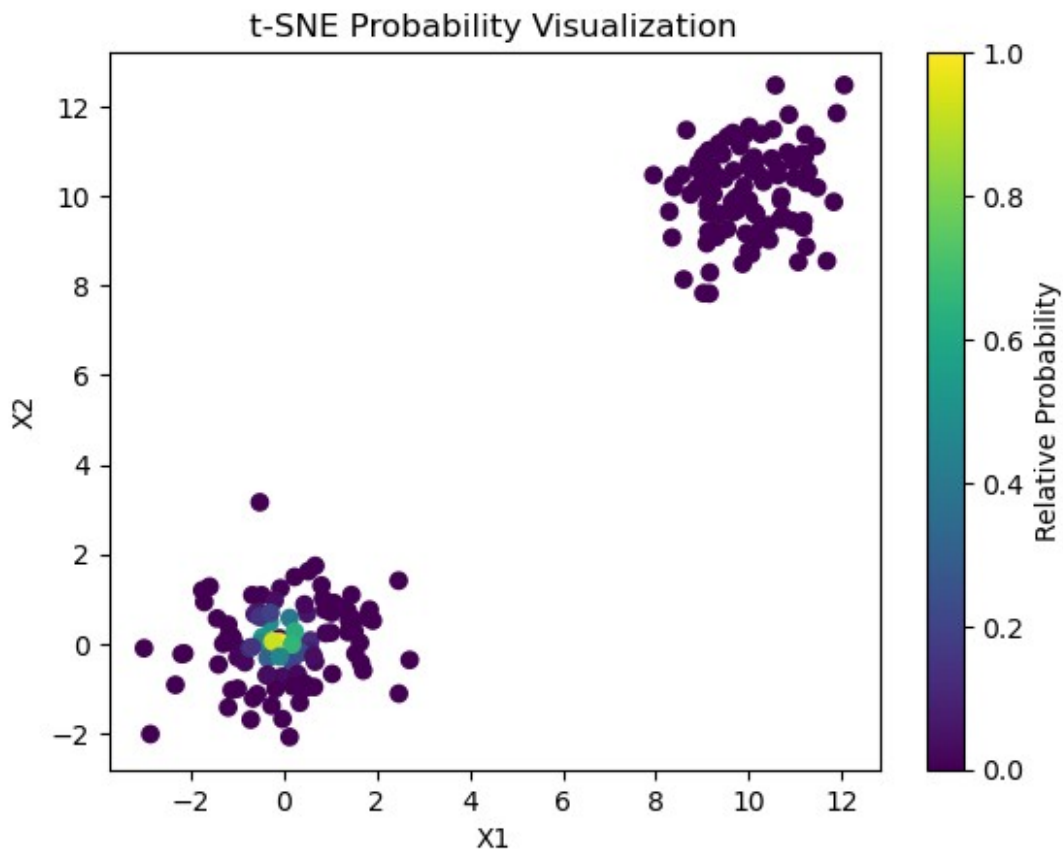



```
plot_probabilities(data, pij_matrix_sigma_2)

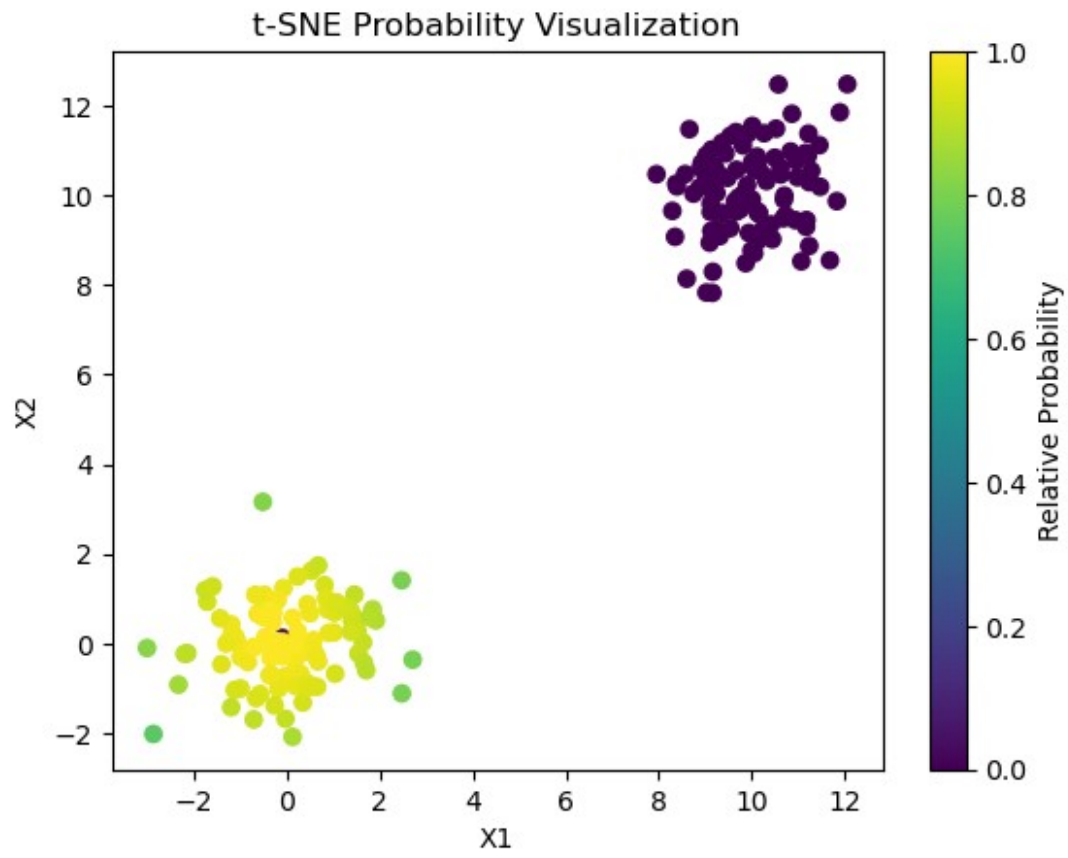
print("The value for sigma^2 = 100")
plot_probabilities(data, pij_matrix_sigma_3)

print("The value for sigma^2 = 1")
plot_probabilities(data, pij_matrix_sigma_4)

