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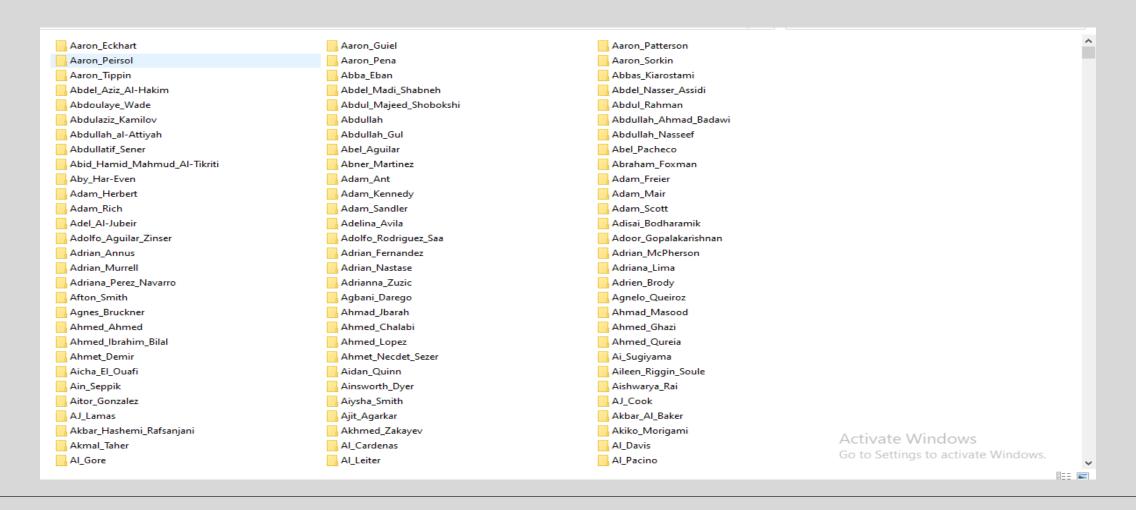
- Dataset Description
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Dataset Description

- Labeled Faces in the Wild (LFW), a database of face photographs designed for studying the problem of unconstrained face recognition. The data set contains more than 13,000 images of faces collected from the web. Each face has been labeled with the name of the person pictured. 1680 of the people pictured have two or more distinct photos in the data set.
- You can download the dataset from here: http://vis-www.cs.umass.edu/lfw/

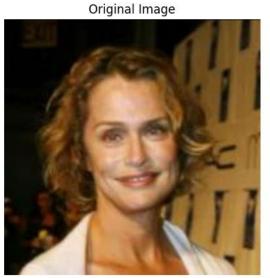
Sample of the Dataset





Main objectives of the analysis:

- This analysis has the following goals:
- 1. Load and Align Faces.
- 2. Find the Embeddings of Face.
- 3. Use Euclidian Distance to Find the Closest Embeddings and Cluster Similar Face Together.







