

Machine Learning Homework 7

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a. code with detailed explanations

Kernel Eigenfaces

Part 1.

- PCA

I use SVD to solve the original eigen problem $Sw = \lambda w$, where $S = \frac{1}{N} \sum_x (x - \bar{x})(x - \bar{x})^T$.

Steps:

1. Compute centered training data X_{centered} .
2. Use SVD to solve for the centered training data, $X_{\text{centered}} = U \times S \times V^T$.
3. $V^T_{1...n}$ is what we want.

```
def PCA(X, output_dim=2):
    X_mean = np.mean(X, axis=0)
    X_centered = X - X_mean
    U, S, VT = np.linalg.svd(X_centered, full_matrices=False,
compute_uv=True)
    X_projected = U[:, :output_dim] * S[:output_dim]

    return X_projected, VT[:output_dim], X_mean
```

- LDA

I follow the formula from the slides.

Steps:

1. Compute $S_W = \sum_{j=1}^k S_j$, where $S_j = \sum_{i \in C_j} (x_i - m_j)(x_i - m_j)^T$ and $m_j = \frac{1}{n_j} \sum_{i \in C_j} x_i$.
2. Compute $S_B = \sum_{j=1}^k S_{B_j} = \sum_{j=1}^k n_j (m_j - m)(m_j - m)^T$ where $m = \frac{1}{n} \sum x$.
3. Solve the eigen problem $S_W^{-1} S_B w = \lambda w$.
4. The resulting $w^{1...n}$ is what we want.

```
def LDA(X, y, output_dim=2):
    n_feat = X.shape[1]
    n_class = len(np.unique(y))
    Sw = np.zeros((n_feat, n_feat))
    Sb = np.zeros((n_feat, n_feat))
    X_mean = np.mean(X, axis=0)

    for c in range(n_class):
        mask = (y == c+1)
```

```

mean_c = np.mean(X[mask], axis=0)
dw = X[mask] - mean_c
Sw += dw.T @ dw
db = (mean_c - X_mean).reshape(-1, 1)
Sb += np.sum(mask) * db @ db.T

A = np.linalg.pinv(Sw) @ Sb
eigenvalue, eigenvector = np.linalg.eigh(A)
W = eigenvector[:, :-1]
X_projected = X @ W[:, :output_dim]

return X_projected, W[:, :output_dim].T, X_mean

```

- Eigenfaces I directly take the eigenvectors corresponding to the top 25 largest eigenvalues from solving PCA.

```

X_train_projected_pca, X_eigenfaces, X_train_mean = PCA(X_train,
output_dim=100)
fig, axes = plt.subplots(5, 5)
for idx, ax in enumerate(axes.flatten()):
    ax.imshow(X_eigenfaces[idx].reshape(H, W), cmap='gray')

```

- Fisherfaces

I follow the steps from [Belhumeur, Hespanha and Kriegman](#).

First, I apply PCA to the data, and then apply LDA to the PCA-projected data. Take the eigenvectors corresponding to the top 25 largest eigenvalues from solving PCA and LDA $W_{\{opt\}} = W_{\{LDA\}}^T W_{\{PCA\}}$

```

def fisherface(X, y, output_dim=2):
    n_feat = X.shape[1]
    n_class = len(np.unique(y))
    X_projected_pca, W_pca, _ = PCA(X, output_dim=n_feat-n_class)
    X_projected_lda, W_lda, _ = LDA(X_projected_pca, y,
output_dim=output_dim)
    W_opt = W_lda @ W_pca

    return X_projected_lda, W_opt

X_train_projected_lda, X_fisherfaces = fisherface(X_train, y_train,
output_dim=100)
fig, axes = plt.subplots(5, 5)
for idx, ax in enumerate(axes.flatten()):
    ax.imshow(X_fisherfaces[idx].reshape(H, W), cmap='gray')

```

- Reconstruction

Use the eigenvectors w we solved from PCA or LDA, multiply it to the projected data $X_{\{projected\}} = X_{\{centered\}} w$, so that $X_{\{reconstruct\}} = X_{\{projected\}} w^T$.

```

reconstruction_pca = X_train_projected_pca @ X_eigenfaces + X_train_mean
fig, axes = plt.subplots(2, 5)
for idx, ax in enumerate(axes.flatten()):
    ax.imshow(reconstruction_pca[random_idx[idx]].reshape(H, W),
              cmap='gray')

reconstruction_lda = X_train_projected_lda @ X_fisherfaces + X_train_mean
fig, axes = plt.subplots(2, 5)
for idx, ax in enumerate(axes.flatten()):
    ax.imshow(reconstruction_lda[random_idx[idx]].reshape(H, W),
              cmap='gray')

```

Part 2.

- KNN

Here is my implementation of k nearest neighbors, which use euclidean distance as criterion.

```

def KNN(X_test, X_train, y_test, y_train, k=5):
    dist = cdist(X_test, X_train, 'euclidean')
    sorted_index = np.argsort(dist, axis=1)
    k_nearest = sorted_index[:, :k]
    y_pred = np.zeros(len(X_test))

    for i in range(len(X_test)):
        unique, counts = np.unique(y_train[k_nearest[i]],
                                   return_counts=True)
        y_pred[i] = unique[np.argmax(counts)]

    acc = np.sum(y_pred == y_test) / len(y_test)

    return y_pred, acc

```

- Face Recognition

To do face recognition, I first projected testing data to the space same as training data by the eigenfaces and fisherfaces, and then use KNN to classify testing data.

```

X_test_mean = np.mean(X_test, axis=0)
X_test_centered = X_test - X_test_mean
X_test_projected_pca = X_test_centered @ X_eigenfaces.T
y_pred_pca, acc = KNN(X_test_projected_pca, X_train_projected_pca, y_test,
                      y_train, k=5)
print(acc)

X_test_projected_lda = X_test_centered @ X_fisherfaces.T
y_pred_lda, acc = KNN(X_test_projected_lda, X_train_projected_lda, y_test,
                      y_train, k=5)
print(acc)

```

Part 3.

- RBF kernel

```
def RBF_kernel(X, X_, gamma):  
    return np.exp(-gamma * cdist(X, X_, 'sqeuclidean'))
```

- Polynomial kernel

```
def polynomial_kernel(X, X_, gamma, coef, degree):  
    return np.power(gamma * X @ X_.T + coef, degree)
```

- Kernel PCA

I follow the formula from the slides.

For centering in high dimensional space, I follow the formula from [Schölkopf, Smola and Müller](#)
Steps:

1. Given similarity matrix K , compute $K_{\text{centered}} = K - \frac{1}{N} K - K \frac{1}{N} + \frac{1}{N} K \frac{1}{N}$, where $\frac{1}{N} = \text{diag}(1/N)$
2. Solve the eigen problem $K_{\text{centered}} w = \lambda w$
3. $w^{1\dots n}$ is what we want.

```
def kernel_PCA(K, output_dim=2):  
    oneN = np.diag(np.full(len(K), 1./len(K)))  
    K_centered = K - oneN @ K - K @ oneN + oneN @ K @ oneN  
    eigenvalue, eigenvector = np.linalg.eig(K_centered)  
    a = eigenvector[:, np.argsort(eigenvalue)[::-1]]  
    K_projected = K_centered @ a[:, :output_dim]  
  
    return K_projected, a[:, :output_dim].T, K_centered
```

- Kernel LDA I follow the formula from [Ghojogh, Karray and Crowley](#). Steps:

1. Compute $N = \sum_{j=1}^c K_j H_j K_j^T$, where $H_j = I - \frac{1}{n_j} \mathbf{1} \mathbf{1}^T$.
2. Compute $M = \sum_{j=1}^c c_j (m_j - m)(m_j - m)^T$.
3. Solve the eigen problem $N^{-1} M w = \lambda w$.
4. $w^{1\dots n}$ is what we want.

```
def kernel_LDA(K, y, output_dim=2):  
    y_onehot = OneHotEncode(y)  
    m_classes = y_onehot.T @ K / np.sum(y_onehot, axis=0)[:, np.newaxis]  
    indices = y - 1  
    N = K @ (K - m_classes[indices])  
    # N += np.eye(len(K)) * 1e-8  
    m_classes_centered = m_classes - np.mean(K, axis=1)
```

```

M = m_classes_centered.T @ m_classes_centered
A = np.linalg.pinv(N) @ M
eigenvalue, eigenvector = np.linalg.eigh(A)
a = eigenvector[:, ::-1]
K_projected = K @ a[:, :output_dim]

return K_projected, a[:, :output_dim].T