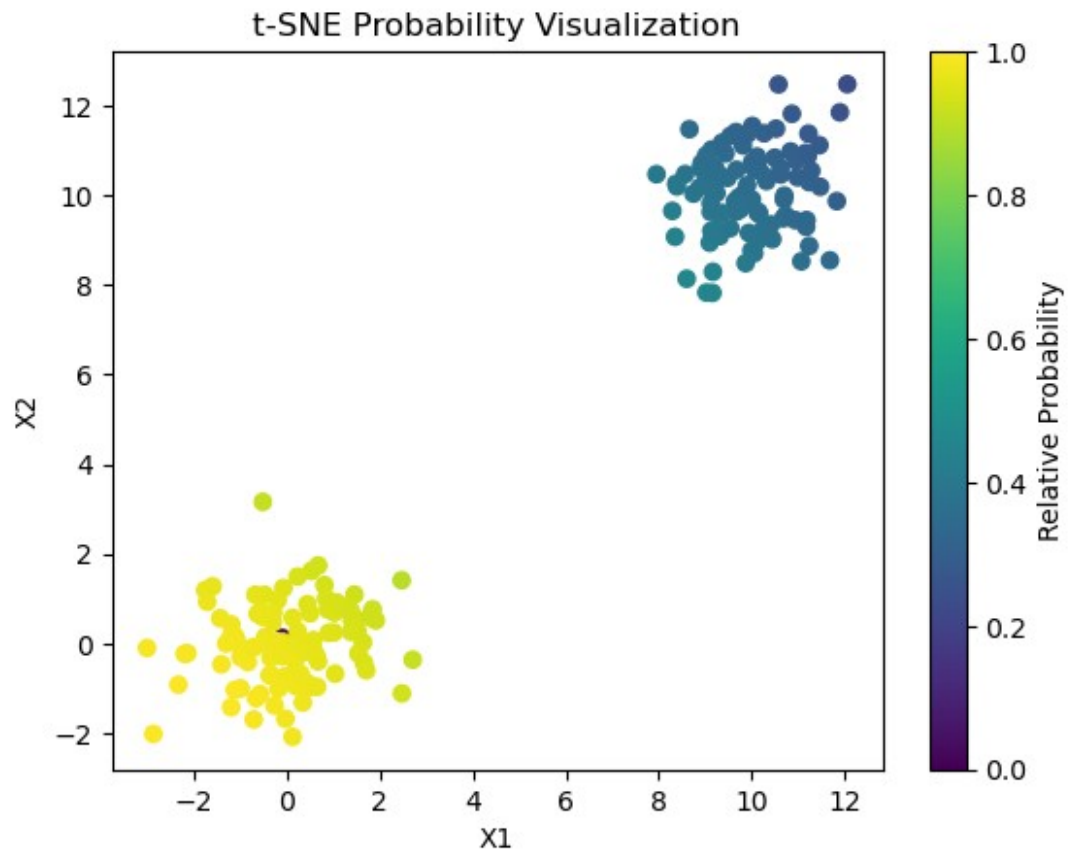
The value for sigma^2 = 0.1
```



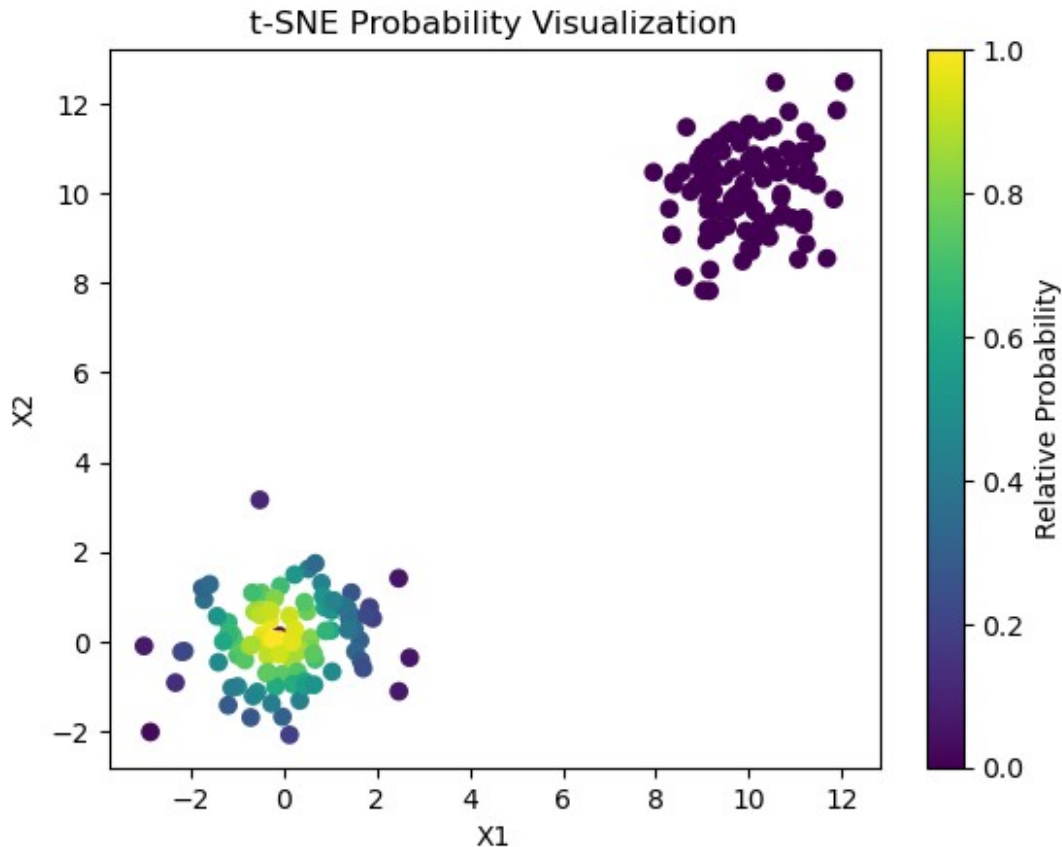
The value for sigma^2 = 10



The value for $\sigma^2 = 100$



The value for $\sigma^2 = 1$



```
def compute_qij(y):
    n = y.shape[0]
    qij = np.zeros((n, n))

    for i in range(n):
