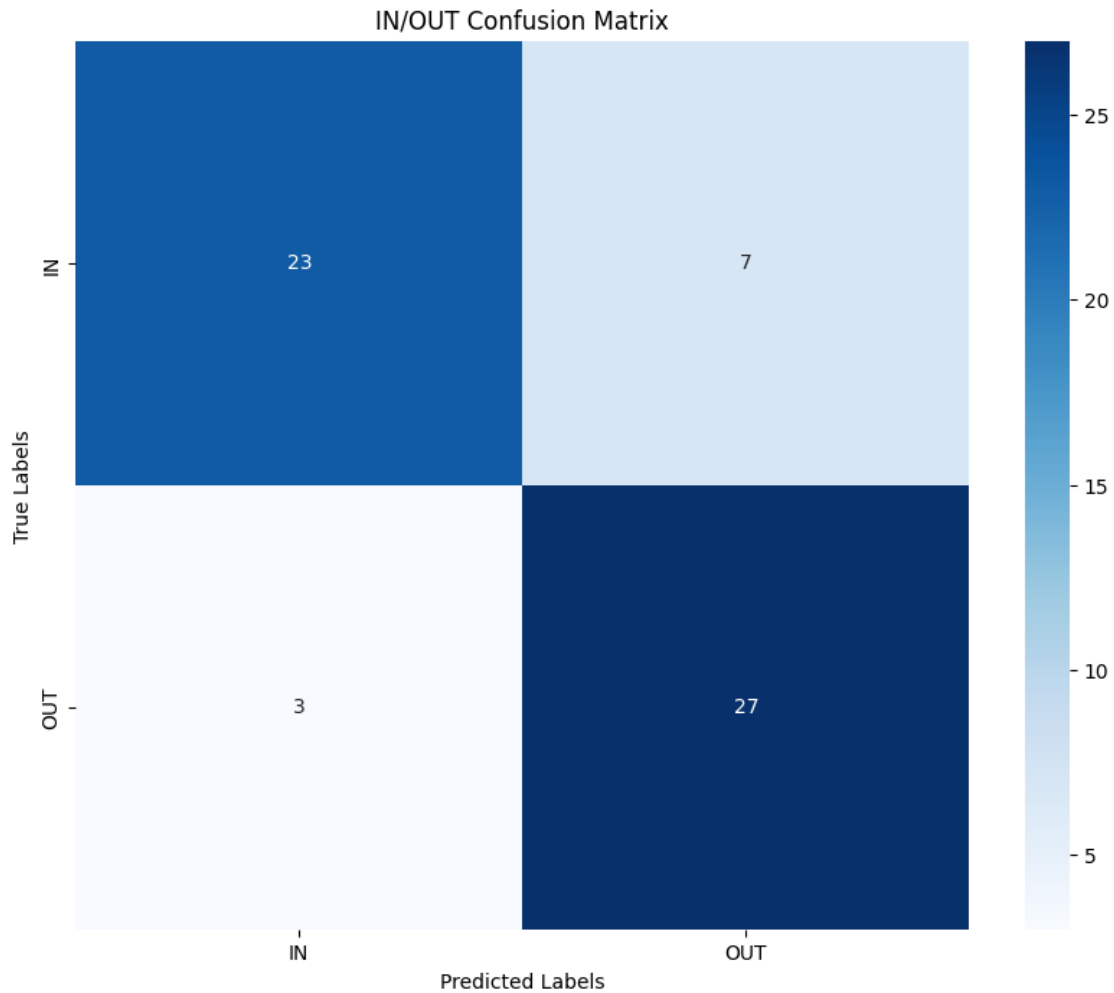
```

split_data/train
split_data/test
split_data/test_out

      : (60, 4096),      : (30, 4096),      : (30, 4096)
PCA      50      ...
PCA
SVM      ...
SVM
      : 0.737

====      =====
      : 0.83
      : 0.88
      : 0.77
      : 0.90
F1 : 0.82
Kappa : 0.67
AUC : 0.86

