AMATH797 PSet4 David Lieberman

```
In [1]:
        import os
        import numpy as np
        import sklearn
        from sklearn import manifold
        from sklearn import datasets
        from sklearn.datasets import fetch_openml
        from sklearn.utils import check_random_state
        from sklearn.cluster import KMeans
        from sklearn.neighbors import kneighbors_graph
        from sklearn.cluster import SpectralClustering
        np.random.seed(0)
        import networkx as nx
        import scipy
        from scipy.sparse import csgraph
        import matplotlib
        import matplotlib.pyplot as plt
        %matplotlib inline
        %config InlineBackend.figure_format = 'svg'
        # from google.colab import drive
        # drive.mount('/content/drive')
        # cd 'drive/My Drive/2020S_AMATH797/PSet3'
        os.chdir(os.path.expanduser(os.sep.join(["/mnt","c","Users","darkg","Desktop", "Homework
        Scans", "2020S_AMATH797", "PSet4"])))
        # os.chdir(os.path.expanduser(os.sep.join(["~","Desktop", "Homework Scans", "2020S_AMATH
        797", "PSet3"])))
```

Problem 2

Part A

Isomap is a non-linear dimensionality reduction method method with the goal of preserving the geodesic distance in the low-dimensional embedding. We use the k nearest neighbors algorithm to construct a neighborhood graph with the edge length as the Euclidean distance, and by computing the shortest path between two nodes we can approximate the geodesic distance on the manifold. The last step is to apply multi-dimensional scaling by taking the eigendecomposition of the geodesic proximity matrix to obtain the low dimensional embedding of the data. We try to minimize the "Strain" cost function in classical MDS:

$$\min_{x_1,\dots,x_M} \left(rac{\sum_{i,j} \left(b_{ij} - \langle x_i, x_j
angle
ight)^2}{\sum_{i,j} b_{ij}^2}
ight)^{1/2}$$

or the "Stress" in metric MDS:

$$\min_{x_1,\ldots,x_M} \left(\sum_{i < j} \left(\|x_i - x_j\| - d_{i,j}
ight)^2
ight)^{1/2}$$

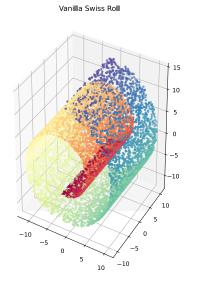
which is just the residual sum of squares.

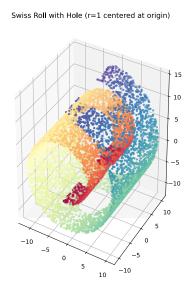
t-SNE is a non-linear dimensionality reduction method that constructs probability distributions over high dimensional objects such that similar objects are modeled as being close while dissimilar objects are modeled as being far away, with high probability. First, in high-dimensional space, t-SNE uses the Gaussian distribution to define relationships between the points, then recreates the probability distribution in low-dimensional space using the t-distribution with 1 degree of freedom (Cauchy). This helps to combat the curse of dimensionality often encountered when going from high to low-dimensional data, where moderate spacing in high dimensions clumps dramatically in lower dimensions. t-SNE uses gradient descent to obtain locations of the mapped points in lower-dimensional space by minimizing the KL divergence between the Gaussian and the t-distribution:

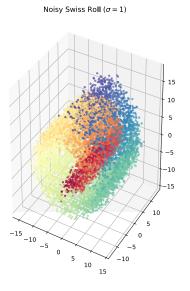
$$ext{KL}(P\|Q) = \sum_{i
eq j} p_{ij} \log rac{p_{ij}}{q_{ij}}$$

```
In [2]: # Vanilla
        vanilla_swiss_roll, vanilla_color = datasets.make_swiss_roll(n_samples=10000)
        vanilla_swiss_roll = vanilla_swiss_roll - np.mean(vanilla_swiss_roll, axis=0)
        # Hole
        hole_swiss_roll, hole_color = datasets.make_swiss_roll(n_samples=10000)
        hole_swiss_roll = hole_swiss_roll - np.mean(hole_swiss_roll, axis=0)
        keep = np.array([], dtype=bool)
        for point in zip(hole_swiss_roll):
          keep = np.append(keep, np.linalg.norm(point[0][0:2]) > 4)
        hole_swiss_roll = hole_swiss_roll[keep, ...]
        hole_color = hole_color[keep]
        # Noise
        noisy_swiss_roll, noisy_color = datasets.make_swiss_roll(n_samples=10000, noise=1.0)
        noisy_swiss_roll = noisy_swiss_roll - np.mean(noisy_swiss_roll, axis=0)
        fig = plt.figure(figsize=(20, 8))
        ax = fig.add_subplot(1, 3, 1, projection='3d')
        ax.scatter(vanilla_swiss_roll[:, 0], vanilla_swiss_roll[:, 1], vanilla_swiss_roll[:, 2],
        c=vanilla_color, cmap='Spectral', s=8)
        ax.title.set_text("Vanilla Swiss Roll")
        ax = fig.add_subplot(1, 3, 2, projection='3d')
        ax.scatter(hole_swiss_roll[:, 0], hole_swiss_roll[:, 1], hole_swiss_roll[:, 2], c=hole_c
        olor, cmap='Spectral', s=8)
        ax.title.set_text("Swiss Roll with Hole (r=1 centered at origin)")
        ax = fig.add_subplot(1, 3, 3, projection='3d')
        ax.scatter(noisy_swiss_roll[:, 0], noisy_swiss_roll[:, 1], noisy_swiss_roll[:, 2], c=noi
        sy_color, cmap='Spectral', s=8)
        ax.title.set text("Noisy Swiss Roll ($\sigma = 1$)")
        fig.suptitle("Original Data", fontsize=22)
        plt.show()
```