        for j in range(n):
            if i != j:
                qij[i, j] = (1 + np.linalg.norm(y[i] - y[j])**2)**-1

    qij /= np.sum(qij)

    return qij

y = data # Using yi = xi
qij_matrix = compute_qij(y)
# print(qij_matrix)

[[0.00000000e+00 3.67091796e-05 1.27288028e-04 ... 7.14311664e-07
 8.50790570e-07 6.50990816e-07]
 [3.67091796e-05 0.00000000e+00 5.09964700e-05 ... 8.58208464e-07
 9.80335848e-07 7.58573976e-07]
 [1.27288028e-04 5.09964700e-05 0.00000000e+00 ... 7.32793199e-07
 8.62331998e-07 6.63278281e-07]]
```

```

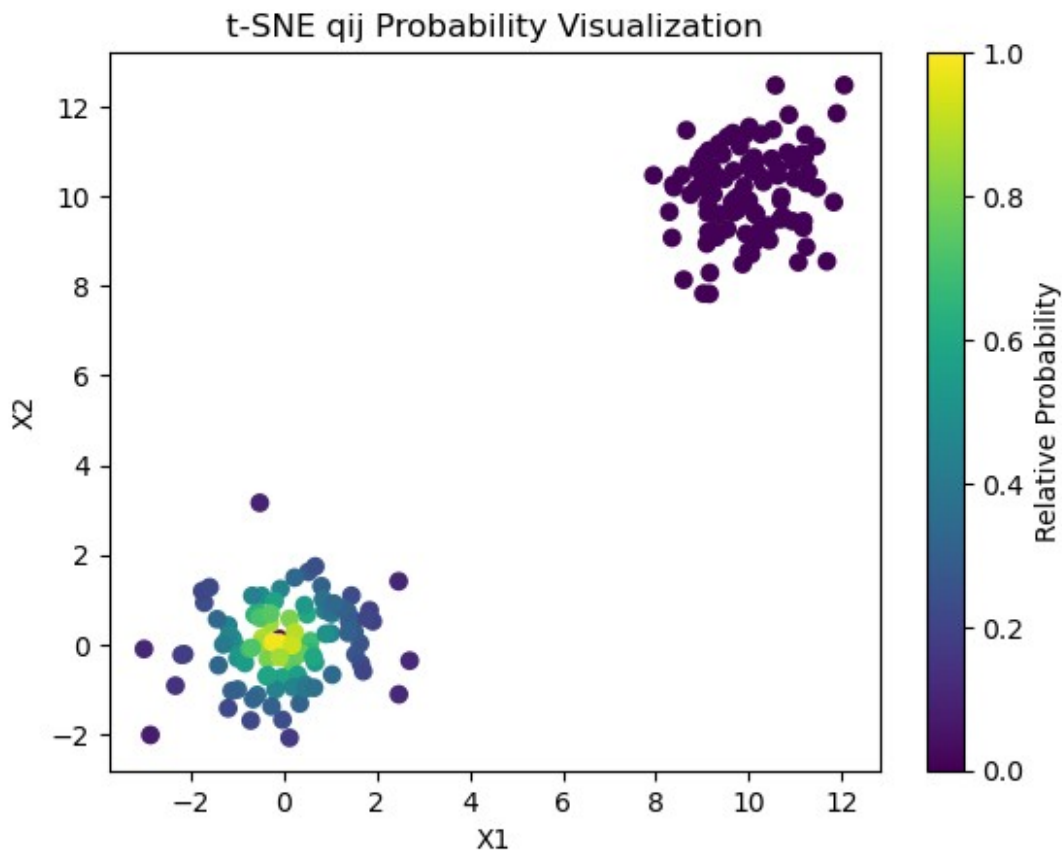
...