```

1.10

```
[156]: #
person_predictions = predict_person(classifiers, test_data_pca)

#
person_cm = confusion_matrix(test_labels, person_predictions)
person_labels = sorted(np.unique(train_labels))

#
person_accuracy = accuracy_score(test_labels, person_predictions)
person_f1_weighted = f1_score(test_labels, person_predictions,
    ↪average='weighted')
person_f1_macro = f1_score(test_labels, person_predictions, average='macro')
person_precision_macro = precision_score(test_labels, person_predictions,
    ↪average='macro')
```

```

person_recall_macro = recall_score(test_labels, person_predictions,
    ↪average='macro')
person_kappa = cohen_kappa_score(test_labels, person_predictions)

#
print("\n====      =====")
print(f"      : {person_accuracy:.2f}")
print(f" F1 : {person_f1_weighted:.2f}")
print(f" F1 : {person_f1_macro:.2f}")
print(f"      : {person_precision_macro:.2f}")
print(f"      : {person_recall_macro:.2f}")
print(f"Kappa : {person_kappa:.2f}")

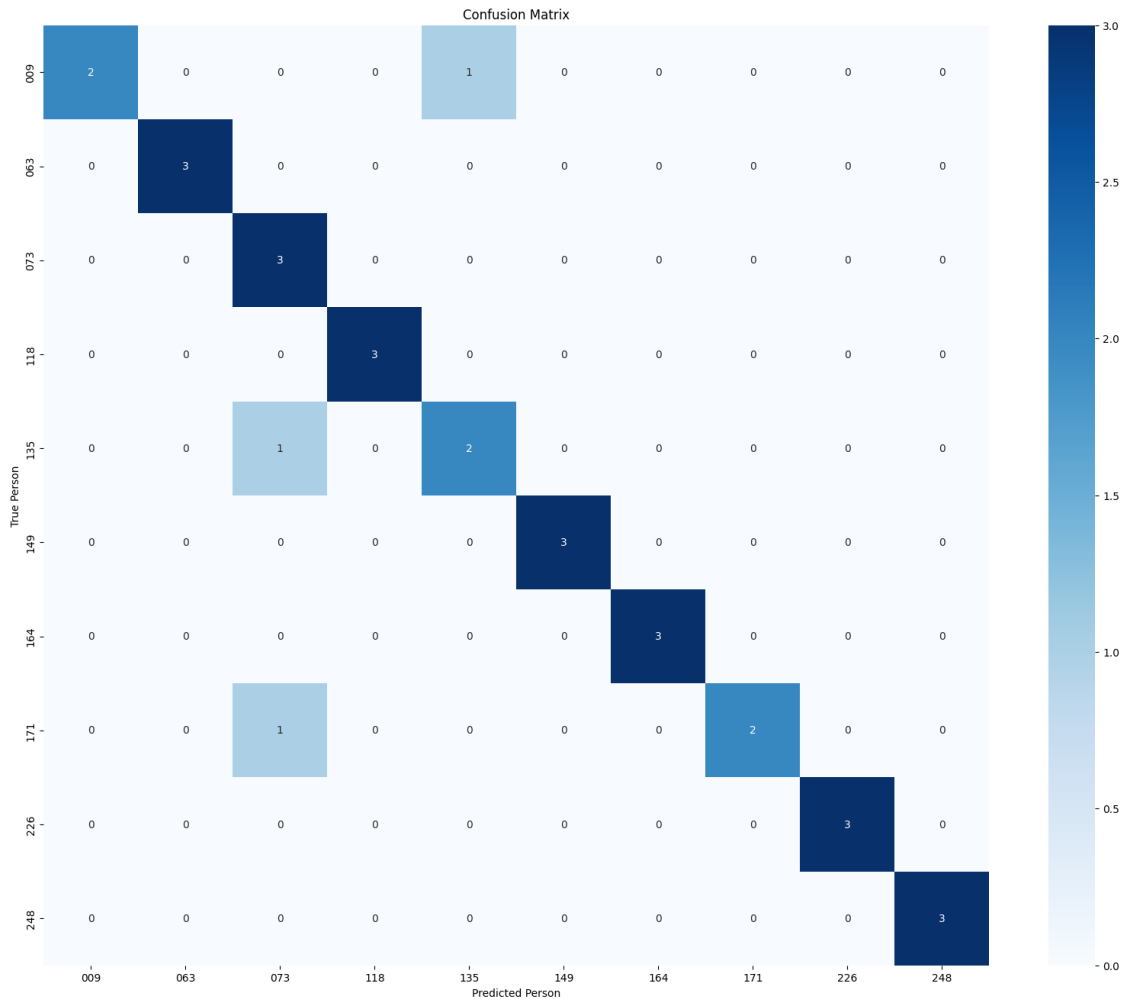
#
plt.figure(figsize=(20, 16))
sns.heatmap(person_cm, annot=True, fmt='d', cmap='Blues',
            xticklabels=person_labels, yticklabels=person_labels)
plt.xlabel('Predicted Person')
plt.ylabel('True Person')
plt.title('Confusion Matrix')
plt.show()

