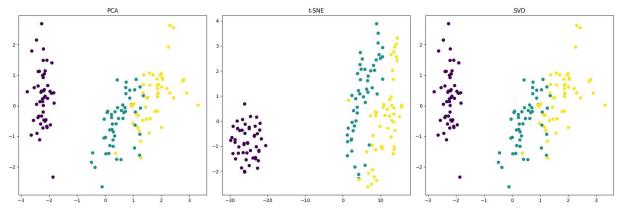
```
In [1]: import numpy as np
        import pandas as pd
        import matplotlib.pyplot as plt
        from sklearn.decomposition import PCA, TruncatedSVD
        from sklearn.datasets import load iris
        from sklearn.preprocessing import StandardScaler
        from sklearn.manifold import TSNE
        import time
In [2]: #Load selected dataset
        data = load iris()
        X = data.data
        v = data.target
In [3]: # Load the dataset
        data = load iris()
        X = data.data
        y = data.target
        # Step 3: Standardize the data
        scaler = StandardScaler()
        X_scaled = scaler.fit_transform(X)
        # Step 4: Apply PCA for dimensionality reduction
        pca = PCA(n_components=2)
        start_time = time.time()
        X pca = pca.fit transform(X scaled)
        pca_time = time.time() - start_time
        # Step 5: Apply other dimensionality reduction methods for comparison
        # TSNE for comparison
        tsne = TSNE(n components=2)
        start_time = time.time()
        X_tsne = tsne.fit_transform(X_scaled)
        tsne_time = time.time() - start_time
        # TruncatedSVD for comparison (alternative to PCA)
        svd = TruncatedSVD(n components=2)
        start_time = time.time()
        X_svd = svd.fit_transform(X_scaled)
        svd time = time.time() - start time
        # Step 6: Visualize the results
        plt.figure(figsize=(18, 6))
        # Plotting PCA
        plt.subplot(1, 3, 1)
        plt.scatter(X_pca[:, 0], X_pca[:, 1], c=y, cmap='viridis')
        plt.title('PCA')
        # Plotting t-SNE
        plt.subplot(1, 3, 2)
        plt.scatter(X_tsne[:, 0], X_tsne[:, 1], c=y, cmap='viridis')
```

```
plt.title('t-SNE')
# Plotting SVD
plt.subplot(1, 3, 3)
plt.scatter(X svd[:, 0], X svd[:, 1], c=y, cmap='viridis')
plt.title('SVD')
plt.tight layout()
plt.show()
# Step 7: Show explained variance ratio for PCA
print("Explained Variance Ratio for PCA:", pca.explained variance ratio )
# Step 8: Discuss the advantages and disadvantages based on the explained variance
# Calculate the cumulative explained variance to assess how much total variance is
cumulative_variance = np.cumsum(pca.explained_variance_ratio_)
# Print out the cumulative explained variance for a clearer view
print(f"Cumulative Explained Variance (for 2 components): {cumulative_variance[-1]:
# Discussion of Advantages and Disadvantages based on the results
if cumulative_variance[-1] < 0.90:</pre>
   print("PCA does not capture enough variance with the chosen components (2D). It
else:
   print("PCA captures a large portion of the variance with just the chosen compon
# Advantages based on explained variance ratio:
print("\nAdvantages of PCA based on explained variance:")
if cumulative_variance[-1] > 0.80:
   print("PCA is effective in reducing dimensionality while retaining most of the
else:
   print("PCA may lose too much information when reducing the dimensions to 2 comp
# Disadvantages based on explained variance ratio:
print("\nDisadvantages of PCA based on explained variance:")
if cumulative_variance[-1] < 0.90:</pre>
   print("PCA may not be ideal when a large amount of data variance is lost in the
else:
   print("PCA effectively preserves the variance, but it may still have limitation
# **Comparison of computational time for each technique**
print("\nComputation Time Comparison:")
print(f"PCA computation time: {pca_time:.4f} seconds")
print(f"t-SNE computation time: {tsne time:.4f} seconds")
print(f"SVD computation time: {svd time:.4f} seconds")
# **Visual Comparison:**
print("\nVisual Comparison:")
print("Observe the scatter plots to evaluate how well each method separates the dat
print("- PCA: Linear dimensionality reduction, captures the largest variance in the
print("- t-SNE: Non-linear method, good for preserving local structure and clusters
print("- SVD: Similar to PCA but can be used in contexts like sparse matrices.")
```



Explained Variance Ratio for PCA: [0.72962445 0.22850762]

Cumulative Explained Variance (for 2 components): 0.96

PCA captures a large portion of the variance with just the chosen components (2D).