```

- Kernel Eigenfaces

I directly take the eigenvectors corresponding to the top 25 largest eigenvalues from solving kernel PCA.

```

K_train_projected, X_kernel_eigenfaces, K_train_centered =
kernel_PCA(K_train, output_dim=100)

```

- Kernel Fisherfaces

Same as fisherfaces, first apply kernel PCA to training data and use kernel LDA to get the kernel fisherfaces. $W_{opt} = W_{kLDA}^T W_{kPCA}$

```

def kernel_fisherfaces(K, y, output_dim):
    n_feat = len(K)
    n_class = len(np.unique(y))
    K_projected_kpca, W_kpca, K_centered = kernel_PCA(K, output_dim=n_feat)
    K_projected_klda, W_klda = kernel_LDA(K_projected_kpca, y,
output_dim=output_dim)
    W_opt = W_klda @ W_kpca

    return K_projected_klda, W_opt

K_train_projected_klda, X_kernel_fisherfaces, _ = LDA(K_train_projected,
y_train, output_dim=100)
W_opt = X_kernel_fisherfaces @ X_kernel_eigenfaces

```

- Face Recognition Same as the non-kernel method, first projected similarity matrix of testing data to the spaces, and adopt KNN to classify.

```

K_test_projected = K_test @ X_kernel_eigenfaces.T
y_pred_kpca, acc = KNN(K_test_projected, K_train_projected, y_test,
y_train, k=5)
print(acc)

K_test_projected = K_test @ W_opt.T
y_pred_klda, acc = KNN(K_test_projected, K_train_projected, y_test,
y_train, k=5)
print(acc)

```

t-SNE

Part 1.

Here is how I modified the original tsne.py code to make it back to symmetric SNE, the commented code is the original t-SNE.

The only 2 different between tsne and symmetric SNE are the Q matrix, which is the low dimensional pairwise affinities, and of course the gradient would be different.

t-SNE:

$$q_{ij} = \frac{(1 + \|y_i - y_j\|^2)^{-1}}{\sum_{k \neq l} (1 + \|y_l - y_k\|^2)^{-1}}$$
$$\frac{\partial C}{\partial y_i} = 4 \sum_j (p_{ij} - q_{ij})(y_i - y_j)(1 + \|y_i - y_j\|^2)^{-1}$$

Symmetric SNE:

$$q_{ij} = \frac{\exp(-\|y_i - y_j\|^2)}{\sum_{k \neq l} \exp(-\|y_l - y_k\|^2)} \quad \frac{\partial C}{\partial y_i} = 2 \sum_j (p_{ij} - q_{ij})(y_i - y_j)$$

```
# Compute pairwise affinities
sum_Y = np.sum(np.square(Y), 1)
num = -2. * np.dot(Y, Y.T)
# num = 1. / (1. + np.add(np.add(num, sum_Y).T, sum_Y))
num = np.exp(-np.add(np.add(num, sum_Y).T, sum_Y))
num[range(n), range(n)] = 0.
Q = num / np.sum(num)
Q = np.maximum(Q, 1e-12)

# Compute gradient
PQ = P - Q
for i in range(n):
    # dY[i, :] = np.sum(np.tile(PQ[:, i] * num[:, i], (no_dims, 1)).T *
    (Y[i, :] - Y), 0)
    dY[i, :] = np.sum(np.tile(PQ[:, i], (no_dims, 1)).T * (Y[i, :] - Y), 0)
```

Part 2.

Every 20 iterations, I save the visualization of the 2D space, and use imagemagick to make all the pictures to gif.

```
if (iter + 1) % 20 == 0:
    pylab.title(f'iter:{iter+1:004}')
    pylab.scatter(Y[:, 0], Y[:, 1], 5, labels)
    pylab.savefig(f'./images/symmetric_sne/{iter+1:004}', dpi=100,
facecolor='white')
    pylab.close()
```

```
$ convert -delay 10 images/symmetric_sne/*.png symmetric_sne.gif
```

Part 3.

I defined a function called `plot_similarity` to visualize the distribution of pairwise similarities in both high dimensional space and low dimensional space, which is the P and Q matrix.

```
def plot_similarity(P, Q):  
    pylab.title("symmetric sne high dim")  
    pylab.hist(P.flatten(), bins=1000, log=True)  
    pylab.savefig("symmetric_sne_high_dim", dpi=100, facecolor='white')  
    pylab.close()  
  
    pylab.title("symmetric sne low dim")  
    pylab.hist(Q.flatten(), bins=1000, log=True)  
    pylab.savefig("symmetric_sne_low_dim", dpi=100, facecolor='white')  
    pylab.close()  
  
plot_similarity(P, Q)
```

Part 4.

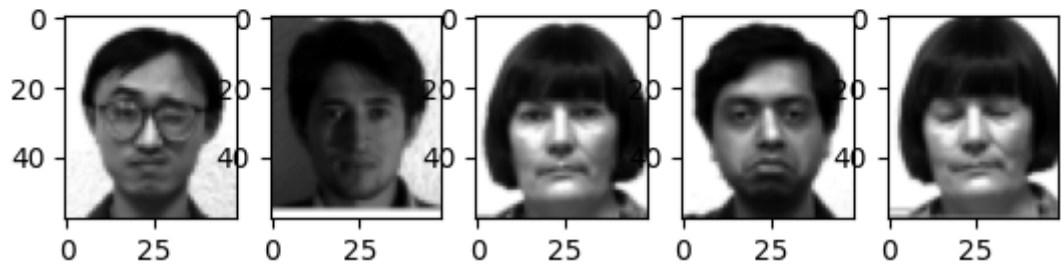
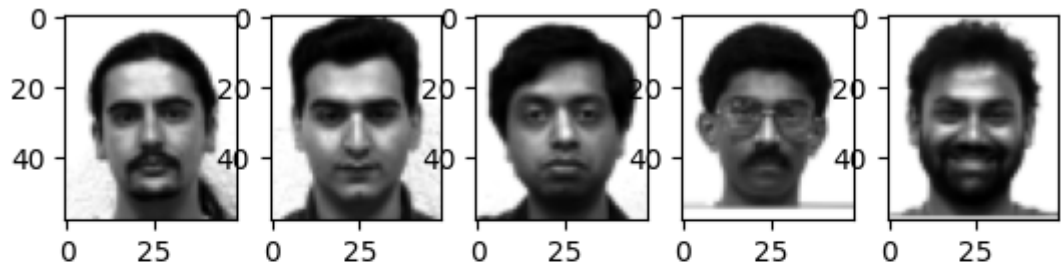
I directly modified the last parameter of the tsne function.

```
Y, P, Q = tsne(X, labels, 2, 50, 5.0)
```