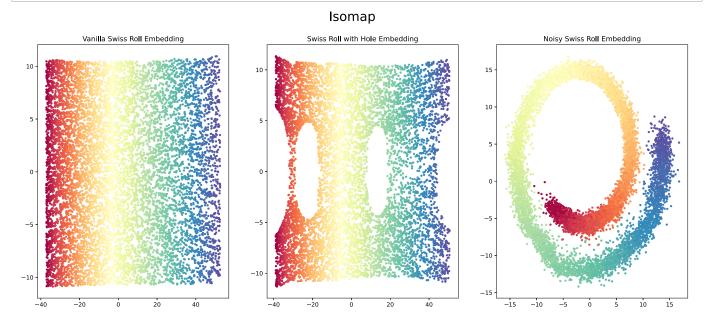






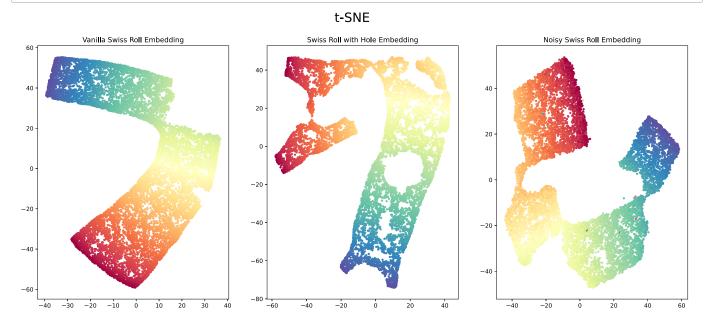


```
In [3]:
        # Isomap Vanilla
        Vanilla Iso = manifold.Isomap(n neighbors=30, n components=2).fit transform(vanilla swis
        s_roll)
        # Isomap Hole
        Hole_Iso = manifold.Isomap(n_neighbors=30, n_components=2).fit_transform(hole_swiss_roll
        # Isomap Noisy
        Noisy_Iso = manifold.Isomap(n_neighbors=30, n_components=2).fit_transform(noisy_swiss_ro
        11)
        fig = plt.figure(figsize=(20, 8))
        ax = fig.add_subplot(1, 3, 1)
        ax.scatter(Vanilla_Iso[:, 0], Vanilla_Iso[:, 1], c=vanilla_color, cmap='Spectral', s=8)
        ax.title.set_text("Vanilla Swiss Roll Embedding")
        ax = fig.add_subplot(1, 3, 2)
        ax.scatter(Hole_Iso[:, 0], Hole_Iso[:, 1], c=hole_color, cmap='Spectral', s=8)
        ax.title.set_text("Swiss Roll with Hole Embedding")
        ax = fig.add_subplot(1, 3, 3)
        ax.scatter(Noisy_Iso[:, 0], Noisy_Iso[:, 1], c=noisy_color, cmap='Spectral', s=8)
        ax.title.set_text("Noisy Swiss Roll Embedding")
        fig.suptitle("Isomap", fontsize=22)
        plt.show()
```