Original Image



Aligned Image



Original Image



Aligned Image

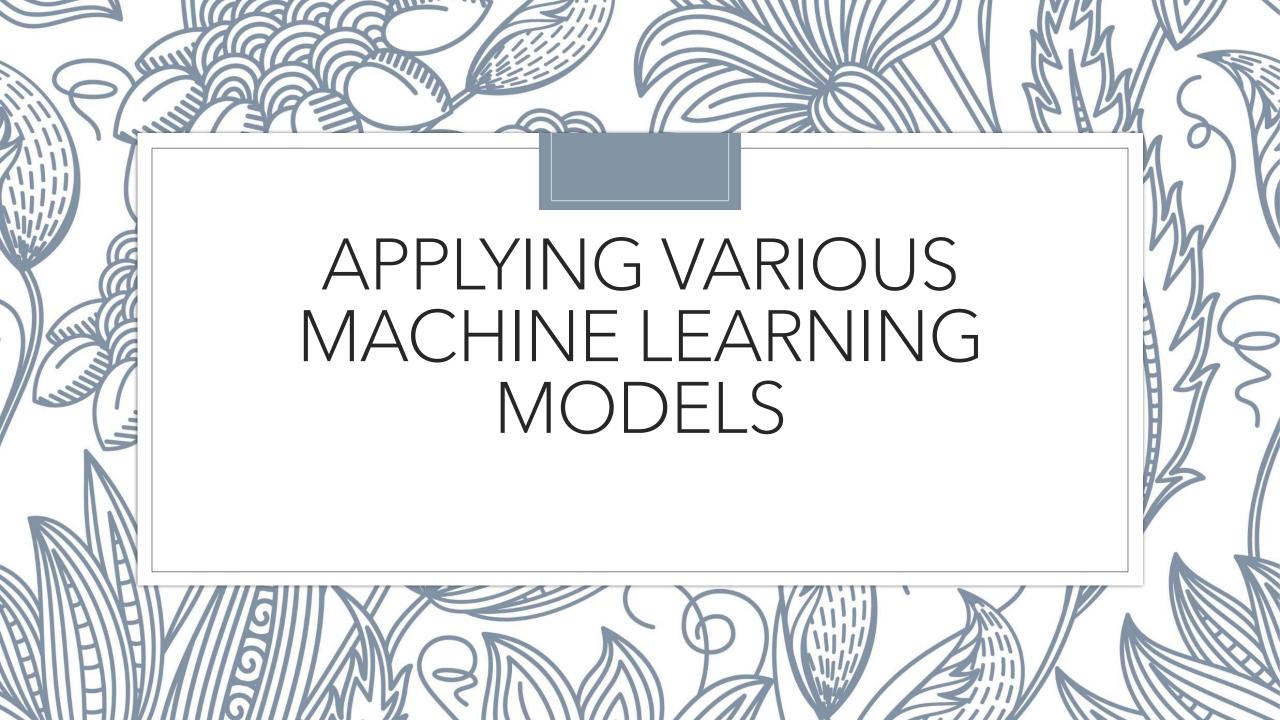


DATA ANALYSIS:

Original Faces Vs Aligned Faces

Main Functions Implemented:

- **1. load_and_align_data:** Initialize the MTCNN face detector and align images using specified parameters.
- **2. generate_embeddings:** Load images, then generate embeddings using a pretrained InceptionResnetV1 model.
- **3. calculate_distance_matrix:** Calculate the Euclidean distance matrix between embeddings.
- 4. sort_faces: Sort faces based on their distances to create a sorted list.
- 5. Cluster Faces: Cluster Facing using K-means and DBSCAN Clustering Algorithms.
- **6. Visualizing The Embeddings:** Use PCA to reduce the dimensionality of face embeddings to 2D and visualizes the embeddings with cluster labels using a scatter plot.



Unsupervised Machine Learning Models Used:

K-means:

K-means is a clustering algorithm that aims to partition data points, in this case face embeddings, into a specified number of clusters based on minimizing the within-cluster sum of squares. It assigns each embedding to the nearest centroid, which effectively groups similar faces together based on their embeddings. By specifying the number of clusters (n_clusters=8), the algorithm identifies distinct groups of similar faces.

DBSCAN:

DBSCAN (Density-Based Spatial Clustering of Applications with Noise) is a density-based clustering algorithm that groups together data points, or face embeddings, that are close to each other and separates regions of higher density from regions of lower density. It identifies dense regions of embeddings and considers them as clusters, effectively grouping similar faces together. The eps parameter (0.55 in this analysis) defines the maximum distance between two embeddings for them to be considered part of the same neighborhood, while the min_samples parameter (4 in this analysis) specifies the minimum number of embeddings required to form a dense region. Unlike K-means, DBSCAN does not require specifying the number of clusters in advance and can automatically detect the number of clusters based on the density of the embeddings.

