



```
eigvals = eigvals[::-1]
eigvecs = eigvecs[:,::-1]
```

```
return (eigvecs,eigvals)
```

```
#takes data with rows as features, columns as samples
```

```
def pcaReduce(data,n_components): #return eigvals and reduced data, and mean it subtrac
    mean = np.mean(data,axis=1).reshape(-1,1)
    vecs, vals = pca(data - mean)
    PCAmat = vecs.T[:n_components]
    reduced = PCAmat @ (data - mean)
    return reduced,vals,mean, PCAmat
```

```
In [ ]: X_train, X_test, y_train, y_test = train_test_split(audio_data, labels, test_size=.25,
sort_indices = np.argsort(y_train))
```

```
# Sort X_train and y_train by the labels
```

```
X_train = [X_train[i] for i in sort_indices]
y_train = y_train[sort_indices]
```

```
# Verify the shapes of the resulting sets
```

```
print("X_train_sorted shape:", len(X_train))
print("y_train_sorted shape:", y_train.shape)
```

```
X_train_sorted shape: 150
y_train_sorted shape: (150,)
```

```
In [ ]: nPCA = 5
reducedTrain,eigvals,PCAmat = pcaReduce(np.hstack(X_train),nPCA)
temp = []
startidx =0
for i in range(len(X_train)):
    temp.append(reducedTrain[:,startidx:startidx+X_train[i].shape[1]])
    startidx += X_train[i].shape[1]
reducedTrain=temp
```

```
In [ ]: def fitHmm(spectrogramList, n_states, n_gaussians):
    model = hmm.GMMHMM(n_components=n_states, n_mix=n_gaussians, covariance_type='diag