[7.14311664e-07 8.58208464e-07 7.32793199e-07 ... 0.00000000e+00
 1.02517234e-05 3.31086556e-05]
[8.50790570e-07 9.80335848e-07 8.62331998e-07 ... 1.02517234e-05
 0.00000000e+00 1.81070447e-05]
[6.50990816e-07 7.58573976e-07 6.63278281e-07 ... 3.31086556e-05
 1.81070447e-05 0.00000000e+00]]

def plot_qij_probabilities(data, qij_matrix, base_index=0):
    n = data.shape[0]
    probabilities = qij_matrix[base_index]
    max_prob = np.max(probabilities)
    colors = probabilities / max_prob

    plt.scatter(data[:, 0], data[:, 1], c=colors, cmap='viridis')
    plt.colorbar(label='Relative Probability')
    plt.xlabel('X1')
    plt.ylabel('X2')
    plt.title('t-SNE qij Probability Visualization')
    plt.show()

plot_qij_probabilities(data, qij_matrix, 0)

```



Comparing the plots for $\sigma = 1$ and the plot for q_{ij} above, we can see that for the $\sigma = 1$ graph, the probability spread is much smoother than the one for q_{ij} . The graphs are really similar because the KL divergence is really low

```
def compute_kl_divergence(pij, qij):
    kl_divergence = 0.0
    n = pij.shape[0]

    for i in range(n):
        for j in range(n):
            if pij[i, j] > 0:
                kl_divergence += pij[i, j] * np.log(pij[i, j] / qij[i, j])

    return kl_divergence
```

```
kl_divergence = compute_kl_divergence(pij_matrix_sigma_4, qij_matrix)
print("KL-Divergence:", kl_divergence)
```

```
for sigma2 in [0.1, 1, 100]:
    pji_matrix = compute_pji(data, sigma2)
    pij_matrix = compute_pij(pji_matrix)
    qij_matrix = compute_qij(data)
    kl_divergence = compute_kl_divergence(pij_matrix, qij_matrix)
    print(f"KL-Divergence for sigma^2 = {sigma2}: {kl_divergence}")
```

```
KL-Divergence: 0.10608591256867625
KL-Divergence for sigma^2 = 0.1: 1.2987869475472107
KL-Divergence for sigma^2 = 1: 0.10608591256867625
KL-Divergence for sigma^2 = 100: 0.7470215079907863
```

Written Part

g)

The smaller the KL divergence, the more accurate the model is. The sigma values give the standard deviation of the probabilities, and a larger sigma gives a larger threshold for the probability. For the clusters above, a sigma of 1 gives the lowest KL-divergence, thus giving the best model. For these clusters, a σ^2 value of 0.1 gives the largest divergence and thus the worst fit.