```

```

====      =====
      : 0.90
F1 : 0.90
F1 : 0.90
      : 0.93
      : 0.90
Kappa : 0.89

```



1.11

```
[154]: # n
def evaluate_different_n(train_dir, test_data, test_labels, test_out_data,
    test_out_labels, n_components, n_values, num_repeats=10):
    print("      n      ...")
    results = []
    for n in n_values:
        # print(f"\n  n = {n}")
        accuracies = []
        for repeat in range(num_repeats):
            # print(f"    {repeat + 1}/{num_repeats}    ...")
            #
            train_data, train_labels, _ = load_dataset(train_dir,
                max_images_per_person=n)
```

```

train_data_pca, test_data_pca, test_out_data_pca =
↳pca_reduction(train_data, test_data, test_out_data, n_components)

#
classifiers = train_svm_classifiers(train_data_pca, train_labels)

#
all_predictions, all_scores = vote_predict(classifiers, np.
↳concatenate([test_data_pca, test_out_data_pca]), 0.5)
true_labels = np.concatenate([np.ones(len(test_labels)), np.
↳zeros(len(test_out_labels))])
fpr, tpr, thresholds = roc_curve(true_labels, all_scores)
optimal_threshold = thresholds[np.argmax(tpr - fpr)]

#
test_predictions, _ = vote_predict(classifiers, test_data_pca,
↳optimal_threshold)
test_out_predictions, _ = vote_predict(classifiers,
↳test_out_data_pca, optimal_threshold)
all_predictions = np.concatenate([test_predictions,
↳test_out_predictions])
all_true_labels = np.concatenate([["IN"] * len(test_labels),
↳["OUT"] * len(test_out_labels)])

#
in_out_accuracy = accuracy_score(all_true_labels, all_predictions)
accuracies.append(in_out_accuracy)

#
mean_accuracy = np.mean(accuracies)
std_accuracy = np.std(accuracies)
print(f"n = {n},      : {mean_accuracy:.2f},      : {std_accuracy:.2f}")
results.append((n, mean_accuracy, std_accuracy))

#
n_values, mean_accuracies, std_accuracies = zip(*results)
plt.figure(figsize=(8, 6))
plt.errorbar(n_values, mean_accuracies, yerr=std_accuracies, fmt='o-',
↳color='b', ecolor='r', capsize=5)
plt.xlabel("Number of Images per Person (n)")
plt.ylabel("Accuracy")
plt.title("Accuracy vs Number of Images per Person")
plt.grid()
plt.show()
print("      n      ")

```

```

#         n_components
def evaluate_different_components(train_data, train_labels, test_data,
    ↪test_labels, test_out_data, test_out_labels, n_components_values,
    ↪num_repeats=10):
    print("         n_components         ...")
    results = []
    for n_components in n_components_values:
        # print(f"\n  n_components = {n_components}")
        accuracies = []
        for repeat in range(num_repeats):
            # print(f"    {repeat + 1}/{num_repeats}    ...")
            train_data_pca, test_data_pca, test_out_data_pca =
    ↪pca_reduction(train_data, test_data, test_out_data, n_components)

            #
            classifiers = train_svm_classifiers(train_data_pca, train_labels)

            #
            all_predictions, all_scores = vote_predict(classifiers, np.
    ↪concatenate([test_data_pca, test_out_data_pca]), 0.5)
            true_labels = np.concatenate([np.ones(len(test_labels)), np.
    ↪zeros(len(test_out_labels))])
            fpr, tpr, thresholds = roc_curve(true_labels, all_scores)
            optimal_threshold = thresholds[np.argmax(tpr - fpr)]

            #
            test_predictions, _ = vote_predict(classifiers, test_data_pca,
    ↪optimal_threshold)
            test_out_predictions, _ = vote_predict(classifiers,
    ↪test_out_data_pca, optimal_threshold)
            all_predictions = np.concatenate([test_predictions,
    ↪test_out_predictions])
            all_true_labels = np.concatenate(["IN" * len(test_labels),
    ↪["OUT" * len(test_out_labels)])

            #
            in_out_accuracy = accuracy_score(all_true_labels, all_predictions)
            accuracies.append(in_out_accuracy)

            #
            mean_accuracy = np.mean(accuracies)
            std_accuracy = np.std(accuracies)
            print(f"n_components = {n_components},      : {mean_accuracy:.2f},      :
    ↪{std_accuracy:.2f}")
            results.append((n_components, mean_accuracy, std_accuracy))