The isomap did an excellent job here at "unrolling" our swiss roll dataset into two dimensions, and both keeping connected neighborhoods of points together and preserving the geodesic distance along the manifold. We see in the noisy swiss roll embedding this geodesic distance preservation might have slightly lapsed as the red/orange points are very far from the blue points on the manifold while they appear close here in Euclidean space, but the spiral/manifold geometric structure is very well preserved, so we retain good intuition regarding the distance geodesic distance.

```
In [4]:
        # t-SNE Vanilla
        Vanilla_tSNE = manifold.TSNE(n_components=2, perplexity=200, learning_rate=250, n_iter=2
        500, init='random').fit_transform(vanilla_swiss_roll)
        # t-SNE Hole
        Hole_tSNE = manifold.TSNE(n_components=2, perplexity=95, init='random').fit_transform(ho
        le_swiss_roll)
        # t-SNE Noisy
        Noisy_tSNE = manifold.TSNE(n_components=2, perplexity=150, learning_rate=250, n_iter=200
        0, init='random').fit_transform(noisy_swiss_roll)
        fig = plt.figure(figsize=(20, 8))
        ax = fig.add_subplot(1, 3, 1)
        ax.scatter(Vanilla_tSNE[:, 0], Vanilla_tSNE[:, 1], c=vanilla_color, cmap='Spectral', s=8
        ax.title.set_text("Vanilla Swiss Roll Embedding")
        ax = fig.add_subplot(1, 3, 2)
        ax.scatter(Hole_tSNE[:, 0], Hole_tSNE[:, 1], c=hole_color, cmap='Spectral', s=8)
        ax.title.set_text("Swiss Roll with Hole Embedding")
        ax = fig.add_subplot(1, 3, 3)
        ax.scatter(Noisy_tSNE[:, 0], Noisy_tSNE[:, 1], c=noisy_color, cmap='Spectral', s=8)
        ax.title.set_text("Noisy Swiss Roll Embedding")
        fig.suptitle("t-SNE", fontsize=22)
        plt.show()
```



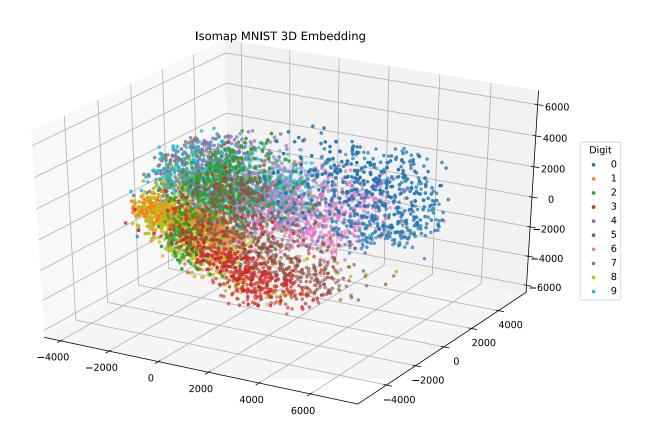
The swiss roll is a canonically pathological distribution for t-SNE, however we see here that with some tuning of the parameters, we still manage to retain decent results. Although there is some tearing along the region of the manifold with the high curvature, for the most part, the data remains connected in a single cluster. We note that the swiss roll with hole was particularly successful as it gives a good visualization of where the hole was excised from the dataset. Although there are a few scattered datapoints in the noisy swiss roll embedding, t-SNE was largely successful in keeping neighboring points in the manifold close to one another in the embedding, while keep distant points far apart.

Part C

```
In [5]: MNIST, labels = fetch_openml('mnist_784', version=1, return_X_y=True)
        random_state = check_random_state(0)
        permutation = random state.permutation(MNIST.shape[0])
        MNIST = MNIST[permutation]
        labels = labels[permutation]
        MNIST = MNIST[:6000, :]
        labels = labels[:6000]
        labels = [int(label) for label in labels]
        groups = []
        for i in np.arange(10):
          groups.append(np.where(labels == i))
In [6]: MNIST_Iso = manifold.Isomap(n_neighbors=15, n_components=3).fit_transform(MNIST)
        fig = plt.figure(figsize=(12,8))
        ax = plt.axes(projection='3d')
        for idx, group in enumerate(groups):
          ax.scatter(MNIST_Iso[group, 0], MNIST_Iso[group, 1], MNIST_Iso[group, 2], label=idx, s
        =8)
        ax.legend(title = "Digit", loc = "center right", borderaxespad=-2.)
```

plt.title("Isomap MNIST 3D Embedding")