h)

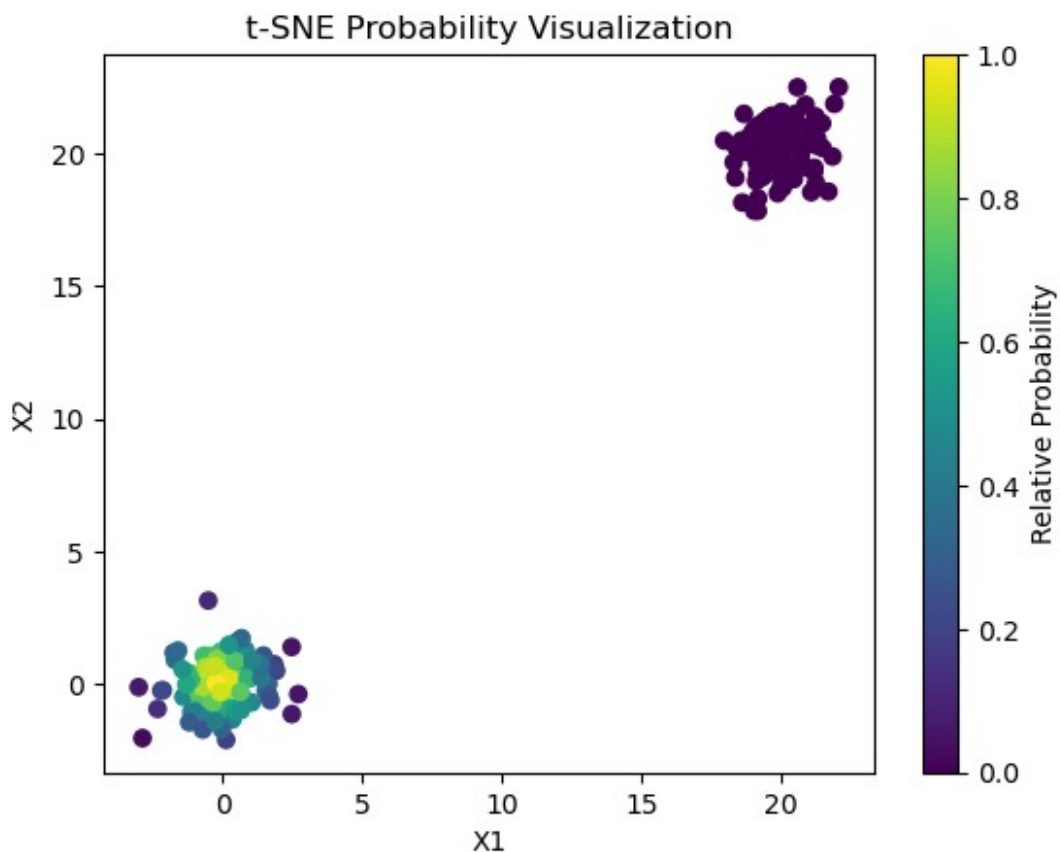
How the parameter matter? As explained above, the sigma gives the threshold for the probabilities of a point being a neighbor of another point. If the points are closer to each other, then a lower sigma would be more suitable when fitting the cluster, and a larger sigma if the points are further away.