b. experiments settings and results & discussions

Kernel Eigenfaces

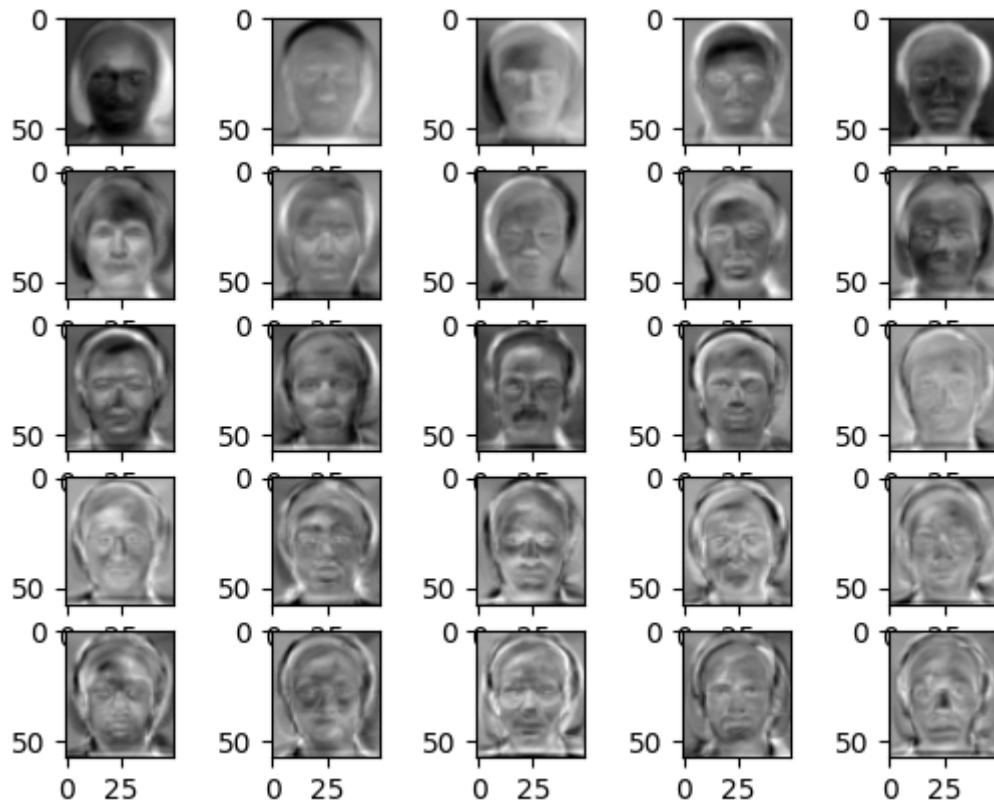
Part 1.



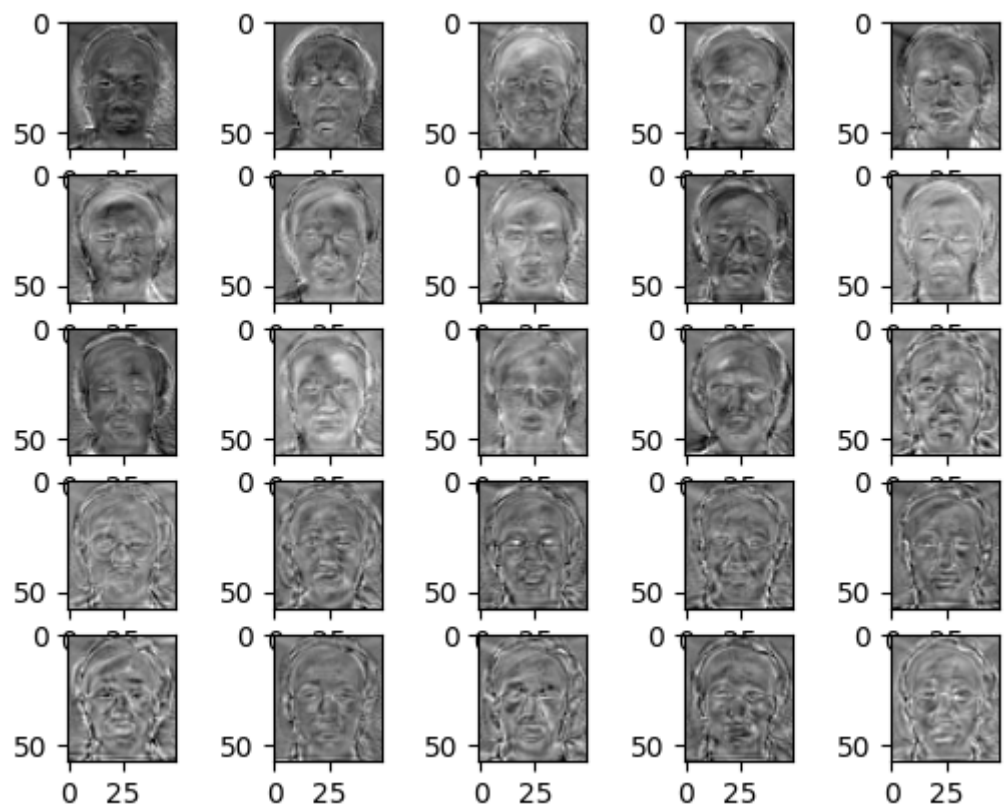
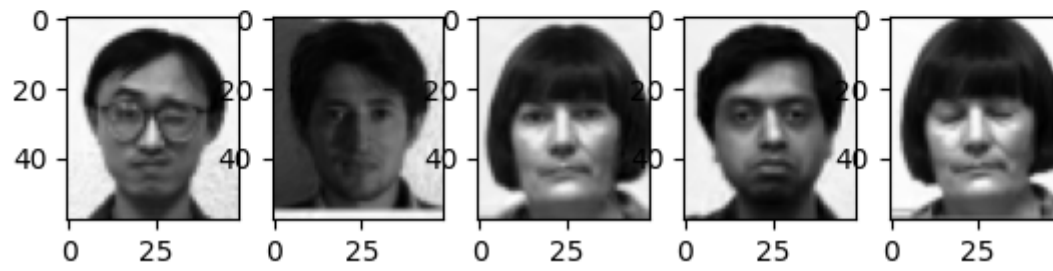
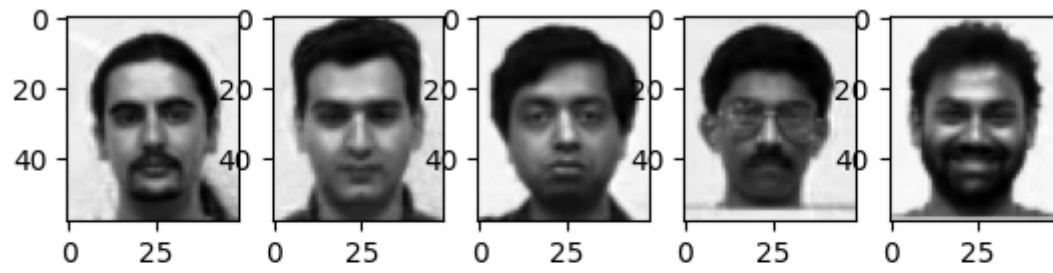
• Original:

• PCA:

Eigenfaces:

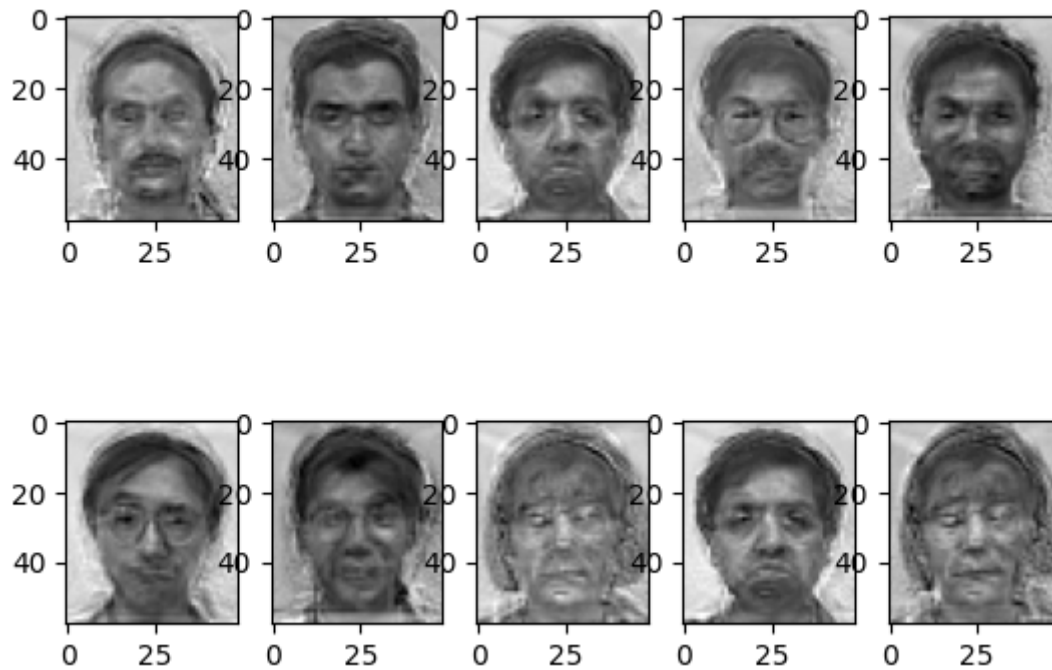


Reconstruction:



- LDA: Fisherfaces:

Reconstruction:



Part 2.

