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## MINI PROJECT 2

## Tasks done:

- 1. Implemented random and stratified sampling
- 2. Implemented K-means algorithm and chose the optimal value of k using elbow graph.
- 3. Found intrinsic dimensionality of the data (random and stratified) using PCA.
- 4. Produced scree plot visualization (for random and stratified data).
- 5. Plotted MDS (for random and stratified data).
- 6. Obtained the three attributes with highest PCA loadings.
- 7. Visualized the data (random and stratified) projected into the top two PCA vectors via 2D scatterplot.
- 8. Visualized the data via MDS (Euclidian & correlation distance) in 2D scatterplots for random and stratified data.
- 9. Visualized scatterplot matrix of the three highest PCA loaded attributes(for random and stratified data).

# Implementation:

```
    Random points -
        random_points=500
        random_row=np.asarray(random.sample(range(2000),random_points))
        df_random=df.loc[random_row]
    Optimal value of K in K-means
        distortions = []
        K = range(1,20)
        for k in K:
            kmeanModel = KMeans(n_clusters=k).fit(df_normalized)
            kmeanModel.fit(df_normalized)
            distortions.append(kmeanModel.inertia_)
            plt.plot(K, distortions, 'bx-')
            plt.xlabel('k')
```

```
plt.ylabel('Distortion')
   plt.show()
3. Stratified points-
   df_stratified=df
    kmeanModel = KMeans(n_clusters=3).fit(df_normalized)
   kmeanModel.fit(df_normalized)
   groups=kmeanModel.labels_
   df_stratified["groups"]=groups
   stratified_points=500
    percentage=stratified_points/len(df)
   pd_result=pd.DataFrame()
   for i in range(0,3):
   x=df_stratified[df_stratified["groups"]==i]
   no_random_points=math.ceil(percentage*len(x))
    random_row=np.asarray(random.sample(range(len(x)),no_random_points))
   points=x.iloc[random_row,:]
    pd_result=pd.concat([pd_result,points])
4. Eigen values-
    pca_stratified = PCA(n_components=10).fit(standard_data_stratified)
    eigenvalues=pca_stratified.explained_variance_
5. PCA values-
   pca_random = PCA(n_components=2).fit_transform(df_random)
   PCA1=pca_random[:,0]
   PCA2=pca_random[:,1]
```

6. Euclidean distance-

```
mds_data = manifold.MDS(n_components=2, dissimilarity='precomputed')
similarity = pairwise_distances(df_random, metric='euclidean') or similarity =
pairwise_distances(df_random, metric='correlation')
components = mds_data.fit_transform(similarity)
data={'X':components[:,0],'Y':components[:,1]}
```

#### 7. MDS matrix-

```
sq_loading = get_intrinsic_dimensionality(df_random, 3)
top_columns = sorted(range(len(sq_loading)), key=lambda k: sq_loading[k], reverse=True)[:3]
def get_intrinsic_dimensionality(data, k):
std_data = StandardScaler().fit_transform(data);
cov_mat = np.cov(std_data.T)
eigenValues, eigenVectors = np.linalg.eig(cov_mat)
squaredLoadings = []
ftrCount = len(eigenVectors)
for ftrld in range(0, ftrCount):
loadings = 0
for compld in range(0, k):
loadings = loadings + eigenVectors[compld][ftrld] * eigenVectors[compld][ftrld]
squaredLoadings.append(loadings)
return squaredLoadings
```

## Observations:

- 1. Random data has more bias then stratified sampling when compared to the original data.
- 2. Eigen values from random data has big variance when run multiple times, whereas the eigen values from stratified data fairly remains the same.
- 3. PCA2 in random data has more range than in stratified data.
- 4. MDS(Euclidean) data shows clearly segregated groups in stratified data.
- 5. MDS(Correlation) data is very compact and closely related.

# Alternative solutions:

- 1. Task 1: To randomize the data, either random function from pandas can be used to random numbers can be generated from 0 to nrow(df) and then these values as index, can be used to extract data from the data.
- 2. Task 2 : To find Eigen values, either the correlation matrix can be used or variance\_explained\_variable from PCA object.

## Youtube Video link:

https://youtu.be/OnWOU4fVJeA

# References:

- 1. TA's sample code(Code from Ayush Kumar's GitHub repo).
- 2. <a href="https://bl.ocks.org/mbostock/4063663">https://bl.ocks.org/mbostock/4063663</a>
- 3. <a href="https://pythonprogramminglanguage.com/kmeans-elbow-method/">https://pythonprogramminglanguage.com/kmeans-elbow-method/</a>
- 4. <a href="https://support.minitab.com/en-us/minitab/18/help-and-how-to/modeling-statistics/multivariate/how-to/factor-analysis/methods-and-formulas/method