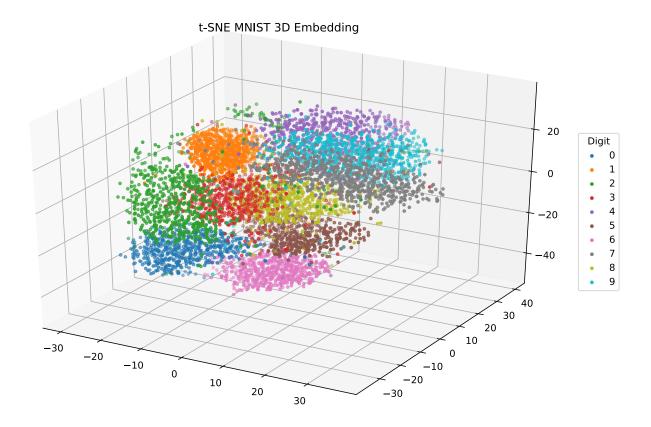
plt.show()



This embedding was largely unsuccessful. I tried many parameters, but this was just not a good embedding technique for this dataset. Just appears to be a scatter of points, with no clear structure maintained.

```
In [7]: MNIST_tSNE = manifold.TSNE(n_components=3, perplexity=150, learning_rate=250, n_iter=200
0).fit_transform(MNIST)

fig = plt.figure(figsize=(12,8))
    ax = plt.axes(projection='3d')
    for idx, group in enumerate(groups):
        ax.scatter(MNIST_tSNE[group, 0], MNIST_tSNE[group, 1], MNIST_tSNE[group, 2], label=idx
        , s=8)
    ax.legend(title = "Digit", loc = "center right", borderaxespad=-2.)
    plt.title("t-SNE MNIST 3D Embedding")
    plt.show()
```



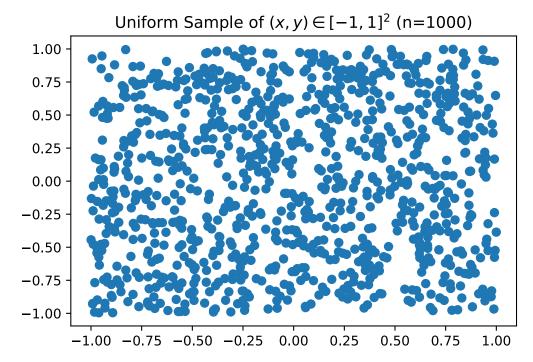
After much tuning, t-SNE was largely successful in seperating the MNIST data into clusters of the same digits which are largely disjoint from the other digit clusters.

Problem 3

Part A

```
In [8]: points = np.random.uniform(-1., 1., (1000,2))

plt.scatter(points[:,0], points[:,1])
plt.title("Uniform Sample of $(x,y) \in [-1,1]^2$ (n=1000)")
plt.show()
```



Part B

```
In [9]: Adjacency = kneighbors_graph(points, n_neighbors=30).toarray()
Adjacency_symmetric = np.maximum(Adjacency, Adjacency.T)

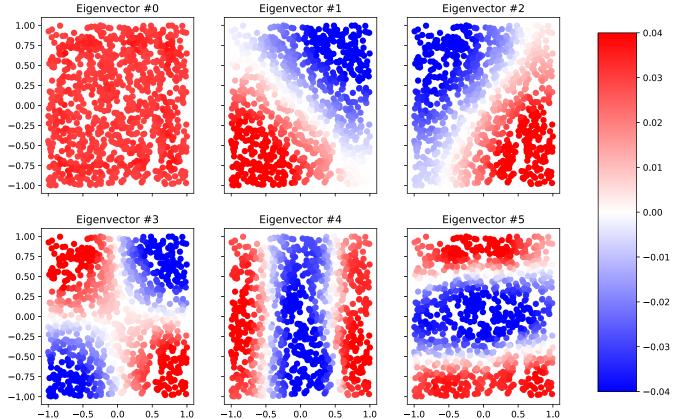
sigma = 1
Weighted_Adjacency = np.multiply(np.exp(-Adjacency_symmetric**2 / sigma**2), Adjacency_symmetric)
Weighted_Graph = nx.from_scipy_sparse_matrix(csgraph.csgraph_from_dense(Weighted_Adjacency))
```

