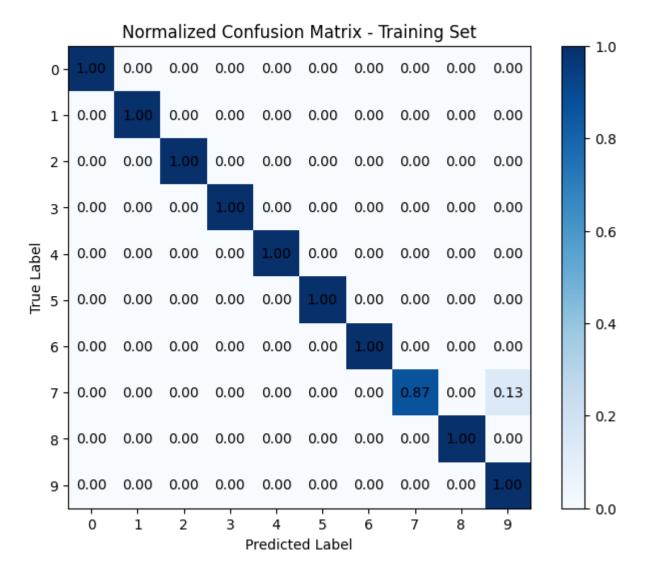
```
In [ ]: from scipy.io import wavfile
      from scipy.signal import stft
      import matplotlib.pyplot as plt
      import numpy as np
      # import librosa as librosa
      from hmmlearn import hmm
      import os
      from sklearn.model selection import train test split
      np.random.seed(1)
In [ ]: def load audio data(folder path, num recordings per digit=20):
          data = []
          labels = []
         for digit in range(10):
             digit count = 0
             # maxlen = 0
             for filename in os.listdir(folder path):
                if filename.endswith(".wav") and filename.startswith('digit'+str(digit)):
                   file_path = os.path.join(folder_path, filename)
                   _ , audio_data = wavfile.read(file_path)
                   data.append(audio data)
                   labels.append(digit)
                   digit count += 1
                   if digit_count == num_recordings_per_digit:
          return data, np.array(labels)
      folder path = './digitRecordings/'
      audio_data, labels = load_audio_data(folder_path)
      n fft = 1024
      noverlap = 512
      # n mfcc = 13
      for i in range(len(audio data)):
         __,_ audio_data[i] = stft(audio_data[i], nperseg=n_fft, noverlap=noverlap)
         audio_data[i] = 10*np.log(np.abs(audio_data[i]) + 1e-14)
          # audio_data[i] = librosa.feature.mfcc(audio_data[i].astype(float),sr = 10,n_me=n_
      print("Audio Data Shape:", len(audio_data))
      print("Labels:", labels)
      Audio Data Shape: 200
      In [ ]: def pca(data): #takes input data as columns and expects it to be 0 mean (add mean back
          covariance matrix = np.dot(data,data.T)/(data.shape[1])
          eigvals, eigvecs = np.linalg.eigh(covariance_matrix)
```

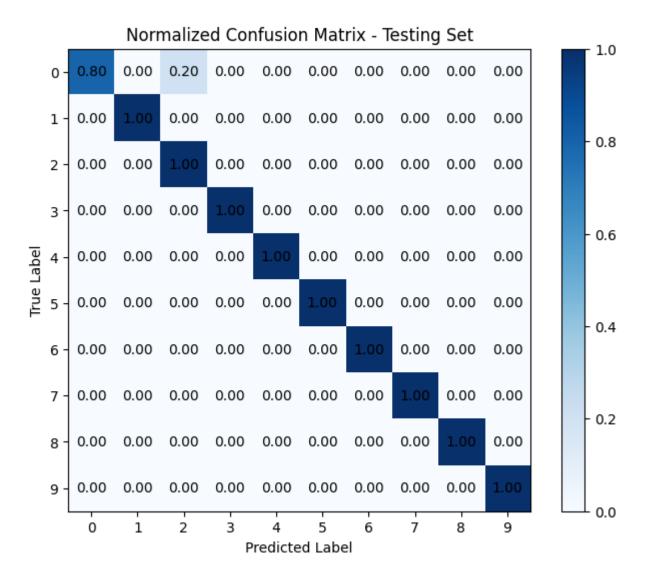
```
eigvals = eigvals[::-1]
             eigvecs = eigvecs[:,::-1]
             return (eigvecs, eigvals)
        #takes data with rows as features, columns as samples
        def pcaReduce(data, n components): #return eigvals and reduced data, and mean it subtra
            mean = np.mean(data,axis=1).reshape(-1,1)
            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,PCAmean,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 gaussia
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
        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 mode
        plt.show()
```

Using 5 PCA components and 5 HMM states per model



Using 5 PCA components and 5 HMM states per model



In []: