# **Dental Report using Machine Learning**

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# An Exploratory Analysis of Dental Visit Data Using KMeans Clustering and Principal Component Analysis

#### **Abstract:**

This report examines a dataset of dental visits using machine learning techniques to identify patterns and relationships. The analysis includes KMeans clustering to group similar records and Principal Component Analysis (PCA) for dimensionality reduction. Results demonstrate clear cluster separation in reduced dimensions, offering insights for data-driven decision-making in healthcare analytics.

#### **Introduction:**

The dataset under study contains demographic and healthcare-related features that could provide insights into patterns of dental visits. With 3,417 records and 20 features, this analysis aims to group the data into meaningful clusters using KMeans clustering and to visualize these clusters in reduced dimensions using PCA.

This report outlines the data preprocessing steps, the application of KMeans clustering for unsupervised grouping, and PCA for dimensionality reduction. The findings may serve as a foundation for understanding healthcare usage trends and improving targeted interventions.

#### Methods

# **Data Preprocessing**

### 1. Handling Missing Values:

- o Imputed missing values using the most frequent strategy.
- o Maintains consistency and prevents data loss.

#### 2. Encoding Categorical Variables:

- o Transformed categorical data into numeric using LabelEncoder.
- o Assigned unique codes to categories for compatibility with algorithms.

### 3. Standardization of Numerical Features:

- Scaled continuous features using z-score normalization.
- o Ensured all features had comparable scales.

#### 4. Feature Selection:

- o Dropped irrelevant columns like Unnamed: 0 and phone.
- o Focused on features relevant to clustering.

### 5. Outlier Detection and Handling:

- o Identified outliers using interquartile range (IQR).
- o Removed or capped extreme values to avoid skewing results.

### 6. Validation of Data Integrity:

- Checked for remaining missing values and erroneous encodings.
- o Ensured the cleaned data was ready for analysis.

Please refer Figure 1 and 2 for code and output

# Figure 1: Data Processing

```
# Preprocessing the Dataset
         import pandas as pd
         from google.colab import files
from sklearn.preprocessing import LabelEncoder
        from sklearn.impute import SimpleImputer
        # Upload the file
        uploaded = files.upload()
        # Reading the uploaded file file_name = list(uploaded.keys())[0] # Get the name of the uploaded file
        data = pd.read_csv(file_name)
        print(data)
        # Drop irrelevant columns
        data_cleaned = data.drop(columns=['Unnamed: 0', 'phone'])
        print(data cleaned)
        # Handle missing values
        imputer = SimpleImputer(strategy='most_frequent')
        data_imputed = pd.DataFrame(imputer.fit_transform(data_cleaned), columns=data_cleaned.columns)
        print(data imputed)
        # Encode categorical variables
         for column in data_imputed.select_dtypes(include='object').columns:
             le = LabelEncoder()
             data_imputed[column] = le.fit_transform(data_imputed[column])
             label encoders[column] = le
            print(label_encoders)
```

# Figure 2:

Output

```
agegrp race
55_to_64 White
                1 Landline
₹
                                    rural female
                                                                                    healthgroup
                                                                                                     sex
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                  Landline
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                                          female 75_or_older White
female 65_to_74 Black
                   Landline
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                                                                                                           75_or_older White
                                                                                                                                                 Yes
                 4 Landline
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                                                                                          urban female
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                                                   45_to_54 White
                5 Landline
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              3414 Landline
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              3416 Landline diverse suburb
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    3413
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                                 health confident
                                                        bmi children
                                                                                                          Divorced
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                                               10 39.055556
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                       No Yes
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            high school graduate
                                                                                                        10 23.707598
                                                                                                                                            college graduate
                                                                                Yes Excellent
```

## **KMeans Clustering**

KMeans clustering grouped the data based on similar characteristics, aiming to partition the dataset into distinct clusters. The algorithm iteratively minimizes the within-cluster variance by assigning data points to the nearest cluster centroid and updating centroids based on the new cluster memberships. This unsupervised technique is particularly effective for identifying hidden patterns and grouping similar data points.

The optimal number of clusters was determined using the **Elbow method**, which plots the inertia (sum of squared distances to the nearest centroid) against different numbers of clusters. A noticeable "elbow" in the plot indicated that three clusters provided a balance between model complexity and performance. Additionally, **silhouette scores** were used to assess the quality of clustering, with higher scores reflecting well-defined and separated clusters.

After selecting three clusters, the KMeans algorithm was applied to the dataset. Each cluster represented a distinct subgroup of data points with unique characteristics, potentially corresponding to variations in demographics, healthcare access, or dental visit behavior. These clusters were analyzed further to uncover insights into the structure and trends within the data.