Both PCA and LDA project training data to 100-dim space, and k in KNN is choose to be 5.

- PCA: accuracy: 0.8667
- LDA: accuracy: 0.9667

Part 3.

Both kernel PCA and kernel LDA project training data to 100-dim space, and k in KNN is choose to be 5.

For RBF kernel, the $\gamma=0.01$.

For Polynomial kernel, $\gamma=0.01$, $\text{coef0}=1$, $\text{degree}=2$.

- kernel PCA:
 - RBF kernel: accuracy: 0.8333
 - Polynomial kernel: accuracy: 0.7333
- kernel LDA:
 - RBF kernel: accuracy: 0.8333
 - Polynomial kernel: accuracy: 0.8

Comparing to non-kernel method, the performance of kernel PCA and kernel LDA are both worse than PCA and LDA. And LDA got the best results in this experiment settings.

t-SNE

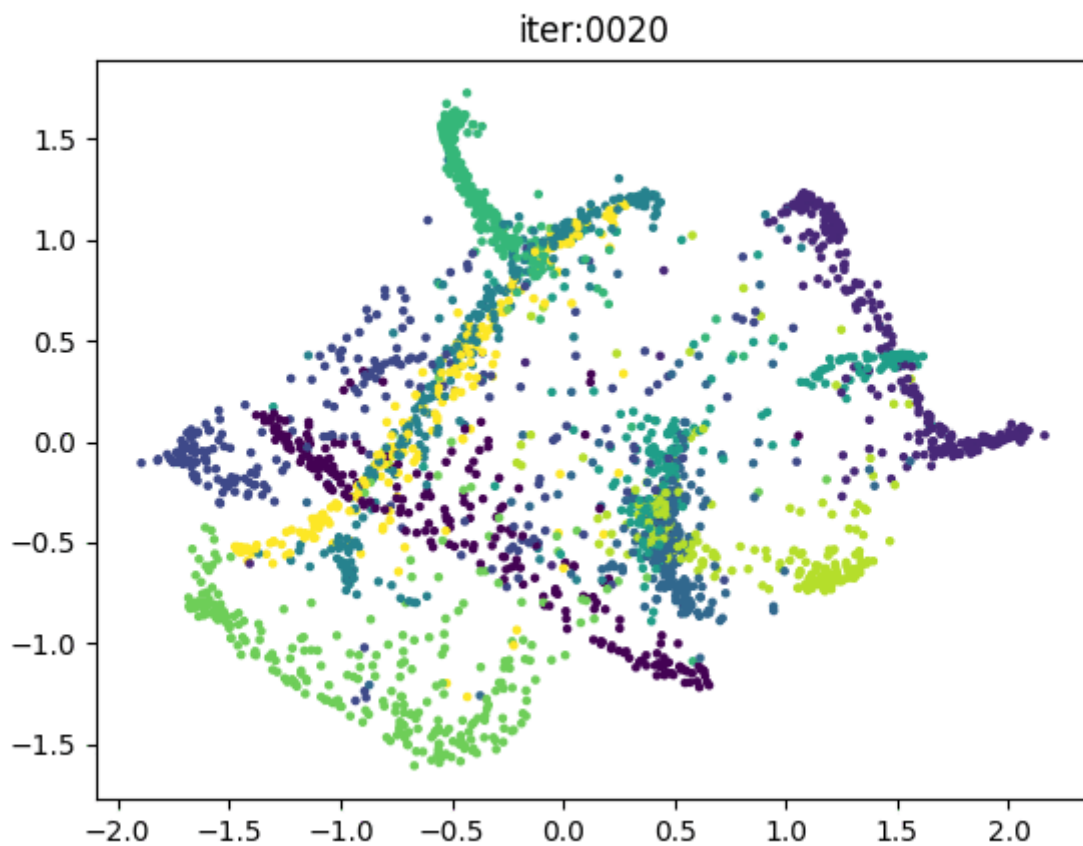
Part 1.

- t-SNE The low dimension distribution of pairwise similarities follows Student's T distribution.
- Symmetric SNE The low dimension distribution of pairwise similarities follows Gaussian distribution.

Since the Student's T distribution is a long-tail distribution compared to Gaussian distribution, it can solve the crowded problem that appears in Symmetric SNE. (The distance between two points has to be larger to achieve the same probability).

Part 2.

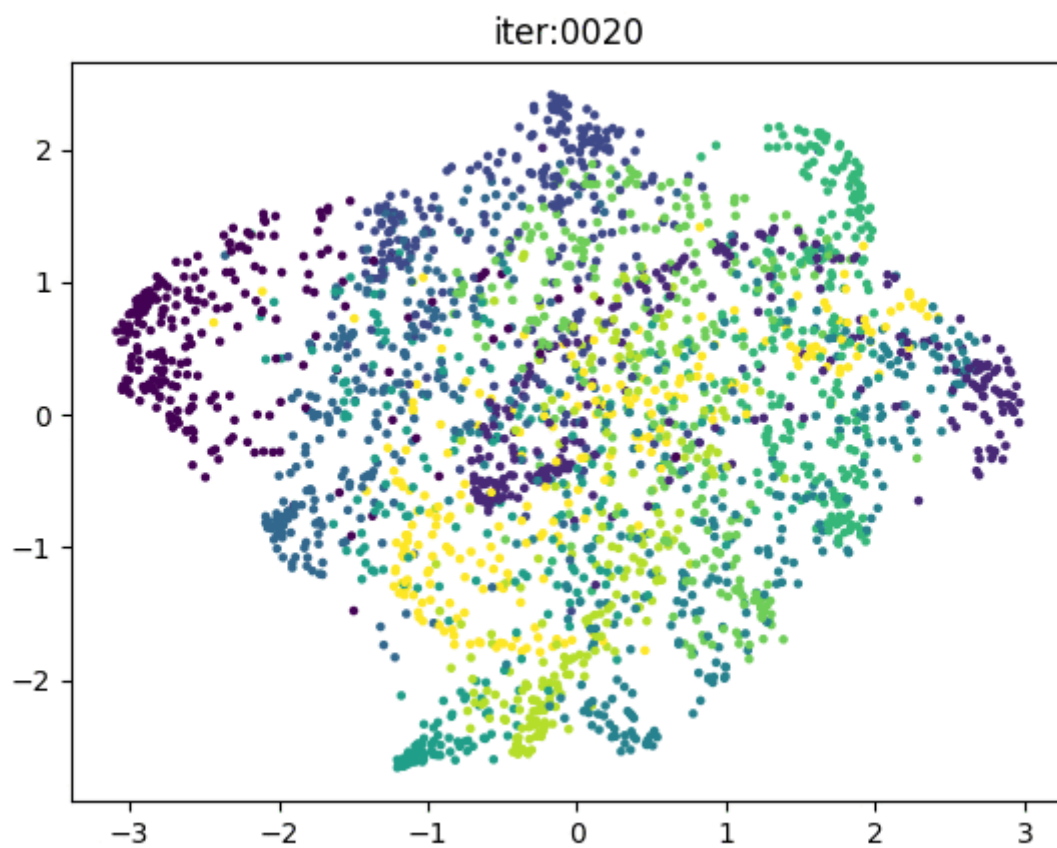
- t-SNE: gif:



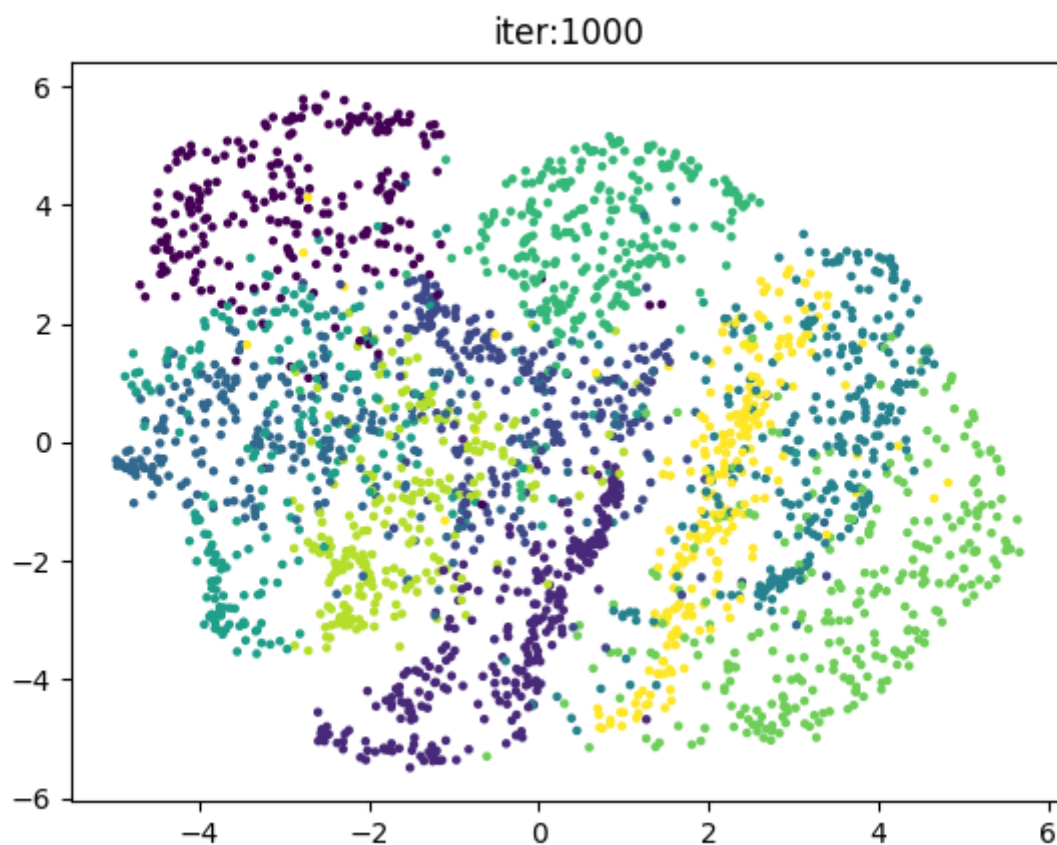
last iteration:



- Symmetric SNE: gif:



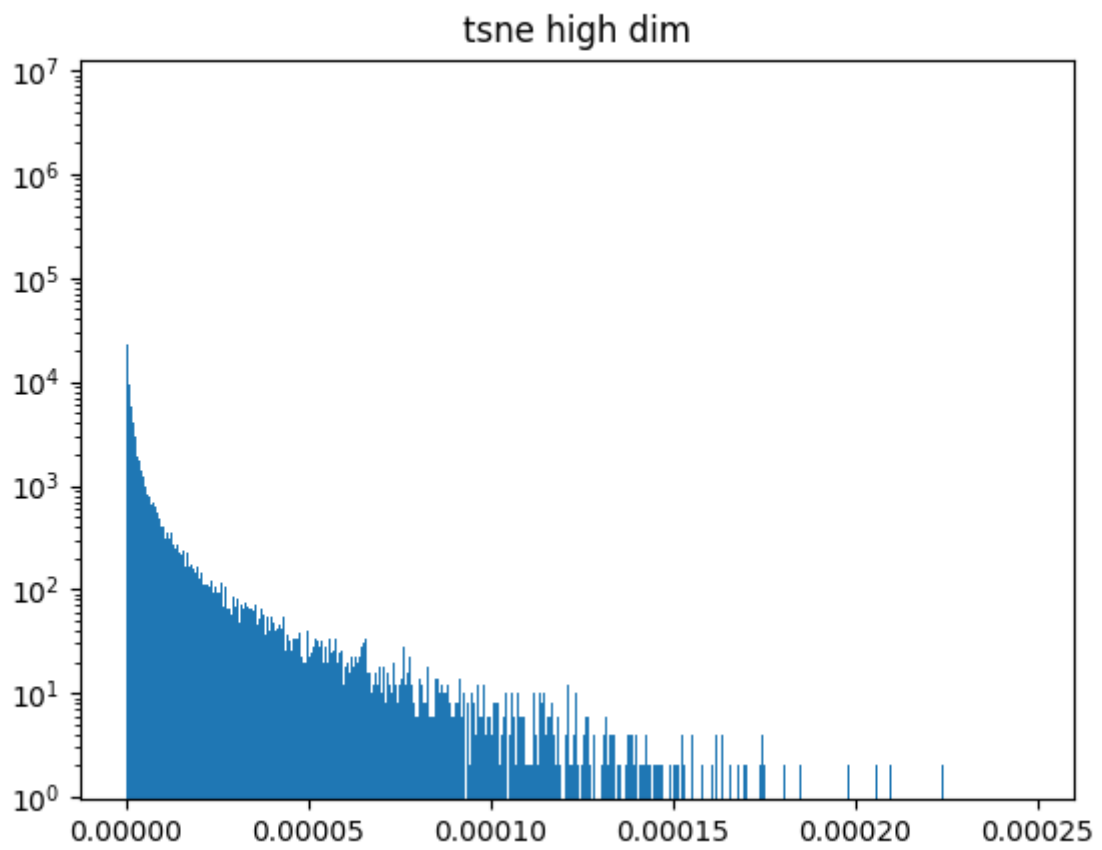
last iteration:



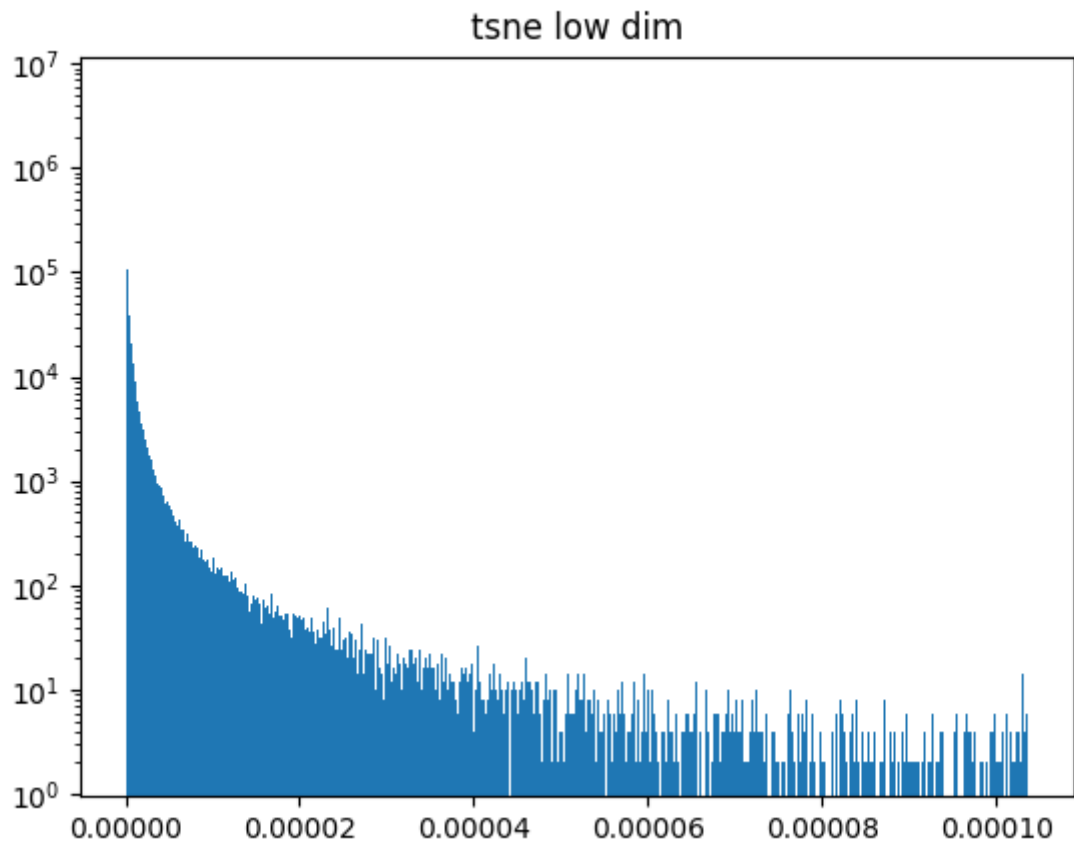
We can observe the clustering result of Symmetric SNE is more crowded.

Part 3.

