

Source Code:

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import numpy as np
import pandas as pd
import seaborn as sns
train = pd.read excel('train.xlsx')
test = pd.read_excel('test.xlsx')
#append train test data
train['type']='train'
test['type']='test'
data = train.append(test, sort=False)
data.head()
data.info()
data.describe()
data.isnull().sum()
#remove null values
data.dropna(subset=['CustomerID'], inplace=True)
data.isnull().sum()
#Remove duplicate values
data[data.duplicated()].shape
data.drop duplicates(inplace=True)
#Reset the index
indexcol=np.array(list(range(0,len(data))))
data.set index(indexcol,inplace=True)
#Remove the orders that were reversed
data = data[data['Quantity'] > 0]
data.shape
data.head()
# Transfer the dataframe into a file for analysis in Tableau
#data.to csv('file.csv')
#Customers who ordered more than once
n orders = data.groupby(['CustomerID'])['InvoiceNo'].nunique()
mult orders perc = np.sum(n orders > 1) / data['CustomerID'].nunique()
print(f {100 * mult orders perc:.2f}% of customers ordered more than once.')
import datetime as dt
#Perform cohort analysis
def get month(x) : return dt.datetime(x.year,x.month,1)
data['InvoiceMonth'] = data['InvoiceDate'].apply(get month)
grouping = data.groupby('CustomerID')['InvoiceMonth']
data['CohortMonth'] = grouping.transform('min')
data.tail()
def get month int (dframe, column):
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year = dframe[column].dt.year
  month = dframe[column].dt.month
  day = dframe[column].dt.day
  return year, month, day
invoice year,invoice month, = get month int(data, 'InvoiceMonth')
cohort year, cohort month, = get month int(data, 'CohortMonth')
year diff = invoice year - cohort year
month diff = invoice month - cohort month
data['CohortIndex'] = year diff * 12 + month diff + 1
#Count monthly active customers from each cohort
grouping = data.groupby(['CohortMonth', 'CohortIndex'])
cohort_data = grouping['CustomerID'].apply(pd.Series.nunique)
# Return number of unique elements in the object.
cohort data = cohort data.reset index()
cohort counts = cohort data.pivot(index='CohortMonth',columns='CohortIndex',values='CustomerID')
cohort counts
# Retention table
cohort size = cohort counts.iloc[:,0]
retention = cohort counts.divide(cohort size,axis=0) #axis=0 to ensure the divide along the row axis
retention.round(3) * 100 #to show the number as percentage
#Build the heatmap
import matplotlib.pyplot as plt
plt.figure(figsize=(15, 8))
plt.title('Retention rates')
sns.heatmap(data=retention,annot = True,fmt = '.0%',vmin = 0.0,vmax = 0.5,cmap="BuPu r")
plt.show()
#Average quantity for each cohort
grouping = data.groupby(['CohortMonth', 'CohortIndex'])
cohort_data = grouping['Quantity'].mean()
cohort data = cohort data.reset index()
average quantity = cohort data.pivot(index='CohortMonth',columns='CohortIndex',values='Quantity')
average quantity.round(1)
average quantity.index = average quantity.index.date
#Build the heatmap
plt.figure(figsize=(15, 8))
plt.title('Average quantity for each cohort')
sns.heatmap(data=average quantity,annot = True,vmin = 0.0,vmax = 20,cmap="BuGn r")
plt.show()
RFM
#Recency
data['Recent']=(pd.to datetime(data['InvoiceDate'].max()) - pd.to datetime(data['InvoiceDate'])).dt.days
data = data[data['Recent'] <= 366]
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Recency=data.groupby(['CustomerID'], as index=False)['Recent'].max()
Recency.columns = ['CustomerID', 'Recency']
Recency.head()
Recency.shape
#Frequency
Frequency = data.groupby(['CustomerID'], as index=False)['InvoiceNo'].count()
Frequency.columns = ['CustomerID', 'Frequency']
Frequency.head()
#Monetary
data['Totalsales'] = data['Quantity']*data['UnitPrice']
Monetary = data.groupby(['CustomerID'], as index=False)['Totalsales'].agg('sum')
