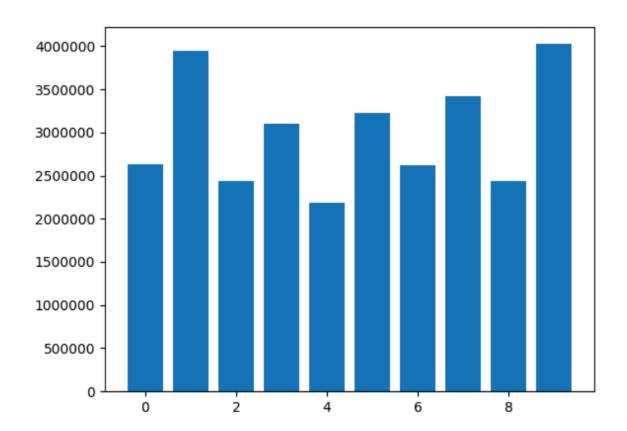
## CS498 AML HW4

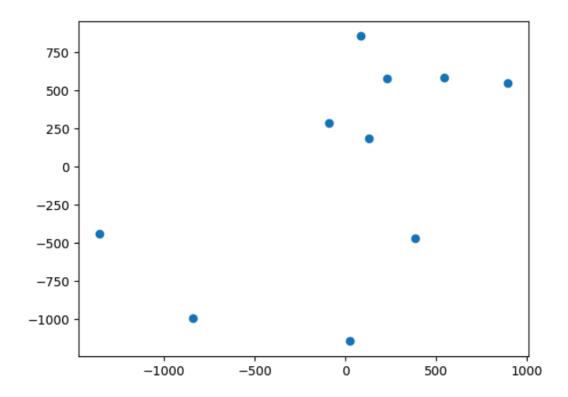
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## 1. MSEs of category $0 \sim 9$ .

We use the MSEs between the training data and the reconstructed **training** data. (We personally think this makes more sense.)



## 2. 2D Map of Principal Coordinates Analysis.



## 3. Code Screenshots.

```
from sklearn.decomposition import PCA
        from scipy.spatial import distance_matrix
 3
        from skbio.stats.ordination import pcoa
 4
        import numpy as np
 5
        import pickle
 6
        from matplotlib import pyplot as plt
        from sklearn import manifold
 8
9
        def unpickle(file):
10
            with open(file, 'rb') as fo:
11
                dict = pickle.load(fo, encoding='bytes')
12
13
            return dict
14
15
        def pca(data):
16
17
            pca = PCA(n_components=20)
18
            components = pca.fit_transform(data)
19
            reconstruct = pca.inverse_transform(components)
20
            return reconstruct
21
22
23
       def divide(raw_data, labels, data):
                                                           # Separate out the 10 categories
24
            for i in range(len(labels)):
                if labels[i] not in data.keys():
25
26
                    data[labels[i]] = [raw_data[i]]
27
                else:
28
                    data[labels[i]].append(raw_data[i])
29
            return data
30
31
32
        def mse(mat1, mat2):
            return 1/5000* np.sum((mat1-mat2)**2)
33
    -After parsing data-----
62
             data = dict()
             data = divide(raw_data1, labels1, data)
data = divide(raw_data2, labels2, data)
63
64
             data = divide(raw_data3, labels3, data)
65
             data = divide(raw_data4, labels4, data)
66
67
             data = divide(raw_data5, labels5, data)
68
69
             error = [0 \text{ for } i \text{ in } range(10)]
             meanImages = np.zeros((10, 3072))
70
71
72
             for i in range(10):
73
                 meanImages[i] = np.mean(data[i], axis=_0)
74
                  recon = pca(data[i])
75
                  for j in range(len(data[i])):
76
                      error[i] += mse(data[i][j], recon[j])
77
78
             plt.figure(0)
79
             plt.bar(range(1, len(error)+1), error)
80
             plt.show()
81
             # 7.7 (b)
82
83
             dis_mat = distance_matrix(meanImages, meanImages)
84
             plt.figure(1)
85
             mds = manifold.MDS(dissimilarity='precomputed')
86
             results = mds.fit(dis_mat)
87
             coords = results.embedding_
88
             plt.scatter(coords[:, 0], coords[:, 1])
89
90
             plt.show()
```