```
# Example of simple manual adjustment (moving one cluster)
data_modified = data.copy()
data_modified[99:] += np.array([10, 10]) # Move second cluster

pji_matrix = compute_pji(data_modified, 1)
pij_matrix = compute_pij(pji_matrix)
qij_matrix = compute_qij(data_modified)
kl_divergence = compute_kl_divergence(pij_matrix, qij_matrix)
print("KL-Divergence after manual adjustment:", kl_divergence)

plot_probabilities(data_modified, pij_matrix)

KL-Divergence after manual adjustment: 0.09482073186203122
```



```

def calculate_perplexity(pij_matrix):
    perplexities = []
    n = pij_matrix.shape[0]

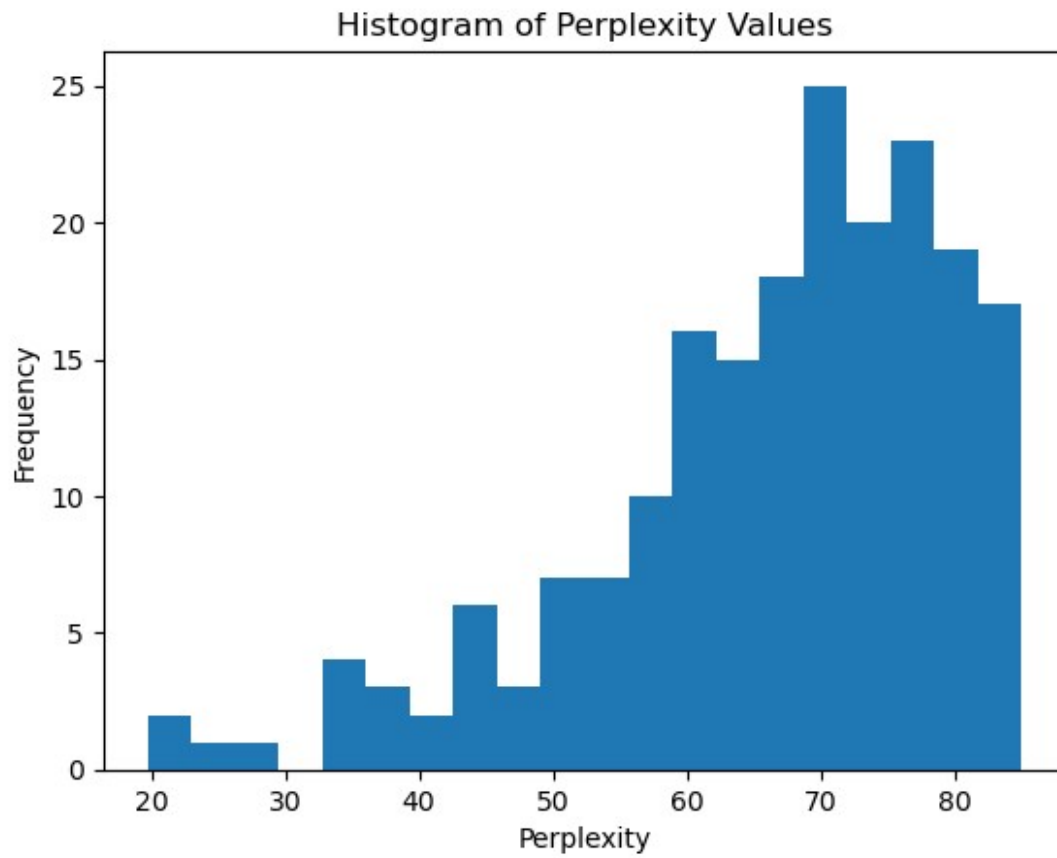
    for i in range(n):
        pi = pij_matrix[i]
        entropy = -np.sum(pi * np.log2(pi + 1e-12))
        perplexity = 2 ** entropy
        perplexities.append(perplexity)

    return perplexities

perplexities = calculate_perplexity(pji_matrix)
plt.hist(perplexities, bins=20)
plt.xlabel('Perplexity')
plt.ylabel('Frequency')
plt.title('Histogram of Perplexity Values')
plt.show()

# Compute KL-divergence for given perplexities
sigma2_values = [5, 25, 50, 100]
for sigma2 in sigma2_values:
    pji_matrix = compute_pji(data, sigma2)
    pij_matrix = compute_pij(pji_matrix)
    qij_matrix = compute_qij(data)
    kl_divergence = compute_kl_divergence(pij_matrix, qij_matrix)
    print(f"KL-Divergence for sigma^2 = {sigma2}: {kl_divergence}")

```



KL-Divergence for $\sigma^2 = 5$: 0.12603272349506978
KL-Divergence for $\sigma^2 = 25$: 0.20667967737238585
KL-Divergence for $\sigma^2 = 50$: 0.36664980915210116
KL-Divergence for $\sigma^2 = 100$: 0.7470215079907863