1. Clustering Fundamentals & Evaluation:

- Why: Clustering is a cornerstone of unsupervised learning. Understanding the core concepts (distance metrics, cluster validity, etc.) is essential.
- Scikit-learn: sklearn.cluster (general concepts), metrics in sklearn.metrics (e.g., silhouette_score, calinski_harabasz_score, davies_bouldin_score). Learn how to evaluate the quality of your clusters.
- Key Concepts: Choosing appropriate distance metrics (Euclidean, Manhattan, cosine, etc.), understanding internal vs. external evaluation methods.

2. K-Means Clustering:

- Why: The most widely used and understood clustering algorithm. A solid starting point.
- Scikit-learn: sklearn.cluster.KMeans
- Key Concepts: Centroids, inertia, the elbow method (for choosing k), initialization methods (K-Means++), limitations of K-Means (spherical clusters, sensitivity to outliers).

3. Hierarchical Clustering:

- Why: Offers a different perspective on clustering, creating a hierarchy of clusters (dendrogram). Useful for understanding relationships at different granularities.
- Scikit-learn: sklearn.cluster.AgglomerativeClustering
- **Key Concepts:** Linkage criteria (single, complete, average, ward), dendrogram interpretation, cutting the dendrogram to form clusters.

4. DBSCAN (Density-Based Spatial Clustering of Applications with Noise):

- Why: Handles clusters of arbitrary shapes and identifies noise points (outliers).
 Excellent for data where K-Means struggles.
- Scikit-learn: sklearn.cluster.DBSCAN
- Key Concepts: Core points, border points, noise points, eps (radius) and
 min_samples parameters, advantages over K-Means for non-spherical clusters.

5. Gaussian Mixture Models (GMMs):

 Why: A probabilistic approach to clustering. Assumes data points are generated from a mixture of Gaussian distributions. Provides "soft" cluster assignments (probabilities).

- Scikit-learn: sklearn.mixture.GaussianMixture
- Key Concepts: Probability distributions, expectation-maximization (EM)
 algorithm, covariance types (full, tied, diag, spherical), model selection (BIC,
 AIC).

6. Principal Component Analysis (PCA):

- **Why:** The *fundamental* dimensionality reduction technique. Finds the directions of greatest variance in the data.
- Scikit-learn: sklearn.decomposition.PCA
- Key Concepts: Eigenvectors, eigenvalues, explained variance ratio, choosing the number of components, data scaling (often crucial before PCA).

7. t-distributed Stochastic Neighbor Embedding (t-SNE):

- Why: Excellent for visualizing high-dimensional data in 2D or 3D. Preserves local structure well.
- Scikit-learn: sklearn.manifold.TSNE
- Key Concepts: Perplexity parameter, understanding that t-SNE is primarily for visualization (not for transforming data for other algorithms), non-deterministic nature.

8. Manifold Learning (Beyond t-SNE):

- **Why:** This is important to know how data can lie on a lower-dimensional *manifold* within the higher-dimensional space.
- Scikit-learn: sklearn.manifold (includes Isomap, LocallyLinearEmbedding, etc.)
- Key Concepts: understanding when linear techniques like PCA will be insufficient.

9. Anomaly Detection with Isolation Forest:

- **Why:** A powerful, tree-based method for identifying outliers. Efficient and works well in high-dimensional spaces.
- Scikit-learn: sklearn.ensemble.IsolationForest
- Key Concepts: Isolation, path length, anomaly score, contamination parameter.

10. Preprocessing for Unsupervised Learning:

- **Why:** Data preparation is *critical* for unsupervised learning. Scaling, normalization, and handling missing values can drastically impact results.
- Scikit-learn: sklearn.preprocessing (e.g., StandardScaler, MinMaxScaler, RobustScaler, Normalizer).
- Key Concepts: Understanding when to use different scaling methods, handling outliers appropriately, impact of preprocessing on distance calculations.