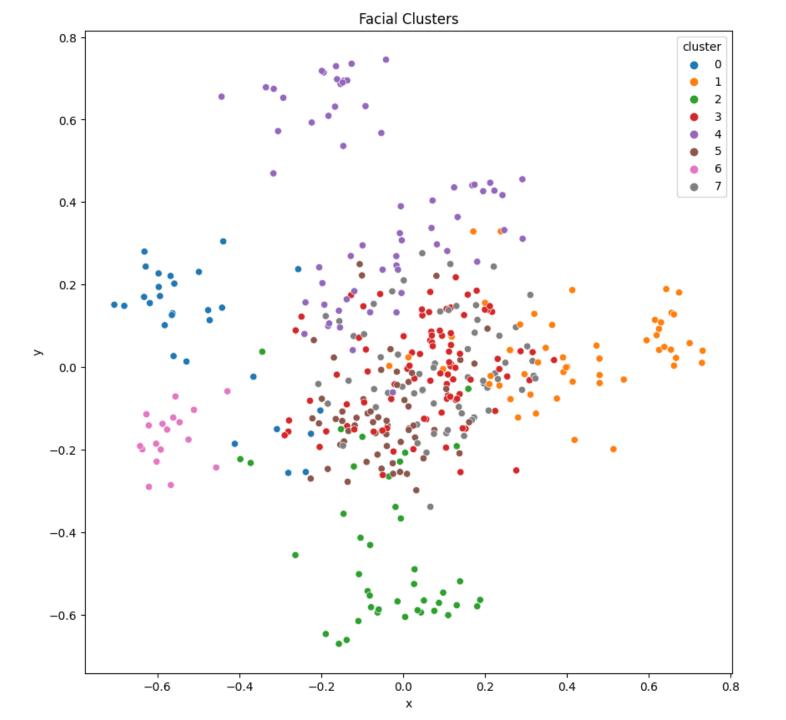
PCA

 It's used to reduce the dimensionality of face embeddings to 2D and visualizes the embeddings with cluster labels using a scatter plot.



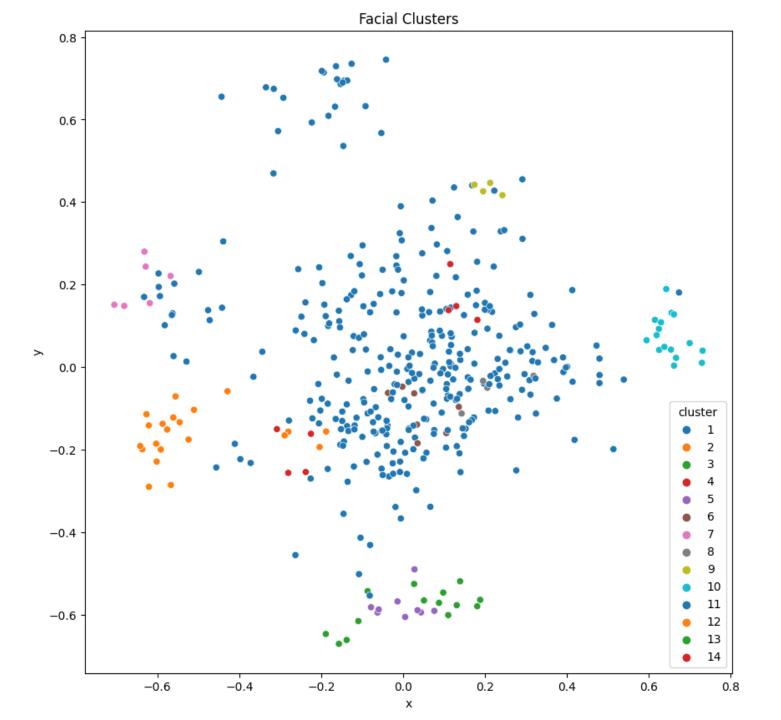
MODELS ANALYSIS AND FINDINGS:

- PCA has successfully reduced the dimensionality of face embeddings to 2D and helped visualizing the embeddings with cluster labels using a scatter plot.
- K-means has demonstrated effective clustering of similar embeddings by assigning each embedding to the nearest centroid, enabling the grouping of similar faces based on their embeddings.
- DBSCAN has shown median success in grouping face embeddings that are close to each other and separating regions of different densities. DBSCAN struggled to differentiate between individuals with slightly similar embeddings, leading to a large portion of the dataset being clustered together as a single group. Nonetheless, DBSCAN tends to successfully identify individuals with highly distinctive embeddings, assigning them to unique clusters.



