customer_segments

March 5, 2016

1 Creating Customer Segments

In this project you, will analyze a dataset containing annual spending amounts for internal structure, to understand the variation in the different types of customers that a wholesale distributor interacts with.

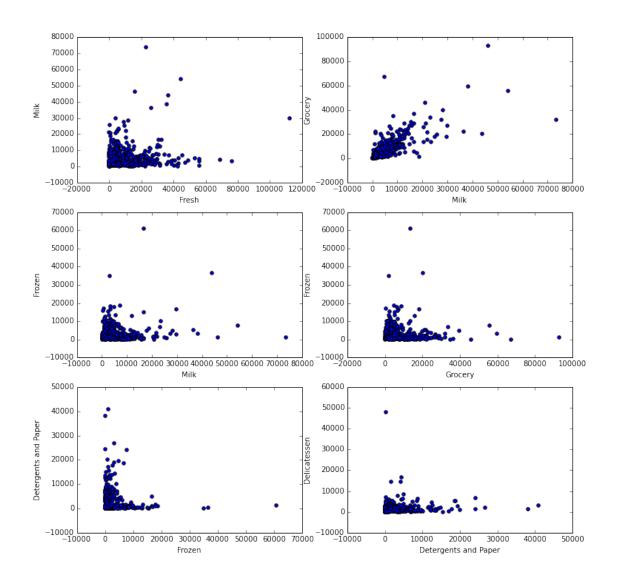
Instructions:

- Run each code block below by pressing **Shift+Enter**, making sure to implement any steps marked with a TODO
- Answer each question in the space provided by editing the blocks labeled "Answer:".
- When you are done, submit the completed notebook (.ipynb) with all code blocks executed, as well as a .pdf version (File > Download as).

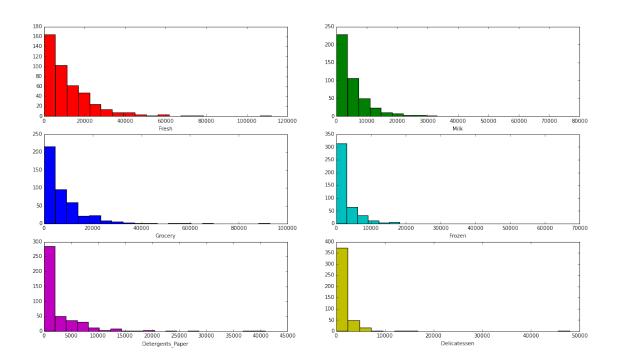
```
In [85]: # Import libraries: NumPy, pandas, matplotlib
         import numpy as np
         import pandas as pd
         import matplotlib.pyplot as plt
         # Tell iPython to include plots inline in the notebook
         %matplotlib inline
         import warnings
         warnings.filterwarnings('ignore')
         pd.options.display.float_format = '{:.5f}'.format
         pd.set_option('display.max_columns', 500)
         pd.set_option('display.width', 1000)
         # Read dataset
         data = pd.read_csv("wholesale-customers.csv")
         num_features=data.shape[1]
         num_data_points=data.shape[0]
         print "Dataset has {} rows, {} columns".format(num_data_points,num_features)
         print data.head() # print the first 5 rows
         print data.describe()
         ,,,
         TODOs:
         1. Create 3-D plot for pca vectors.
         2. Apply PCA on ICA-demixed-transformed data.
         3. Draw elbow graph to identify k in k-means.
Dataset has 440 rows, 6 columns
  Fresh Milk Grocery Frozen Detergents_Paper Delicatessen
```

```
12669 9656
                   7561
                            214
                                             2674
                                                           1338
   7057 9810
                   9568
1
                           1762
                                             3293
                                                           1776
2
   6353 8808
                   7684
                           2405
                                             3516
                                                           7844
3 13265 1196
                   4221
                           6404
                                              507
                                                            1788
 22615 5410
                   7198
                           3915
                                             1777
                                                           5185
                                                Frozen Detergents_Paper Delicatessen
             Fresh
                          Milk
                                   Grocery
count
         440.00000
                     440.00000
                                 440.00000
                                             440.00000
                                                               440.00000
                                                                              440.00000
       12000.29773 5796.26591 7951.27727 3071.93182
mean
                                                               2881.49318
                                                                             1524.87045
std
       12647.32887 7380.37717 9503.16283 4854.67333
                                                               4767.85445
                                                                             2820.10594
min
           3.00000
                      55.00000
                                   3.00000
                                              25.00000
                                                                  3.00000
                                                                                3.00000
25%
       3127.75000 1533.00000 2153.00000
                                             742.25000
                                                                256.75000
                                                                              408.25000
50%
       8504.00000 3627.00000 4755.50000 1526.00000
                                                                816.50000
                                                                              965.50000
75%
      16933.75000 7190.25000 10655.75000 3554.25000
                                                               3922.00000
                                                                             1820.25000
      112151.00000 73498.00000 92780.00000 60869.00000
                                                              40827.00000
                                                                            47943.00000
Out[85]: '\nTODOs:\n1. Create 3-D plot for pca vectors.\n2. Apply PCA on ICA-demixed-transformed data.\n
In [86]: ## Cleanup data, remove outliers.
         f, ((ax1, ax2), (ax3, ax4), (ax5, ax6)) = plt.subplots(3, 2)
         ax1.scatter(data.iloc[:,0],data.iloc[:,1])
         ax1.set_xlabel('Fresh')
         ax1.set_ylabel('Milk')
         ax2.scatter(data.iloc[:,1],data.iloc[:,2])
         ax2.set_xlabel('Milk')
         ax2.set_ylabel('Grocery')
         ax3.scatter(data.iloc[:,1],data.iloc[:,3])
         ax3.set_xlabel('Milk')
         ax3.set_ylabel('Frozen')
         ax4.scatter(data.iloc[:,2],data.iloc[:,3])
         ax4.set_xlabel('Grocery')
         ax4.set_ylabel('Frozen')
         ax5.scatter(data.iloc[:,3],data.iloc[:,4])
         ax5.set_xlabel('Frozen')
         ax5.set_ylabel('Detergents and Paper')
         ax6.scatter(data.iloc[:,4],data.iloc[:,5])
         ax6.set_xlabel('Detergents and Paper')
         ax6.set_ylabel('Delicatessen')
         fig = plt.gcf()
         fig.set_size_inches(12, 12)
         #fig.set_size_inches(18.5, 10.5, forward=True)
         plt.show()
         print "-"*100
```

```
print "Histogram of spending on specific product types."
print "-"*100
f, ((axis1, axis2), (axis3, axis4), (axis5, axis6)) = plt.subplots(3, 2)
### Visualize data spread.
colormap = np.array(['r', 'g', 'b','c','m','y'])
f.axes[0].hist(data.iloc[:,0],bins=20,color=colormap[0])
f.axes[0].set_xlabel(data.columns.values[0]);
f.axes[1].hist(data.iloc[:,1],bins=20,color=colormap[1])
f.axes[1].set_xlabel(data.columns.values[1]);
f.axes[2].hist(data.iloc[:,2],bins=20,color=colormap[2])
f.axes[2].set_xlabel(data.columns.values[2]);
f.axes[3].hist(data.iloc[:,3],bins=20,color=colormap[3])
f.axes[3].set_xlabel(data.columns.values[3]);
f.axes[4].hist(data.iloc[:,4],bins=20,color=colormap[4])
f.axes[4].set_xlabel(data.columns.values[4]);
f.axes[5].hist(data.iloc[:,5],bins=20,color=colormap[5])
f.axes[5].set_xlabel(data.columns.values[5]);
fig = plt.gcf()
fig.set_size_inches(12, 12)
fig.set_size_inches(18.5, 10.5, forward=True)
plt.show()
```