```

```

#
n_components_values, mean accuracies, std accuracies = zip(*results)
plt.figure(figsize=(8, 6))
plt.errorbar(n_components_values, mean accuracies, yerr=std accuracies,
fmt='o-', color='g', ecolor='r', capsize=5)
plt.xlabel("Number of Principal Components (n_components)")
plt.ylabel("Accuracy")
plt.title("Accuracy vs Number of Principal Components")
plt.grid()
plt.show()
print("      n_components      ")

#      n      n_components
if __name__ == "__main__":
    #
    test_data, test_labels, _ = load_dataset(test_dir, max_images_per_person=n)
    test_out_data, test_out_labels, _ = load_dataset(test_out_dir,
max_images_per_person=n)

    #      n
    n_values = [1,2,3,4,5,6] #
    evaluate_different_n(train_dir, test_data, test_labels, test_out_data,
test_out_labels, n_components=50, n_values=n_values)

    #      n = 6
    train_data, train_labels, _ = load_dataset(train_dir,
max_images_per_person=6)

    #      n_components
    n_components_values = [10, 20, 30, 40, 50]
    evaluate_different_components(train_data, train_labels, test_data,
test_labels, test_out_data, test_out_labels, n_components_values)

```

```

split_data/test
split_data/test_out
    n      ...
split_data/train
    PCA      50      ...
: n_components=50      10
PCA
    SVM      ...
SVM
split_data/train
    PCA      50      ...
: n_components=50      10
PCA

```

[illegible]


```

PCA
  SVM ...
SVM
split_data/train
  PCA 50 ...
  : n_components=50      20
PCA
  SVM ...
SVM
split_data/train
  PCA 50 ...
  : n_components=50      20
PCA
  SVM ...
SVM
n = 2,      : 0.63,      : 0.05
split_data/train
  PCA 50 ...
  : n_components=50      30
PCA
  SVM ...
SVM
split_data/train
  PCA 50 ...
  : n_components=50      30
PCA
  SVM ...
SVM
split_data/train
  PCA 50 ...
  : n_components=50      30
PCA
  SVM ...
SVM
split_data/train
  PCA 50 ...
  : n_components=50      30
PCA
  SVM ...
SVM
split_data/train
  PCA 50 ...

```

```

: n_components=50      30
PCA
SVM ...
SVM
split_data/train
PCA 50 ...
: n_components=50      30
PCA
SVM ...
SVM
split_data/train
PCA 50 ...
: n_components=50      30
PCA
SVM ...
SVM
split_data/train
PCA 50 ...
: n_components=50      30
PCA
SVM ...
SVM
split_data/train
PCA 50 ...
: n_components=50      30
PCA
SVM ...
SVM
n = 3, : 0.70, : 0.03
split_data/train
PCA 50 ...
: n_components=50      40
PCA
SVM ...
SVM
split_data/train
PCA 50 ...
: n_components=50      40
PCA
SVM ...
SVM
split_data/train
PCA 50 ...
: n_components=50      40
PCA
SVM ...
SVM
split_data/train

```

[illegible]



```

n
split_data/train
n_components  ...
PCA      10  ...
PCA
SVM      ...
SVM
PCA      10  ...
PCA
SVM      ...
SVM
PCA      10  ...
PCA
SVM      ...
SVM
PCA      10  ...
PCA
SVM      ...
SVM
PCA      10  ...

```

```

PCA
  SVM ...
SVM
  PCA 10 ...
PCA
  SVM ...
SVM
  PCA 10 ...
PCA
  SVM ...
SVM
  PCA 10 ...
PCA
  SVM ...
SVM
  PCA 10 ...
PCA
  SVM ...
SVM
  PCA 10 ...
PCA
  SVM ...
SVM
n_components = 10, : 0.78, : 0.02
  PCA 20 ...
PCA
  SVM ...
SVM
  PCA 20 ...
PCA
  SVM ...
SVM
  PCA 20 ...
PCA
  SVM ...
SVM
  PCA 20 ...
PCA
  SVM ...
SVM
  PCA 20 ...
PCA
  SVM ...
SVM