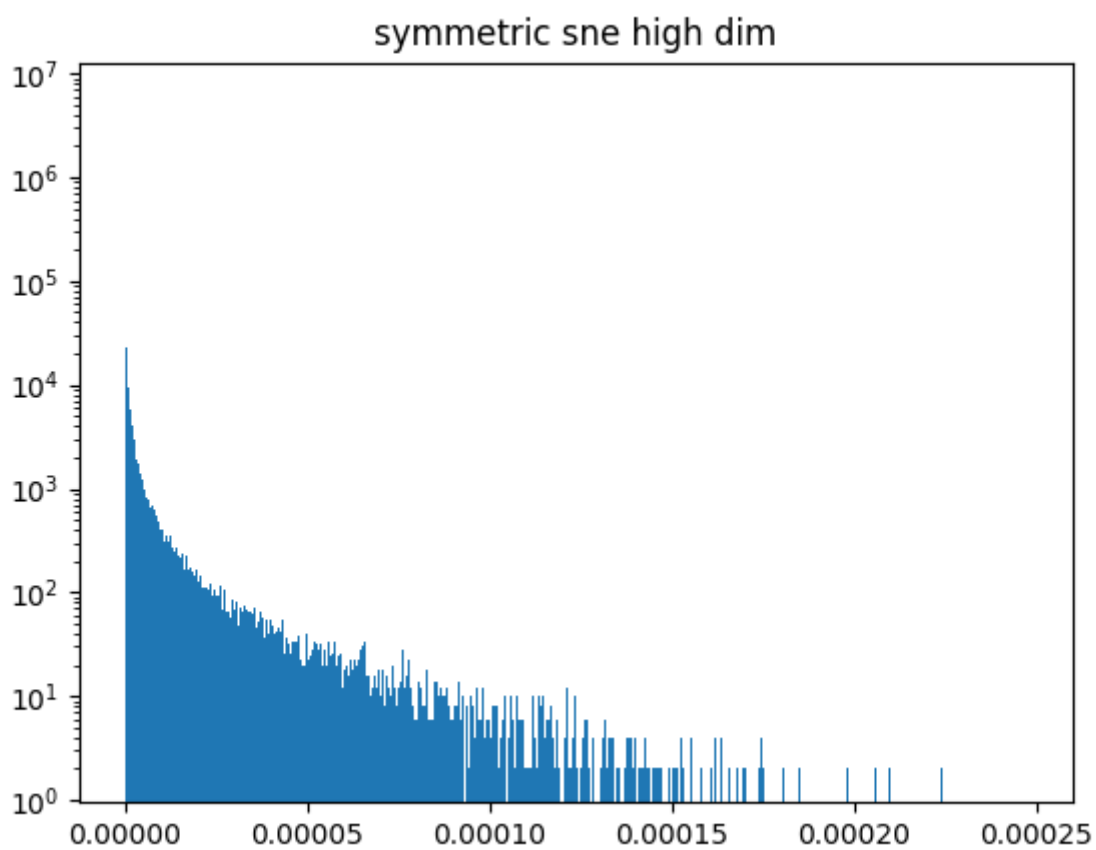
- t-SNE: High dimension:



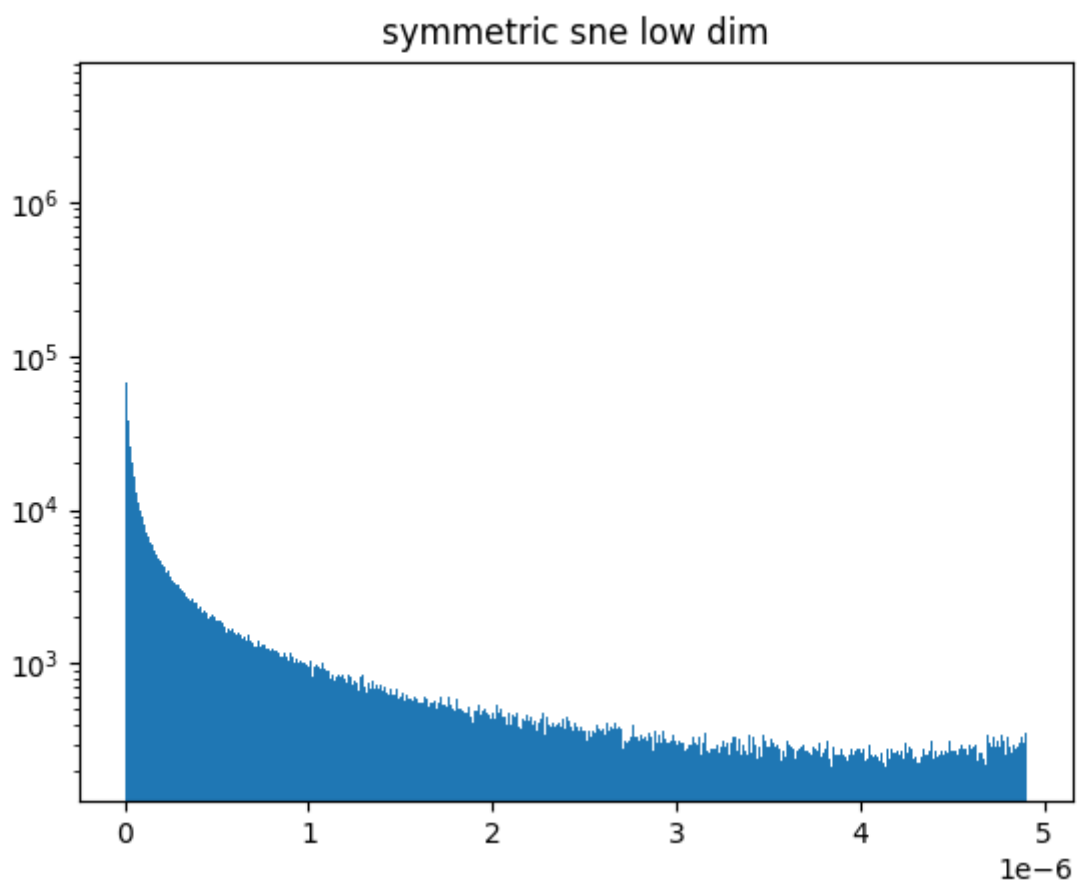
Low dimension:



- Symmetric SNE:
High dimension:



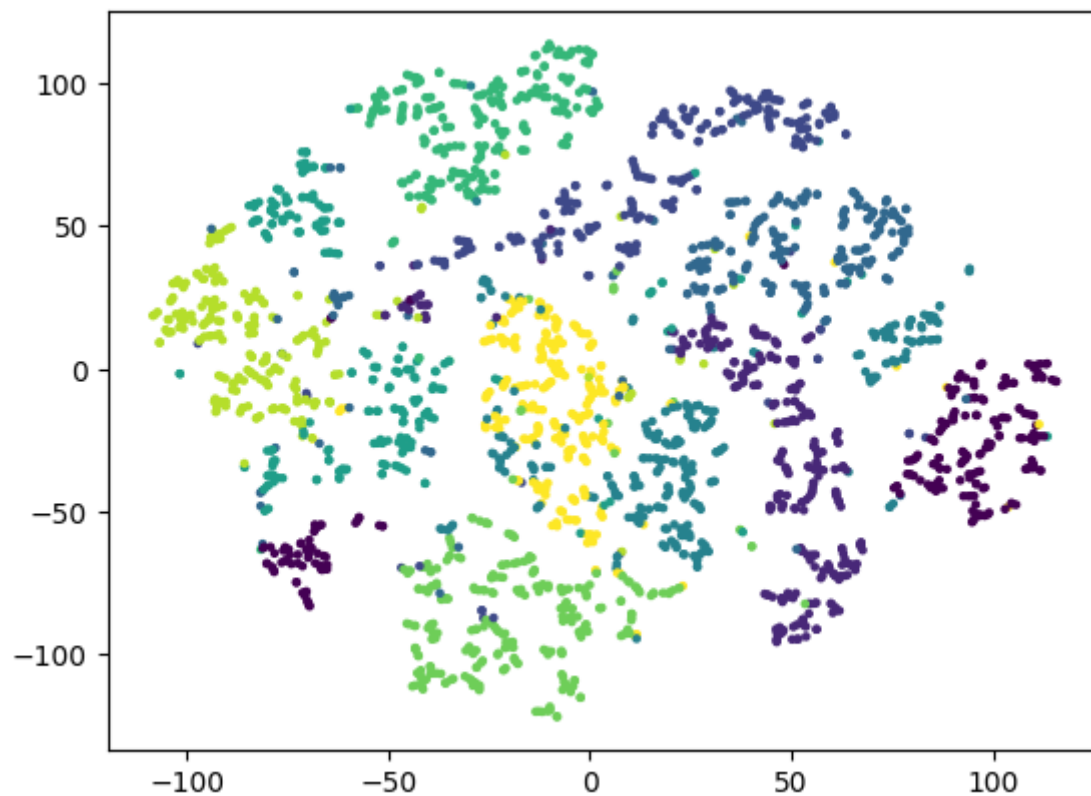
Low dimension:



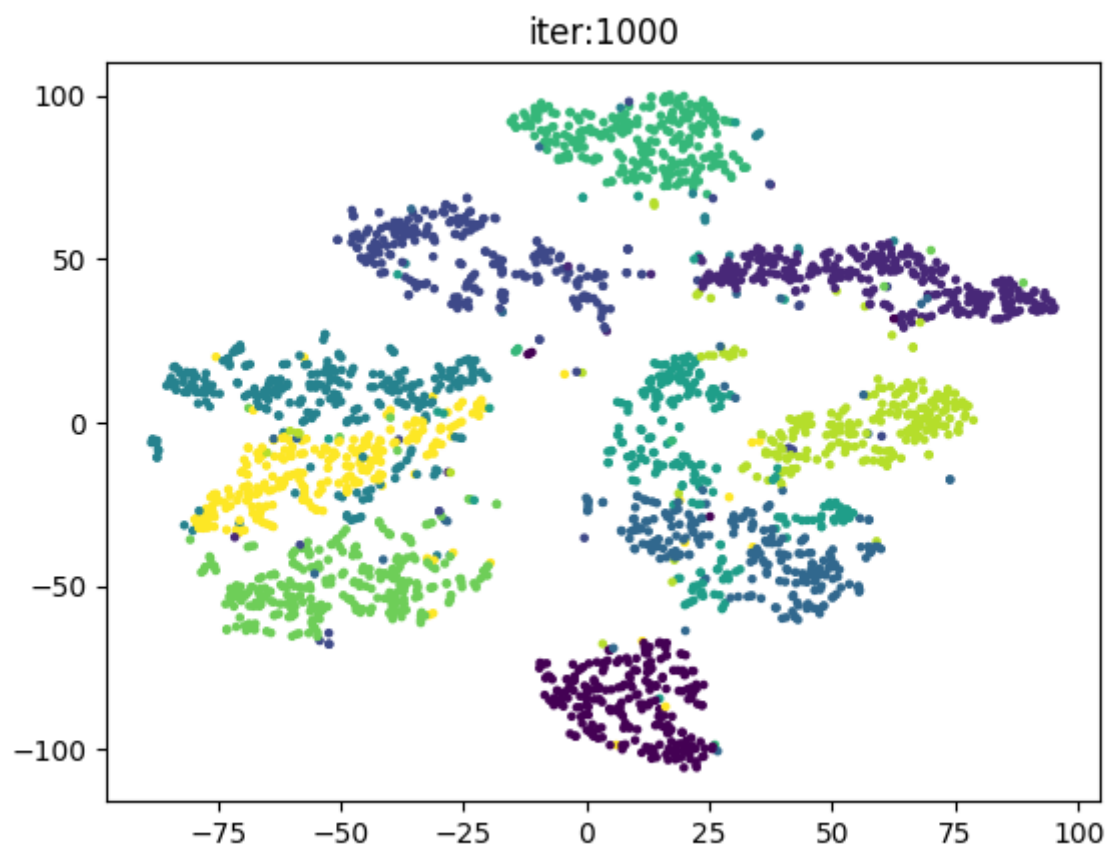
We can observe the range of x in low dimension distribution of Symmetric SNE is smaller than that of t-SNE, hence results in the crowded problem.

Part 4.