Histogram of spending on specific product types.



In [87]: ## Cleaning outliers could be useful, since it would remove noise which is more prevelent in l cleaned_data=data.copy(deep=True) cleaned_data=cleaned_data[cleaned_data['Fresh']<60000]</pre> cleaned_data=cleaned_data[cleaned_data['Milk']<50000]</pre> cleaned_data=cleaned_data[cleaned_data['Grocery']<50000]</pre> cleaned_data=cleaned_data[cleaned_data['Frozen']<30000]</pre> cleaned_data=cleaned_data[cleaned_data['Detergents_Paper']<20000]</pre> cleaned_data=cleaned_data[cleaned_data['Delicatessen']<20000]</pre> # Removed scaling since units are same, and feature-wise expenses are part of same expense, i. #from sklearn import preprocessing #cleaned_data[['Fresh','Milk','Grocery','Frozen','Detergents_Paper','Delicatessen']] = cleaned print '-'*100 print " Cleaned, centered, and normalized data." print '-'*100 f, ((ax1, ax2), (ax3, ax4), (ax5, ax6)) = plt.subplots(3, 2)ax1.scatter(cleaned_data.iloc[:,0],cleaned_data.iloc[:,1])

ax2.scatter(cleaned_data.iloc[:,1],cleaned_data.iloc[:,2])

ax3.scatter(cleaned_data.iloc[:,1],cleaned_data.iloc[:,3])

ax1.set_xlabel('Fresh')
ax1.set_ylabel('Milk')

ax2.set_xlabel('Milk')
ax2.set_ylabel('Grocery')