    X = np.hstack(spectrogramList)

    lengths = [spec.shape[1] for spec in spectrogramList]
    X = X.T

    model.fit(X, lengths=lengths)
    return model

def evaluatefromHmms(model_list, spectrogram):
    scores = np.zeros(len(model_list))
    for i in range(len(model_list)):
        scores[i], _ = model_list[i].decode(spectrogram.T)
    return np.argmax(np.exp(scores))
```

```
In [ ]: digitModels = []
n_states = 5
n_gaussians = 1
for i in range(10):
    # print(i)
    digitModels.append(fitHmm(reducedTrain[i*15:(i+1)*15], n_states=n_states, n_gaussians=n_gaussians))
```

```
In [ ]: %%capture

trainConfMat = np.zeros((10,10))
testConfMat = np.zeros((10,10))

for i in range(len(X_train)):
    yhat = evaluatefromHmms(digitModels, PCAmat @(X_train[i] - PCAmean))
    y = y_train[i]
    trainConfMat[y,yhat] +=1

for i in range(len(X_test)):
    yhat = evaluatefromHmms(digitModels, PCAmat @(X_test[i] - PCAmean))
    y = y_test[i]
    testConfMat[y,yhat] +=1
```

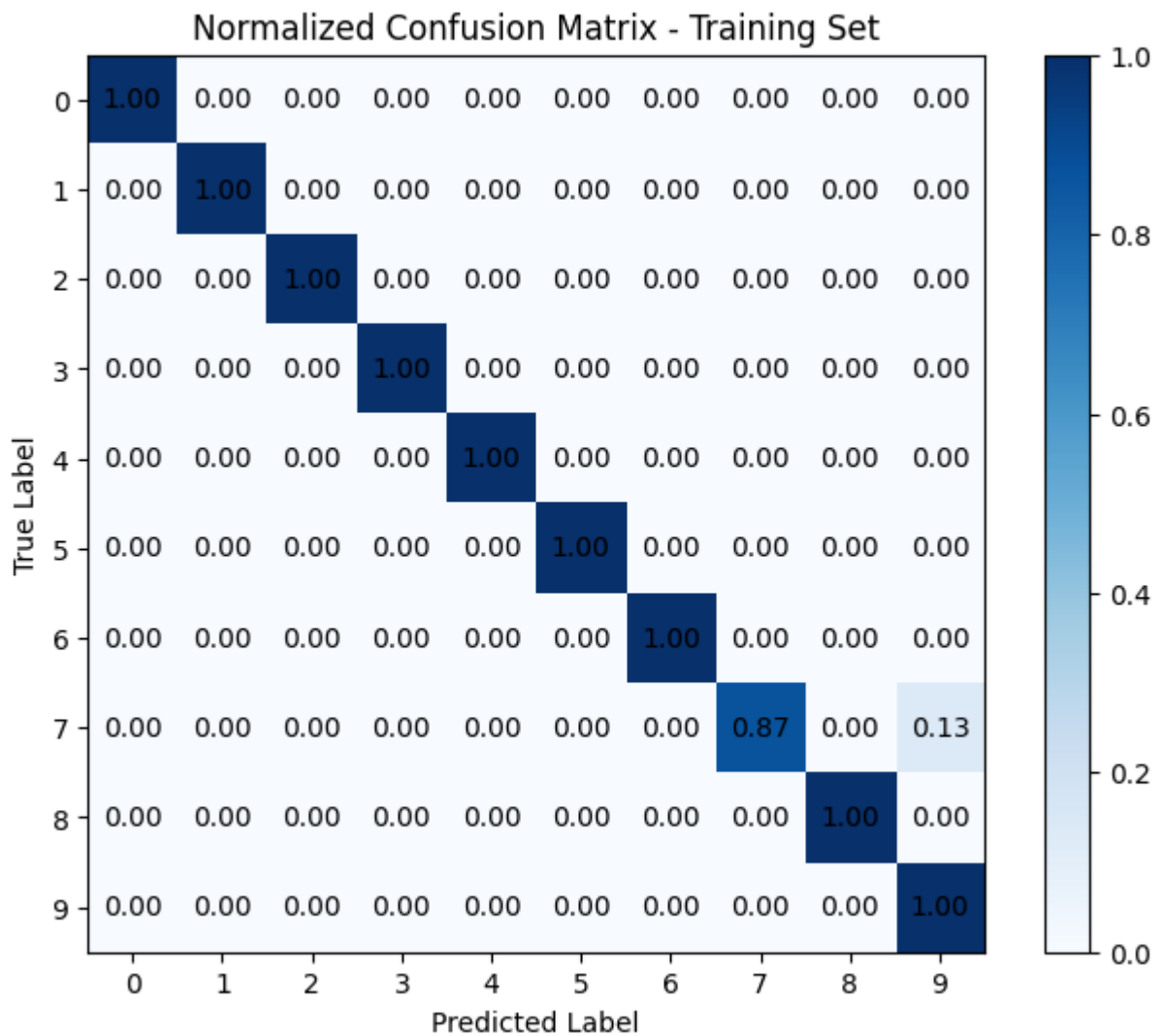
```
In [ ]: trainConfMatNormalized = trainConfMat.astype('float') / trainConfMat.sum(axis=1)[:, np.newaxis]
plt.figure(figsize=(8, 6))
cax = plt.imshow(trainConfMatNormalized, cmap="Blues")
for i in range(trainConfMatNormalized.shape[0]):
    for j in range(trainConfMatNormalized.shape[1]):
        plt.text(j, i, f'{trainConfMatNormalized[i, j]:.2f}', ha='center', va='center')