Advantages of PCA based on explained variance:

PCA is effective in reducing dimensionality while retaining most of the information in the data.

Disadvantages of PCA based on explained variance:

PCA effectively preserves the variance, but it may still have limitations with non-linear relationships or outliers.

Computation Time Comparison:

PCA computation time: 0.0113 seconds t-SNE computation time: 0.6891 seconds SVD computation time: 0.0022 seconds

Visual Comparison:

Observe the scatter plots to evaluate how well each method separates the data:

- PCA: Linear dimensionality reduction, captures the largest variance in the data.
- t-SNE: Non-linear method, good for preserving local structure and clusters but computationally expensive.
- SVD: Similar to PCA but can be used in contexts like sparse matrices.

PCA (Principal Component Analysis)

Advantages:

- **Simplicity**: PCA is relatively easy to implement and understand. It provides an intuitive method for reducing the dimensionality of data.
- **Interpretability**: Since PCA is linear, it is often easier to interpret. It identifies principal components, which can be mapped back to the original features.
- **Speed**: PCA is computationally efficient and works well on datasets with fewer dimensions.
- **Global Structure Preservation**: PCA preserves global variance (i.e., the global structure of the data), making it good for many types of data.

Disadvantages:

- Linear Assumption: PCA assumes that data lies along linear axes, which limits its ability to capture complex, non-linear relationships between features.
- Sensitivity to Outliers: PCA is sensitive to outliers because they can have a large influence on the computed principal components.
- Interpretability of Components: While PCA is interpretable in terms of variance explained, the principal components themselves may not always map to real-world features in a meaningful way.
- No Clustering Insights: PCA does not explicitly aim to group similar data points together, which can make clustering tasks harder if data is not naturally linear.

t-SNE (t-Distributed Stochastic Neighbor Embedding)

Advantages:

- Non-linear Relationships: t-SNE can capture complex non-linear relationships in data, which is something PCA may struggle with.
- Visualizing Clusters: t-SNE is excellent for visualizing clusters or groups within the data, especially when working with high-dimensional data.
- Effective for High-Dimensional Data: t-SNE is especially effective in visualizing the structure of high-dimensional data when reduced to 2 or 3 dimensions.

Disadvantages:

- Computationally Expensive: t-SNE is much slower and more resource-intensive compared to PCA, especially for large datasets.
- Non-Global Structure Preservation: While t-SNE works well for local structure (nearness), it does not preserve the global structure of the data, meaning distances between clusters or groups may not be accurately represented.
- Parameter Sensitivity: t-SNE requires careful tuning of its parameters (like perplexity), which can significantly affect the resulting visualization.
- No Explicit Component Scores: Unlike PCA, t-SNE does not provide a clear mathematical representation of how features contribute to the reduced dimensions.

SVD (Singular Value Decomposition)

Advantages:

- Sparse Data Handling: SVD works well with sparse matrices and large datasets, making it ideal for text data (such as TF-IDF matrices in NLP).
- **Generalizable**: SVD is applicable to any matrix decomposition and can be used in a wide variety of fields, including image processing and collaborative filtering.
- Data Compression: SVD allows for efficient storage and compression of data, retaining the most important singular values and vectors while discarding the less significant ones.

Disadvantages:

- Less Intuitive: Unlike PCA, SVD does not directly provide a clear interpretation of the components in terms of the original features, which can make it harder to interpret.
- **Computationally Heavy**: While SVD is a powerful technique, it can be computationally intensive, especially for large datasets.
- May Not Preserve Global Structure: SVD, like PCA, is linear and may not capture more complex, non-linear patterns in the data.

Summary:

- PCA is great for quick, interpretable dimensionality reduction when you want to preserve the global variance of the data, especially when the data is linear.
- t-SNE is powerful for visualizing non-linear relationships and clustering, but it is slower and more computationally expensive.
- SVD is ideal for large, sparse datasets and compression tasks but lacks interpretability and may not preserve the global structure as effectively as PCA.