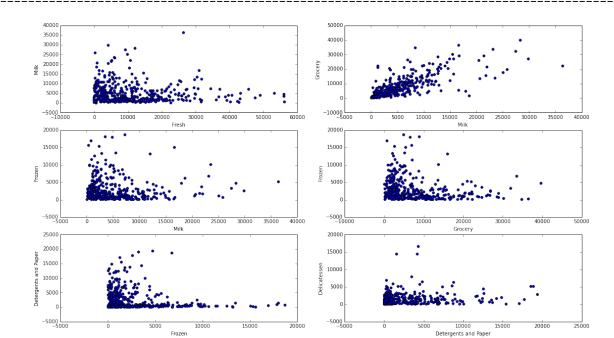
ax3.set_xlabel('Milk')

```
ax3.set_ylabel('Frozen')
ax4.scatter(cleaned_data.iloc[:,2],cleaned_data.iloc[:,3])
ax4.set_xlabel('Grocery')
ax4.set_ylabel('Frozen')
ax5.scatter(cleaned_data.iloc[:,3],cleaned_data.iloc[:,4])
ax5.set_xlabel('Frozen')
ax5.set_ylabel('Detergents and Paper')
ax6.scatter(cleaned_data.iloc[:,4],cleaned_data.iloc[:,5])
ax6.set_xlabel('Detergents and Paper')
ax6.set_ylabel('Delicatessen')
fig = plt.gcf()
fig.set_size_inches(12, 12)
fig.set_size_inches(18.5, 10.5, forward=True)
plt.show()
print " -> Fresh vs Milk, Milk vs Frozen, Grocery vs Frozen, and Detergent_paper vs Frozen all
print "-"*100
print "Histogram of spending on specific product types."
print "-"*100
f, ((axis1, axis2), (axis3, axis4), (axis5, axis6)) = plt.subplots(3, 2)
### Visualize data spread.
colormap = np.array(['r', 'g', 'b','c','m','y'])
f.axes[0].hist(cleaned_data.iloc[:,0],bins=20,color=colormap[0])
f.axes[0].set_xlabel(cleaned_data.columns.values[0]);
f.axes[1].hist(cleaned_data.iloc[:,1],bins=20,color=colormap[1])
f.axes[1].set_xlabel(cleaned_data.columns.values[1]);
f.axes[2].hist(cleaned_data.iloc[:,2],bins=20,color=colormap[2])
f.axes[2].set_xlabel(cleaned_data.columns.values[2]);
f.axes[3].hist(cleaned_data.iloc[:,3],bins=20,color=colormap[3])
f.axes[3].set_xlabel(cleaned_data.columns.values[3]);
f.axes[4].hist(cleaned_data.iloc[:,4],bins=20,color=colormap[4])
f.axes[4].set_xlabel(cleaned_data.columns.values[4]);
f.axes[5].hist(cleaned_data.iloc[:,5],bins=20,color=colormap[5])
f.axes[5].set_xlabel(cleaned_data.columns.values[5]);
fig = plt.gcf()
fig.set_size_inches(12, 12)
fig.set_size_inches(18.5, 10.5, forward=True)
```

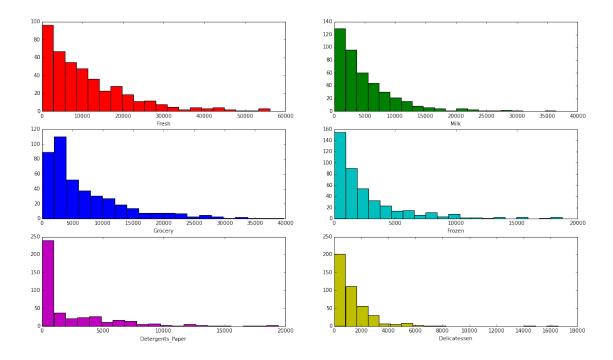
plt.show()

print "-> Plots of Fresh, Milk, Grocery, and Frozen seems to have some similarity in shape and

Cleaned, centered, and normalized data.



-> Fresh vs Milk, Milk vs Frozen, Grocery vs Frozen, and Detergent_paper vs Frozen all seem to have inverted to the seem of spending on specific product types.



-> Plots of Fresh, Milk, Grocery, and Frozen seems to have some similarity in shape and scale.

1.1 Feature Transformation

1) In this section you will be using PCA and ICA to start to understand the structure of the data. Before doing any computations, what do you think will show up in your computations? List one or two ideas for what might show up as the first PCA dimensions, or what type of vectors will show up as ICA dimensions. Answer:

Idea 1. Based on data spread, first PCA would be either fresh, or it could be combination of milk and groceries. Second PCA could include Frozen and Detergent_Paper, and Third PCA could be delicatessen.

Idea 2. ICA could identify perishability as the differentiator in consumables / non-Delicatessen.