Monetary.columns = ['CustomerID','Monetary']
Monetary.head()
temp df = pd.merge(Recency,Frequency,on='CustomerID')
RFM_metrics = pd.merge(temp_df,Monetary,on='CustomerID')
RFM metrics.head()
#RFM metrics.to csv('file 1.csv')
quantiles = RFM metrics.quantile(q=[0.25,0.5,0.75])
quantiles
quantiles.to dict()
#Define function for the most frequent and high spending customer
def FMscore(x,c,d):
  if x \le d[c][0.25]:
     return 1
  elif x \le d[c][0.50]:
     return 2
  elif x \le d[c][0.75]:
     return 3
  else:
     return 4
#Define function for the most Recent customer
def Rscore(x,c,d):
  if x \le d[c][0.25]:
     return 4
  elif x \le d[c][0.50]:
     return 3
  elif x \le d[c][0.75]:
     return 2
  else:
     return 1
rfm segmentation = RFM metrics
rfm segmentation['R Quartile'] = rfm segmentation['Recency'].apply(Rscore, args=('Recency',quantiles))
rfm_segmentation['F_Quartile'] = rfm_segmentation['Frequency'].apply(FMscore, args=('Frequency',quantiles))
rfm segmentation['M Quartile'] = rfm segmentation['Monetary'].apply(FMscore, args=('Monetary',quantiles))
rfm segmentation.head()
#RFM segment
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rfm segmentation['RFMSegment'] = rfm segmentation['R Quartile'].map(str) + rfm segmentation['F Quartile']
.map(str) + rfm segmentation['M Quartile'].map(str)
rfm segmentation['RFMScore'] = rfm segmentation.R Quartile + rfm segmentation.F Quartile + rfm segment
ation.M Quartile
rfm segmentation.head()
rfm segmentation = rfm segmentation.sort values('RFMScore', ascending=False)
#rfm segmentation.to csv('rfm.csv')
rfm segmentation.head()
print("Best Customers: {}".format(len(rfm segmentation[rfm segmentation['RFMScore'] == 12])))
print("Frequent Customers: {}".format(len(rfm segmentation[rfm segmentation['F Quartile'] == 4])))
print("Money spending Customers: {}".format(len(rfm segmentation[rfm segmentation['M Quartile'] == 4])))
print("Lost Customers: {}".format(len(rfm segmentation[rfm segmentation['RFMScore'] == 3])))
#Normalize and standardize the data
rfm segmentation.skew()
#UnitPrice
sns.boxplot(rfm segmentation['Frequency'])
#Quantity
sns.boxplot(rfm segmentation['Monetary'])
#Remove Outliers
rfm_segmentation1 = rfm_segmentation[(rfm_segmentation.Frequency < 3000) & (rfm_segmentation.Monetar
y < 10000)
rfm segmentation1.skew()
rfm segmentation1['Frequency'] = np.log(rfm segmentation1['Frequency']+0.01)
rfm segmentation1['Monetary'] = np.log(rfm segmentation1['Monetary']+0.01)
rfm segmentation1.skew()
rfm segmentation1.shape
sns.pairplot(rfm segmentation1,diag kind='kde');
final = rfm segmentation1.iloc[:,1:4]
#Standardise the data
from sklearn.preprocessing import StandardScaler
scale = StandardScaler()
data2 = scale.fit transform(final)
data2
final.head()
from sklearn.cluster import KMeans
from sklearn.metrics import silhouette samples, silhouette score
import matplotlib.pyplot as plt
import matplotlib.cm as cm
import numpy as np
print( doc )
X = data2
range n clusters = [2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16]
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for n clusters in range n clusters:
  # Create a subplot with 1 row and 2 columns
  fig, (ax1, ax2) = plt.subplots(1, 2)
  fig.set size inches(18, 7)
  # The 1st subplot is the silhouette plot
  # The silhouette coefficient can range from -1, 1 but in this example all
  # lie within [-0.1, 1]
  ax1.set xlim([-0.1, 1])
  # The (n clusters+1)*10 is for inserting blank space between silhouette
  # plots of individual clusters, to demarcate them clearly.
  ax1.set ylim([0, len(X) + (n clusters + 1) * 10])
  # Initialize the clusterer with n_clusters value and a random generator
  # seed of 10 for reproducibility.
  clusterer = KMeans(n clusters=n clusters, random state=10)
  cluster labels = clusterer.fit predict(X)