plt.xticks(np.arange(trainConfMatNormalized.shape[1]))
plt.yticks(np.arange(trainConfMatNormalized.shape[0]))
plt.xlabel('Predicted Label')
plt.ylabel('True Label')
plt.title('Normalized Confusion Matrix - Training Set')

cbar = plt.colorbar(cax)
print("Using", PCAmat.shape[0], " PCA components and ", n_states, "HMM states per model")

plt.show()
```

Using 5 PCA components and 5 HMM states per model



```
In [ ]: testConfMatNormalized = testConfMat.astype('float') / testConfMat.sum(axis=1)[:, np.newaxis]
plt.figure(figsize=(8, 6))
cax = plt.imshow(testConfMatNormalized, cmap="Blues")
for i in range(testConfMatNormalized.shape[0]):
    for j in range(testConfMatNormalized.shape[1]):
        plt.text(j, i, f'{testConfMatNormalized[i, j]:.2f}', ha='center', va='center',
                 color='white' if testConfMatNormalized[i, j] > 0.5 else 'black')

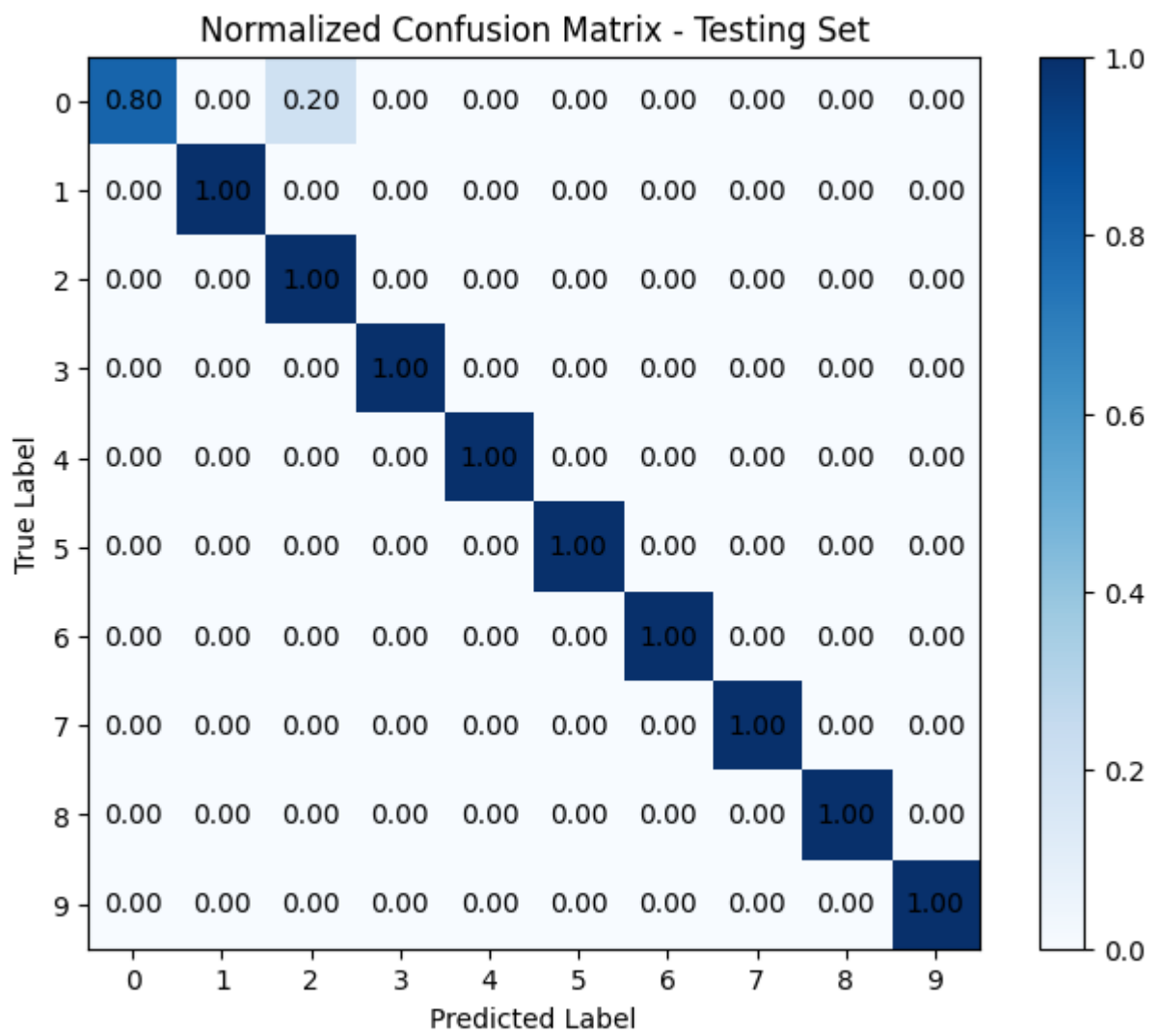
plt.xticks(np.arange(testConfMatNormalized.shape[1]))
plt.yticks(np.arange(testConfMatNormalized.shape[0]))
plt.xlabel('Predicted Label')
plt.ylabel('True Label')
plt.title('Normalized Confusion Matrix - Testing Set')

print("Using", PCAMat.shape[0], " PCA components and ", n_states, "HMM states per model")

cbar = plt.colorbar(cax)

plt.show()
```

Using 5 PCA components and 5 HMM states per model



In [ ]: