## eigenface

#### April 23, 2025

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[48]: import zipfile
      import numpy as np
      import matplotlib.pyplot as plt
      from PIL import Image
      from sklearn.decomposition import PCA
      from sklearn.naive bayes import GaussianNB
      from sklearn.model_selection import KFold
      from sklearn.metrics import confusion matrix, classification report
      from sklearn.preprocessing import StandardScaler, LabelEncoder
      import seaborn as sns
      import pandas as pd
[49]: # Load Dataset
      zip_path = './orl_faces.zip'
[50]: # Lists for image data and labels
      images = []
      labels = []
[51]: # Pre-processing
      with zipfile.ZipFile(zip_path, 'r') as z:
          for file info in z.infolist():
              if file_info.filename.endswith(".pgm"): # ORL dataset images
                  parts = file info.filename.split('/')
                  if len(parts) >= 2:
                      label = parts[1] # Extract label from folder name
                      # Convert to grayscale
                      with z.open(file_info.filename) as file:
                          image = Image.open(file).convert("L")
                          images.append(np.array(image, dtype=np.float32))
                          labels.append(label)
[52]: # Convert images into a matrix where each row is a flattened image vector
      X = np.array([img.flatten() for img in images])
      y = np.array(labels)
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[53]: # Convert labels to numerical values
      le = LabelEncoder()
      y_encoded = le.fit_transform(y)
[54]: # Compute the mean vector (average face)
      mean_face = np.mean(X, axis=0)
[55]: # Subtract the mean from the dataset
      X_centered = X - mean_face
[56]: # Compute the covariance matrix
      cov_matrix = np.cov(X_centered, rowvar=False)
[57]: # Compute eigenvalues and eigenvectors
      eigenvalues, eigenvectors = np.linalg.eigh(cov_matrix)
[58]: # Sort eigenvalues & eigenvectors in descending order
      sorted_indices = np.argsort(eigenvalues)[::-1]
      eigenvalues = eigenvalues[sorted indices]
      eigenvectors = eigenvectors[:, sorted_indices]
[59]: # Normalize eigenvectors using Z-score normalization
      scaler = StandardScaler()
      eigenvectors_normalized = scaler.fit_transform(eigenvectors)
[60]: | # Reduce dimensionality using PCA (keeping top components)
      n_components = 100  # No. of principal components to keep
      pca = PCA(n_components=n_components, whiten=True, random_state=42)
      X_pca = pca.fit_transform(X_centered)
[61]: # 10-fold Cross Validation
      kf = KFold(n_splits=10, shuffle=True, random_state=42)
      conf matrices = []
      reports = []
      for train_index, test_index in kf.split(X_pca):
          X_train, X_test = X_pca[train_index], X_pca[test_index]
          y_train, y_test = y_encoded[train_index], y_encoded[test_index]
          # Compute standard deviation & mean for training data
          train_mean = np.mean(X_train, axis=0)
          train_std = np.std(X_train, axis=0)
          # Compute mean for test data
          test_mean = np.mean(X_test, axis=0)
          # Train Naive Bayes classifier
```

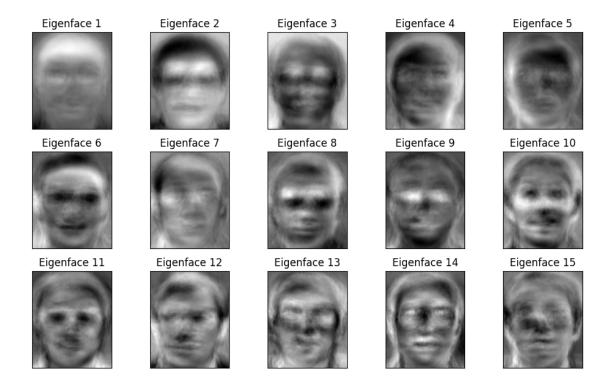
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clf = GaussianNB()
clf.fit(X_train, y_train)

# Confusion matrix

y_pred = clf.predict(X_test)
conf_matrices.append(confusion_matrix(y_test, y_pred, labels=np.

ounique(y_encoded)))
reports.append(classification_report(y_test, y_pred, zero_division=0))
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[65]: # Eigenfaces Visualization
     img_shape = images[0].shape
     eigenfaces = pca.components_reshape((n_components, img_shape[0], img_shape[1]))
     fig, axes = plt.subplots(3, 5, figsize=(10, 6), subplot_kw={'xticks': [],__
      for i, ax in enumerate(axes.flat):
         if i < n_components:</pre>
             ax.imshow(eigenfaces[i], cmap='gray')
             ax.set_title(f"Eigenface {i+1}")
     plt.tight_layout()
     plt.show()
     mean_face = np.mean(images, axis=0)
     plt.figure(figsize=(4, 4))
     plt.imshow(mean_face.reshape(img_shape), cmap='gray')
     plt.title("Average Face")
     plt.axis('off')
     plt.show()
```