  # The silhouette score gives the average value for all the samples.
  # This gives a perspective into the density and separation of the formed
  # clusters
  silhouette avg = silhouette score(X, cluster labels)
  print("For n clusters =", n clusters,
      "The average silhouette score is:", silhouette avg)
  # Compute the silhouette scores for each sample
  sample silhouette values = silhouette samples(X, cluster labels)
  y lower = 10
  for i in range(n clusters):
    # Aggregate the silhouette scores for samples belonging to
    # cluster i, and sort them
    ith cluster silhouette values = \
       sample silhouette values[cluster labels == i]
    ith cluster silhouette values.sort()
    size cluster i = ith cluster silhouette values.shape[0]
    y_upper = y_lower + size_cluster_i
    color = cm.nipy spectral(float(i) / n clusters)
    ax1.fill betweenx(np.arange(y lower, y upper),
                0, ith_cluster_silhouette_values,
                facecolor=color, edgecolor=color, alpha=0.7)
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Label the silhouette plots with their cluster numbers at the middle

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ax1.text(-0.05, y_lower + 0.5 * size_cluster_i, str(i))
     # Compute the new y lower for next plot
     y lower = y upper + 10 \# 10 for the 0 samples
  ax1.set title("The silhouette plot for the various clusters.")
  ax1.set xlabel("The silhouette coefficient values")
  ax1.set ylabel("Cluster label")
  # The vertical line for average silhouette score of all the values
  ax1.axvline(x=silhouette avg, color="red", linestyle="--")
  ax1.set yticks([]) # Clear the yaxis labels / ticks
  ax1.set xticks([-0.1, 0, 0.2, 0.4, 0.6, 0.8, 1])
  # 2nd Plot showing the actual clusters formed
  colors = cm.nipy spectral(cluster labels.astype(float) / n clusters)
  ax2.scatter(X[:, 0], X[:, 1], marker='.', s=30, lw=0, alpha=0.7,
          c=colors, edgecolor='k')
  # Labeling the clusters
  centers = clusterer.cluster centers
  # Draw white circles at cluster centers
  ax2.scatter(centers[:, 0], centers[:, 1], marker='o',
          c="white", alpha=1, s=200, edgecolor='k')
  for i, c in enumerate(centers):
     ax2.scatter(c[0], c[1], marker='$%d$' % i, alpha=1,
             s=50, edgecolor='k')
  ax2.set title("The visualization of the clustered data.")
  ax2.set_xlabel("Feature space for the 1st feature")
  ax2.set ylabel("Feature space for the 2nd feature")
  plt.suptitle(("Silhouette analysis for KMeans clustering on sample data"
           "with n clusters = %d" % n clusters),
          fontsize=14, fontweight='bold')
plt.show()
#Since the silhouette score is more at clusters 2, we will use the value
model = KMeans(n clusters=2)
model.fit(data2)
group = model.predict(data2)
final.head()
final['group'] = group
# to plot in tableau
#final.to csv('cluster data.csv')
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```
sns.pairplot(final,diag_kind='kde',hue='group');
#Elbow method to plot in tableau
wcss=[]
cluster=[]
for i in range(1,10):
    model = KMeans(n_clusters=i)
    model.fit(final)
    wcss.append(model.inertia_)
    cluster.append(i)
from pandas import DataFrame
plot = list(zip(cluster,wcss))
df = DataFrame(plot,columns=['No of cluster','error'])
#df.to_csv('error.csv')
```