MODEL ANALYSIS AND FINDINGS:

2D embeddings visualization using PCA for K-means 8 Clusters:



MODEL ANALYSIS AND FINDINGS:

2D embeddings visualization using PCA for DBSCAN Clustering Algorithm:

Cluster 0 Jennifer_Aniston_0004.jpg Jennifer_Aniston_0006.jpg Jennifer_Aniston_0005.jpg Jennifer_Aniston_0009.jpg Jennifer_Aniston_0002.jpg Jennifer_Aniston_0007.jpg Jennifer_Aniston_0003.jpg Jennifer_Aniston_0008.jpg Cluster 2
Heizo_Takenaka_0005.jpg Heizo_Takenaka_0004.jpg James_Gandolfini_0003.jpg Monica_Seles_0001.jpg Cluster 4
Naomi_Watts_0004.jpg Naomi_Watts_0003.jpg

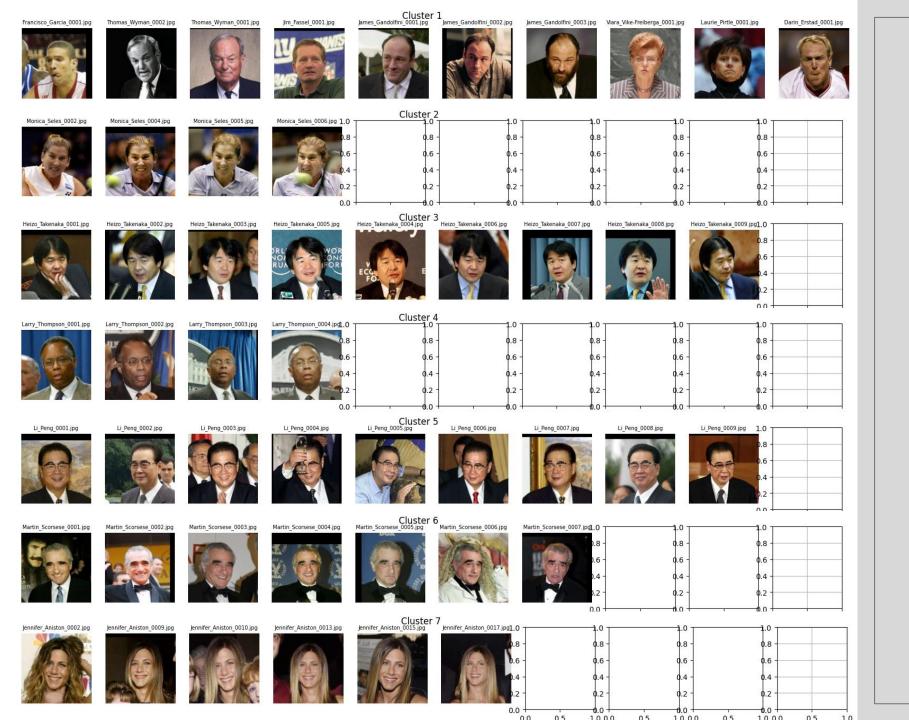


THE 8



Cluster 7
Larry_Brown_0002.jpg

Larry_Brown_0003.jpg



SOME CLUSTERS MADE BY DBSCAN.



Model Flaws and Advanced Steps:

- One potential flaw in the clustering process is that it relies solely on the embeddings to determine similarity. While embeddings capture certain facial features, they may not capture all aspects that humans use to assess similarity, such as hairstyle, facial expressions, or accessories. This could result in misclassifications or grouping of dissimilar faces together.
- Another potential flaw is inaccurate face alignment, which can be addressed by using the original aligned dataset or exploring different alignment techniques for improved accuracy.
- Another potential flaw is the sensitivity of clustering algorithms to the choice of hyperparameters.
 The number of clusters for K-means and the epsilon and min_samples values for DBSCAN need to be carefully selected. Suboptimal choices can lead to overfitting or underfitting, impacting the quality of the clustering results.

Advanced Steps To be Made:

- To mitigate the limitations of relying solely on embeddings, it is beneficial to incorporate additional features, such as facial landmarks or other facial descriptors, in the clustering process. This can provide a more comprehensive representation of facial similarity.
- Exploring alternative clustering algorithms beyond K-means and DBSCAN can be beneficial. Algorithms like spectral clustering, agglomerative clustering, or hierarchical clustering may offer different perspectives on grouping similar faces together and could potentially yield improved results.
- Consider incorporating a validation or evaluation step to assess the quality of the clustering results. This can
 involve using external criteria, such as labeled data or human evaluation, to measure the accuracy and
 effectiveness of the clustering algorithm. This step can help identify potential shortcomings and refine the
 clustering process.