```

PCA	20	...
PCA		
SVM	...	
SVM		
PCA	20	...
PCA		
SVM	...	
SVM		
PCA	20	...
PCA		
SVM	...	
SVM		
PCA	20	...
PCA		
SVM	...	
SVM		
n_components = 20,		: 0.81, : 0.02
PCA	30	...
PCA		
SVM	...	
SVM		
PCA	30	...
PCA		
SVM	...	
SVM		
PCA	30	...
PCA		
SVM	...	
SVM		
PCA	30	...
PCA		
SVM	...	
SVM		
PCA	30	...
PCA		
SVM	...	
SVM		
PCA	30	...
PCA		
SVM	...	
SVM		
PCA	30	...
PCA		
SVM	...	


```

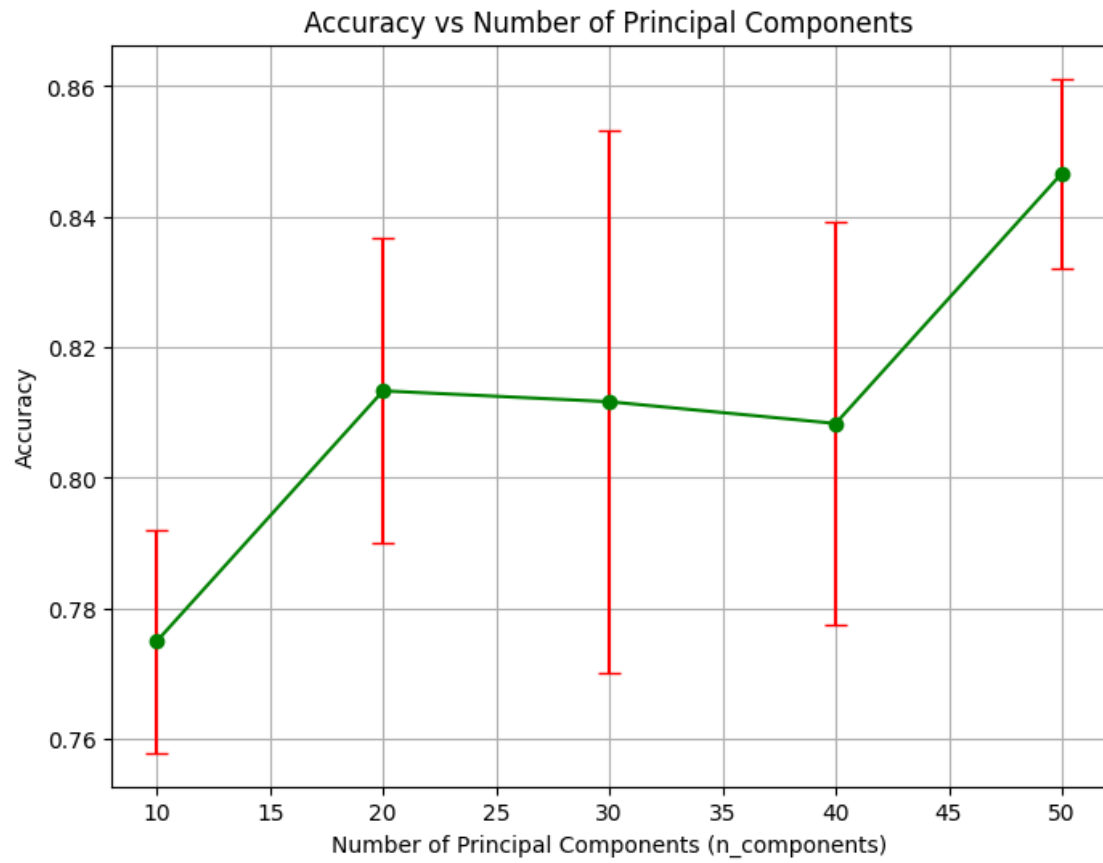
SVM
  PCA      30  ...
PCA
  SVM      ...
SVM
  PCA      30  ...
PCA
  SVM      ...
SVM
n_components = 30,      : 0.81,      : 0.04
  PCA      40  ...
PCA
  SVM      ...
SVM
  PCA      40  ...
PCA
  SVM      ...
SVM
  PCA      40  ...
PCA
  SVM      ...
SVM
  PCA      40  ...
PCA
  SVM      ...
SVM
  PCA      40  ...
PCA
  SVM      ...
SVM
  PCA      40  ...
PCA
  SVM      ...
SVM
  PCA      40  ...
PCA
  SVM      ...
SVM
  PCA      40  ...
PCA
  SVM      ...
SVM
  PCA      40  ...
PCA

```

```

    SVM    ...
SVM
n_components = 40,      : 0.81,      : 0.03
    PCA    50    ...
PCA
    SVM    ...
SVM
    PCA    50    ...
PCA
    SVM    ...
SVM
    PCA    50    ...
PCA
    SVM    ...
SVM
    PCA    50    ...
PCA
    SVM    ...
SVM
    PCA    50    ...
PCA
    SVM    ...
SVM
    PCA    50    ...
PCA
    SVM    ...
SVM
    PCA    50    ...
PCA
    SVM    ...
SVM
n_components = 50,      : 0.85,      : 0.01

```



n_components