Part C

```
In [10]: L_sym = nx.normalized_laplacian_matrix(Weighted_Graph)
    vals, vecs = np.linalg.eigh(L_sym.A)

In [11]: fig, axes = plt.subplots(nrows=2, ncols=3, figsize=(12,8), sharex=True, sharey=True)
    for idx, axes in enumerate(axes.flat):
        axes.set_title("Eigenvector #" + str(idx))
        ax = axes.scatter(points[:,0], points[:,1], c=vecs[:,idx], cmap='bwr', vmin=-0.04, v
        max=0.04)

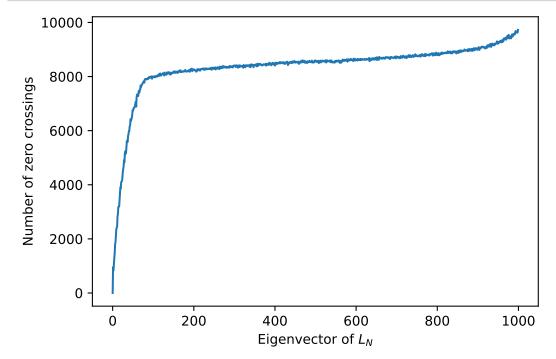
    fig.subplots_adjust(right=0.8)
    cbar_ax = fig.add_axes([0.85, 0.15, 0.05, 0.7])
    fig.colorbar(ax, cax=cbar_ax)
    plt.show()
```



Part D

```
In [12]: zero_crossings = np.zeros(1000)
for i in np.arange(1000):
    for edge in Weighted_Graph.edges:
        zero_crossings[i] += int(vecs[edge[0], i] * vecs[edge[1], i] < 0)

plt.plot(np.arange(1000), zero_crossings)
    plt.xlabel("Eigenvector of $L_N$")
    plt.ylabel("Number of zero crossings")
    plt.show()</pre>
```



Part E & F

```
In [13]:
          sigmas = np.array([1, 0.5, 0.05])
           f, axes = plt.subplots(2, 3, figsize = (12, 8), sharey='row')
           for idx, sigma in enumerate(sigmas):
             signal = [np.exp(-np.linalg.norm(point[0])**2 / sigma) for point in zip(points)]
             g = [np.dot(signal, vec) for vec in vecs.T]
             axes[0, idx].set_title("Fourier Domain ($\sigma$ = " + str(sigma) + ")")
             axes[0, idx].set_ylabel("Fourier coefficient of eigenvector" if idx == 0 else "")
             axes[0, idx].plot(np.arange(len(g)), g, c='C0')
             axes[1, idx].set_title("Graph Signal ($\sigma$ = " + str(sigma) + ")")
             im = axes[1, idx].scatter(points[:, 0], points[:, 1], c=signal, cmap='bwr', s=10, vmin
           =0., vmax=1.)
           plt.subplots adjust(hspace = 0.3)
           fig.colorbar(im, ax=axes[1, 2], pad = 0.025)
           plt.show()
                    Fourier Domain (\sigma = 1.0)
                                                     Fourier Domain (\sigma = 0.5)
                                                                                      Fourier Domain (\sigma = 0.05)
           Fourier coefficient of eigenvector
              15
              10
               5
               0
                      200
                           400
                                600
                                     800
                                         1000
                                                       200
                                                            400
                                                                 600
                                                                      800
                                                                          1000
                                                                                         200
                                                                                             400
                                                                                                  600
                                                                                                       800
                                                                                                            1000
                     Graph Signal (\sigma = 1.0)
                                                       Graph Signal (\sigma = 0.5)
                                                                                     Graph Signal (\sigma = 0.05)
                                                                                                              1.0
             1.00
             0.75
                                                                                                              8.0
             0.50
             0.25
                                                                                                              0.6
             0.00
                                                                                                             0.4
            -0.25
            -0.50
                                                                                                              0.2
            -0.75
            -1.00
```

-0.5

0.0

0.5

-1.0

-0.5

0.0

0.5

1.0

-1.0

-0.5

0.0

0.5

Part G & H

```
In [14]: sigma = 0.5
         signal = [np.exp(-np.linalg.norm(point[0])**2 / sigma) for point in zip(points)]
         signal_tilde = signal + np.random.normal(scale=0.1, size=len(signal))
         errors = np.array([])
         for tau in np.arange(10, step=0.05):
           g_hat = [np.dot(signal_tilde, vec)/(1+tau*vals[idx]) for idx, vec in enumerate(vecs.T
         )]
           s_hat = np.zeros(shape=(1000,))
           for idx in np.arange(1000):
             s_hat += g_hat[idx]*vecs[:, idx]
           error = np.linalg.norm(signal - s_hat)
           errors = np.append(errors, error)
         plt.plot(np.arange(10, step=0.05), errors)
         plt.title("Reconstruction Error ($\sigma$ = " + str(sigma) + ")")
         plt.xlabel("$\\tau$")
         plt.ylabel("$||s - hat{s}||$")
         plt.show()
```

Reconstruction Error ($\sigma = 0.5$)