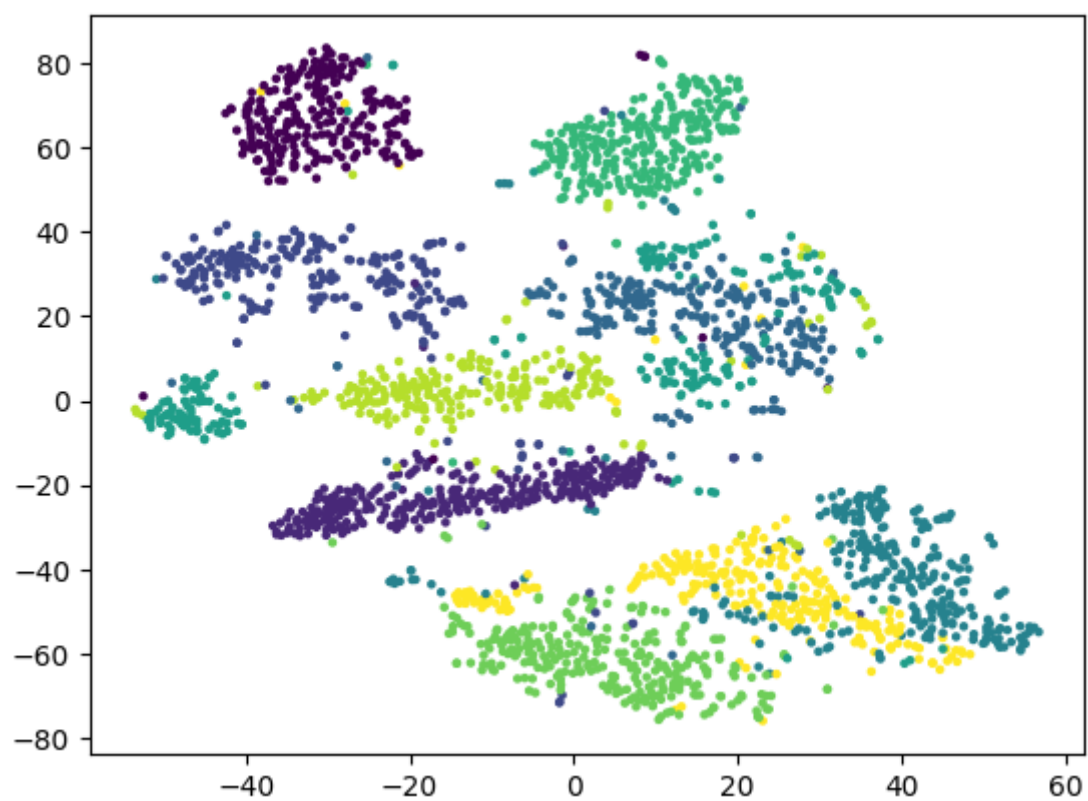
- t-SNE:
perplexity=5



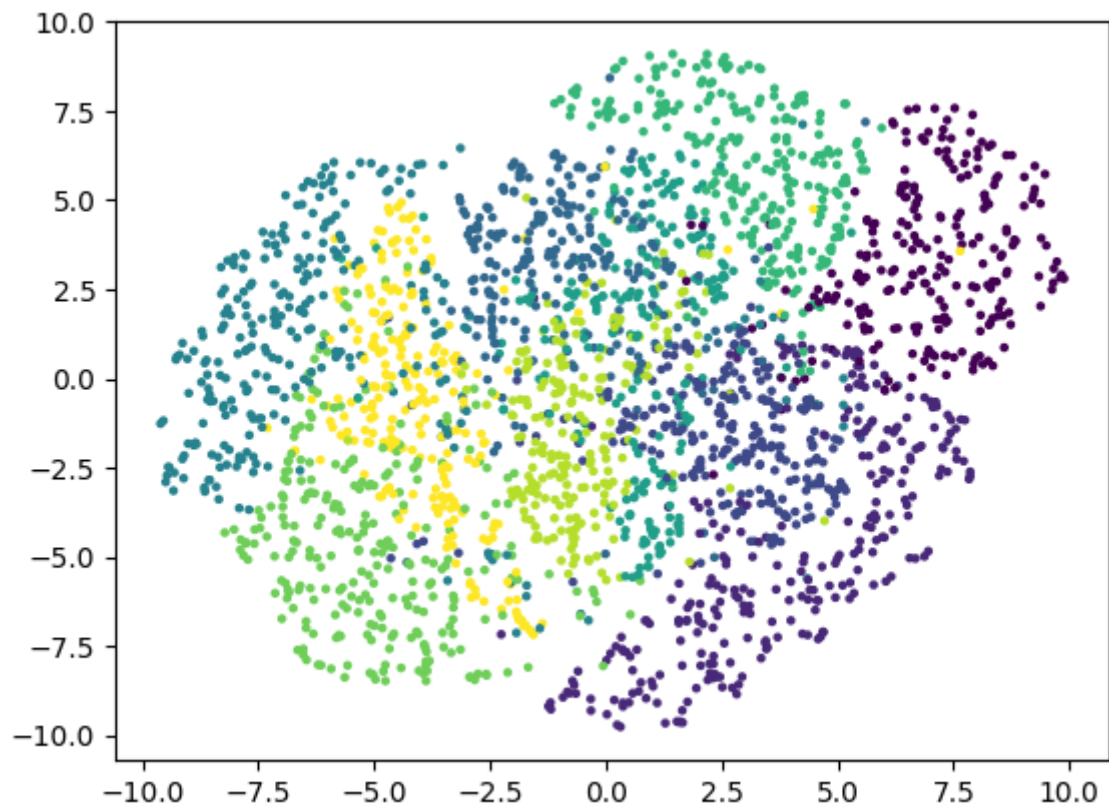
perplexity=20



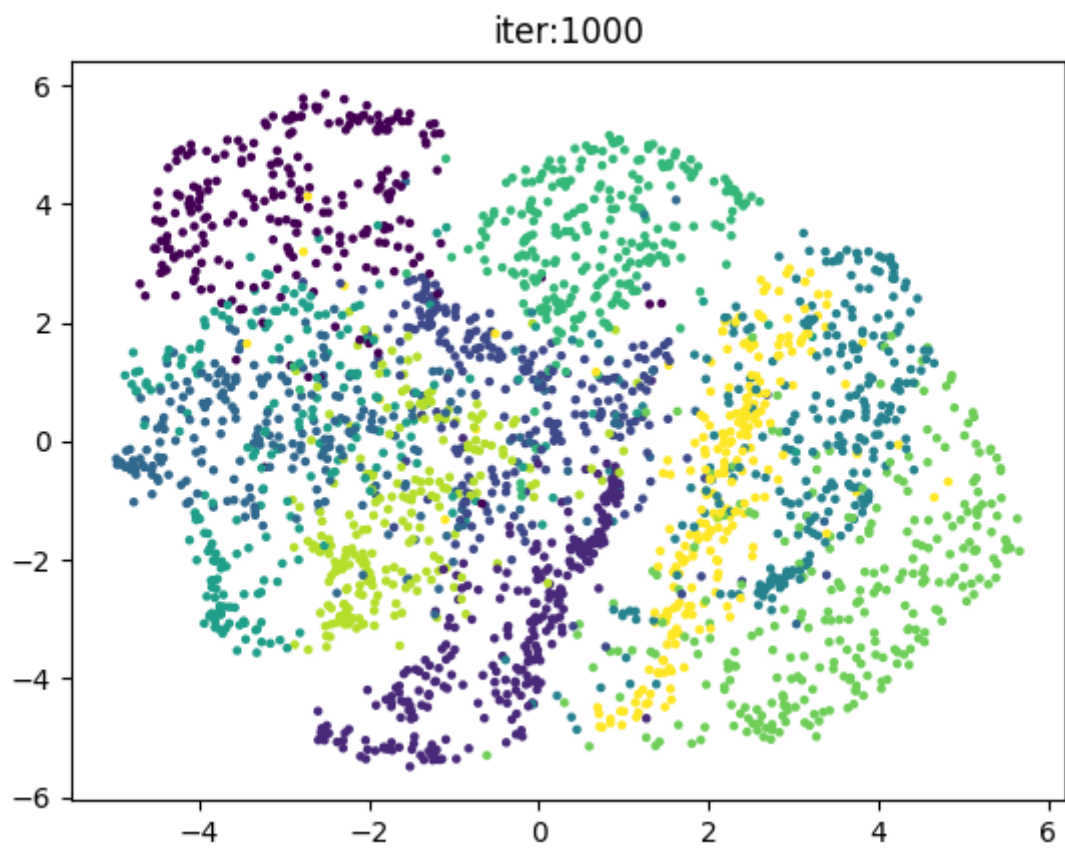
perplexity=50

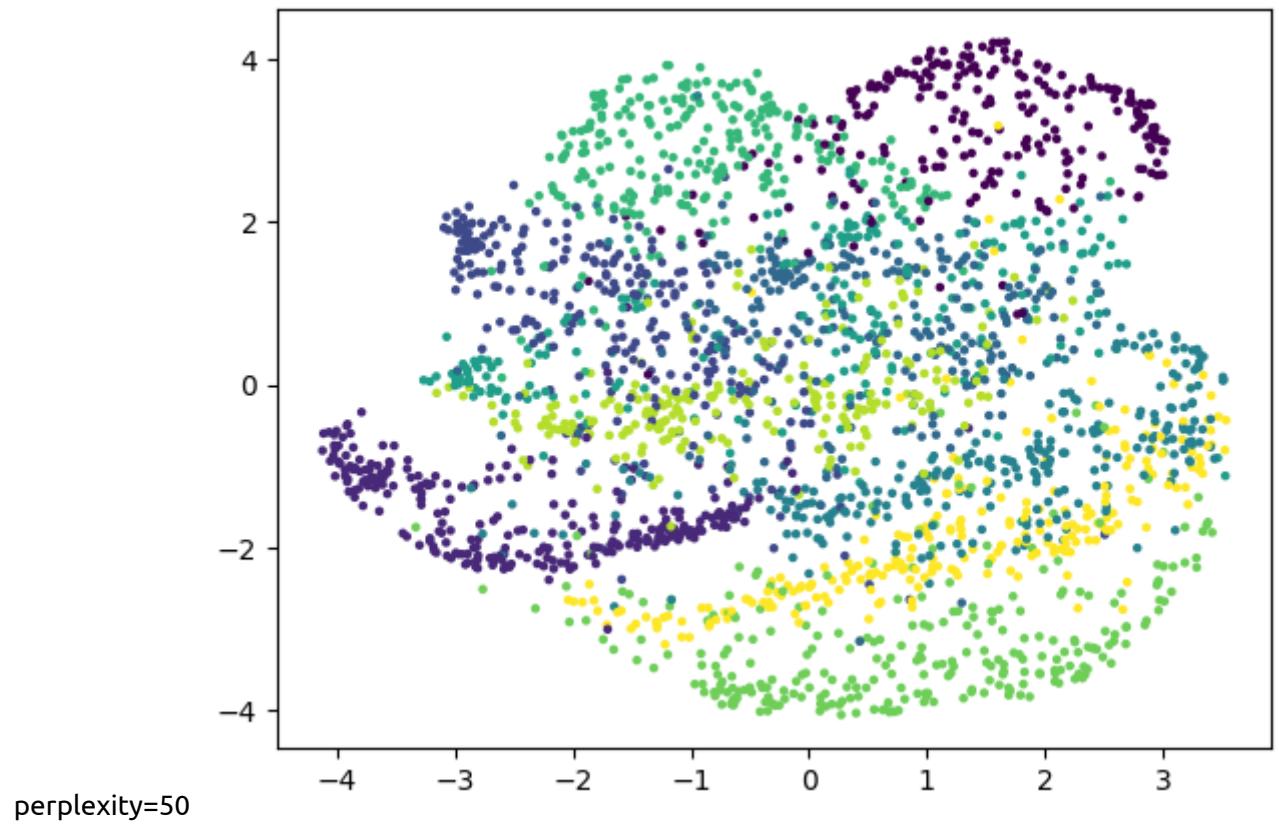


- Symmetric SNE: perplexity=5



perplexity=20





We can observe that with higher perplexity, the overall structure is more obvious, however, it may be harder to distinguish between every clusters. With lower perplexity, only a few neighbors are influential, which may results in one group to be split into mutilple groups.