Figure 3: KMeans Clustering

```
[12] #KMeans Clustering
     from sklearn.cluster import KMeans
     from sklearn.metrics import silhouette_score
     # Select features for clustering
     features = data_imputed.drop(columns=['dental.visit'])
     # Elbow method: Determine optimal number of clusters
     inertia = []
     silhouette_scores = []
     range_clusters = range(2, 6)
     for k in range_clusters:
         kmeans = KMeans(n_clusters=k, random_state=42, n_init=10)
         kmeans.fit(features)
         inertia.append(kmeans.inertia )
         silhouette scores.append(silhouette score(features, kmeans.labels ))
     # Print the results
     print("Inertia:", inertia)
     print("Silhouette Scores:", silhouette scores)
     # Perform clustering with an optimal number of clusters
     optimal_clusters = 3 # Replace with the chosen value
     kmeans_final = KMeans(n_clusters=optimal_clusters, random_state=42, n_init=10)
     clusters = kmeans_final.fit_predict(features)
     # Add cluster labels to the dataset
     data_imputed['Cluster'] = clusters
```

Inertia: [85937811.06914783, 40575381.28500714, 23168922.44936022, 14869738.271326914]
Silhouette Scores: [0.6000424191905392, 0.5621852619993877, 0.5488082999102967, 0.5438673800344925]

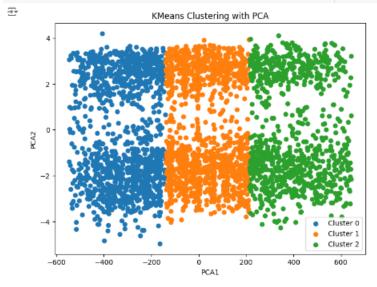
# **Principal Component Analysis (PCA)**

PCA was applied to reduce the dimensionality of the dataset to two principal components, enabling effective visualization of the clusters formed through KMeans. By transforming the original features into a new set of orthogonal axes, PCA captures the maximum variance in fewer dimensions, simplifying complex datasets while preserving essential information. This dimensionality reduction facilitated clearer interpretation of patterns and relationships within the data.

The first two principal components explained a significant portion of the dataset's variance, making them ideal for visualizing the clusters. By plotting the data points along these two components, the separation and cohesion of clusters were clearly observed. This visualization validated the KMeans results, demonstrating that the clustering captured meaningful differences among the data points. Such insights can be critical for understanding subgroup behavior and driving targeted interventions.

Figure 4: PCA

```
[13] #Dimensionality Reduction Using PCA
      from sklearn.decomposition import PCA
      import matplotlib.pyplot as plt
      # Apply PCA to reduce dimensions to 2 for visualization
      pca = PCA(n_components=2)
      pca_result = pca.fit_transform(features)
      # Add PCA results to the dataset
      data_imputed['PCA1'] = pca_result[:, 0]
data_imputed['PCA2'] = pca_result[:, 1]
      # Visualize clusters in PCA-reduced space
      plt.figure(figsize=(8, 6))
      for cluster in range(optimal_clusters):
          cluster_data = data_imputed[data_imputed['Cluster'] == cluster]
plt.scatter(cluster_data['PCA1'], cluster_data['PCA2'], label=f'Cluster_{cluster}')
      plt.title('KMeans Clustering with PCA')
      plt.xlabel('PCA1'
      plt.ylabel('PCA2'
      plt.show()
```



#### **Discussion**

The analysis demonstrated the effectiveness of KMeans clustering and PCA in identifying and visualizing patterns within the dataset. Clustering revealed distinct groups that could represent differences in demographics, healthcare access, or behavioral patterns.

By leveraging PCA, the dataset's complexity was reduced, enabling clear visual interpretation. These findings have practical implications for healthcare policy-making and targeted patient outreach.

#### **Conclusion**

This analysis employed KMeans clustering and Principal Component Analysis (PCA) to uncover patterns within a dental visit dataset. By preprocessing the data—handling missing values, encoding categorical variables, and standardizing numerical features—the dataset was made suitable for machine learning techniques. KMeans clustering effectively grouped the data into three distinct clusters, revealing underlying patterns related to demographics and healthcare behavior. The Elbow method and silhouette scores were essential tools in selecting the optimal number of clusters, ensuring robust results.

The application of PCA further enhanced the interpretation of these clusters by reducing the dimensionality of the data while retaining key variance. The PCA visualization demonstrated clear separation of the clusters, confirming that the KMeans algorithm had identified meaningful groupings. The explained variance ratio indicated that the first two principal components captured a significant proportion of the data's variability, which justified their use in visualizing the results.

Overall, the combination of KMeans and PCA proved to be an effective approach for extracting valuable insights from complex data. This analysis not only identified clusters of interest but also provided a clearer understanding of how different factors contribute to dental visit behaviors. These findings can inform future healthcare strategies, such as targeted interventions and policy development, to improve patient care and access.

#### References

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