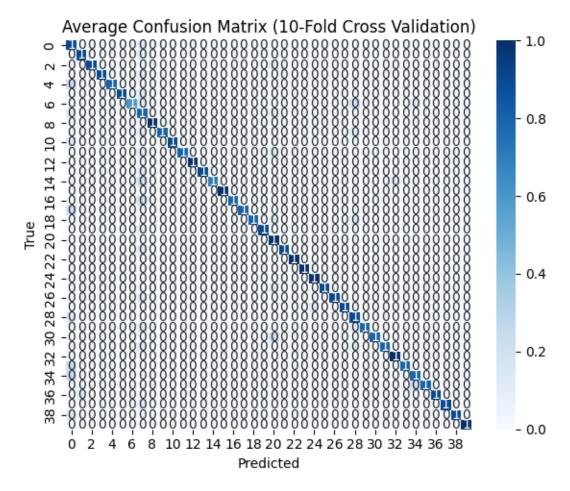
## Average Face



[67]: # Compute & display average confusion matrix avg\_conf\_matrix = np.mean(conf\_matrices, axis=0)

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plt.figure(figsize=(6,5))
sns.heatmap(avg_conf_matrix, annot=True, fmt=".0f", cmap="Blues")
plt.title("Average Confusion Matrix (10-Fold Cross Validation)")
plt.xlabel("Predicted")
plt.ylabel("True")
plt.tight_layout()
plt.show()
# Display final classification report
report_dfs = []
for report_str in reports:
    lines = report_str.split("\n")
    lines = [line.strip() for line in lines if line.strip()]
    data_lines = [line for line in lines if line[0].isdigit()]
    for i, line in enumerate(data_lines):
        tokens = line.split()
        if len(tokens) == 5: # Handle standard rows
            label, prec, rec, f1, support = tokens
            report_dfs.append({
                'label': label,
                'precision': float(prec),
                'recall': float(rec),
                'f1-score': float(f1),
                'support': int(support)
            })
df_reports = pd.DataFrame(report_dfs)
df_accuracy = pd.DataFrame({
    "Principal Components": pcs,
    "Accuracy Without Z-Score (%)": accuracy_no_zscore,
    "Accuracy With Z-Score (%)": accuracy_zscore
})
# Print the table
print("\nAccuracy Comparison Table:")
print(df_accuracy.to_string(index=False))
# Group by label and average metrics
avg_report = df_reports.groupby('label')[['precision', 'recall', 'f1-score']].
 →mean()
print("\nAverage Classification Report (10-Fold Cross Validation):")
print(avg_report.round(2))
pcs = [100, 500, 1000, 2000, 5000, 10000, 15000, 20000, 22500]
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```
accuracy_zscore = [36.5, 44.5, 51.0, 51.5, 74.0, 81.0, 86.0, 88.5, 88.5]
accuracy_no_zscore = [27.0, 31.5, 34.0, 31.5, 43.5, 68.0, 71.0, 71.5, 69.0]
x = np.arange(len(pcs)) # label locations
width = 0.35
                         # width of the bars
plt.figure(figsize=(6, 3))
plt.bar(x - width/2, accuracy_no_zscore, width, label='Without Z-Score')
plt.bar(x + width/2, accuracy_zscore, width, label='With Z-Score')
plt.xlabel('Number of Principal Components')
plt.ylabel('Accuracy (%)')
plt.title('Accuracy Comparison: With vs Without Z-Score Normalization')
plt.xticks(x, pcs)
plt.legend()
plt.grid(True, axis='y')
plt.tight_layout()
plt.show()
```



### Accuracy Comparison Table:

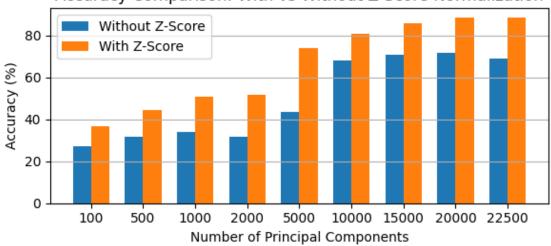
Principal Components	Accuracy Without Z-Score (%)	Accuracy With Z-Score (%)
100	27.0	36.5
500	31.5	44.5
1000	34.0	51.0
2000	31.5	51.5
5000	43.5	74.0
10000	68.0	81.0
15000	71.0	86.0
20000	71.5	88.5
22500	69.0	88.5

# Average Classification Report (10-Fold Cross Validation): precision recall f1-score

	precision	recall	f1-score
label			
0	0.44	0.67	0.50
1	0.75	0.75	0.75
10	1.00	0.94	0.96
11	1.00	0.79	0.87
12	1.00	1.00	1.00
13	0.88	0.88	0.88
14	1.00	0.88	0.90
15	1.00	1.00	1.00
16	1.00	0.83	0.89
17	1.00	0.86	0.91
18	0.83	0.78	0.80
19	1.00	0.92	0.94
2	1.00	0.93	0.95
20	0.65	0.70	0.67
21	0.86	0.86	0.86
22	1.00	1.00	1.00
23	1.00	1.00	1.00
24	0.86	0.86	0.86
25	0.88	0.88	0.88
26	1.00	0.93	0.96
27	0.88	0.88	0.88
28	0.72	0.81	0.75
29	0.86	0.79	0.81
3	0.88	0.88	0.88
30	0.71	0.71	0.71
31	0.86	0.79	0.81
32	0.95	1.00	0.97
33	0.86	0.81	0.83
34	0.69	0.79	0.69
35	0.86	0.79	0.81
36	0.86	0.86	0.86
37	1.00	0.96	0.98

38	0.83	0.83	0.83
39	1.00	1.00	1.00
4	1.00	0.83	0.89
5	1.00	0.92	0.94
6	1.00	0.66	0.75
7	0.44	0.50	0.45
8	1.00	1.00	1.00
9	0.71	0.71	0.71

## Accuracy Comparison: With vs Without Z-Score Normalization



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