1.1.1 PCA

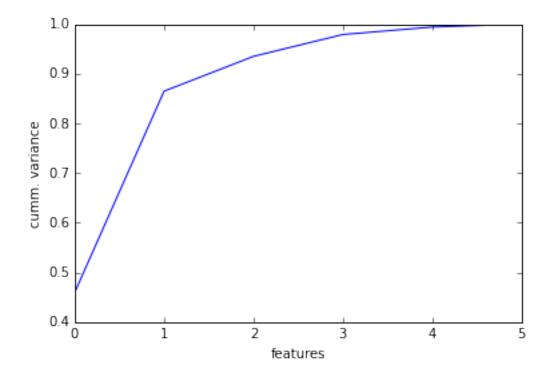
```
In [88]: # TODO: Apply PCA with the same number of dimensions as variables in the dataset
    # Using original data
    from sklearn.decomposition import PCA
    pca = PCA(n_components=num_features,whiten=True)
    pca.fit(data)

# Print the components and the amount of variance in the data contained in each dimension
    print data.columns.values
    print pca.components_
    print pca.explained_variance_ratio_

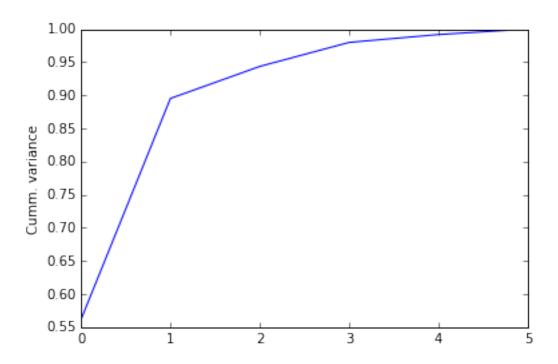
print "\n",'*'*5,"PCA on original data.",'*'*5,"\n"
    pc_df=pd.DataFrame({"pca":pca.explained_variance_ratio_})
    pc_cmf_df=np.cumsum(pc_df)
    print '*'*5," Cumm variance:",'*'*5,"\n",pc_cmf_df
```

```
plt.plot(pc_cmf_df)
         plt.ylabel('cumm. variance')
         plt.xlabel('features')
         plt.show()
['Fresh' 'Milk' 'Grocery' 'Frozen' 'Detergents_Paper' 'Delicatessen']
 \begin{bmatrix} [-0.97653685 & -0.12118407 & -0.06154039 & -0.15236462 & 0.00705417 & -0.06810471] \end{bmatrix} 
 [-0.11061386 \quad 0.51580216 \quad 0.76460638 \quad -0.01872345 \quad 0.36535076 \quad 0.05707921]
 [-0.17855726  0.50988675  -0.27578088  0.71420037  -0.20440987  0.28321747]
 [-0.04187648 -0.64564047 0.37546049 0.64629232 0.14938013 -0.02039579]
 [ 0.015986
                0.20323566 -0.1602915
                                           0.22018612 0.20793016 -0.91707659]
 [-0.01576316 0.03349187 0.41093894 -0.01328898 -0.87128428 -0.26541687]]
[ \ 0.45961362 \ \ 0.40517227 \ \ 0.07003008 \ \ 0.04402344 \ \ 0.01502212 \ \ 0.00613848 ]
**** PCA on original data. ****
**** Cumm variance: ****
      pca
0 0.45961
1 0.86479
2 0.93482
3 0.97884
```

4 0.99386 5 1.00000



```
from sklearn.decomposition import PCA
         pca = PCA(n_components=num_features, whiten=True)
         pca.fit(cleaned_data)
         # Print the components and the amount of variance in the data contained in each dimension
         print cleaned_data.columns.values
         print pca.components_
         print pca.explained_variance_ratio_
         print "\n", '*'*5, "PCA on cleaned data.", '*'*5, "\n"
        pc_df=pd.DataFrame({"pca":pca.explained_variance_ratio_})
        pc_cmf_df=np.cumsum(pc_df)
         print '*'*5," Cumm variance:",'*'*5,"\n",pc_cmf_df
        plt.plot(pc_cmf_df)
        plt.ylabel('Cumm. variance')
['Fresh' 'Milk' 'Grocery' 'Frozen' 'Detergents_Paper' 'Delicatessen']
 \begin{bmatrix} [-0.93484845 & 0.15350373 & 0.26484215 & -0.09783119 & 0.14979567 & -0.01854519] \end{bmatrix} 
 [ 0.33727454  0.49357837  0.72998125  0.00529907  0.32169172  0.0789937 ]
 [-0.09132697  0.61832533  -0.30822162  0.6719085  -0.2013167
                                                               0.14947626]
              0.57071797 -0.34542213 -0.72864576 -0.14113319 0.041873 ]
 [ 0.048662
 [ \ 0.03907126 \ \ 0.12678591 \ -0.31604449 \ \ 0.08770818 \ \ 0.64625958 \ -0.67614383]
 [-0.00831154 \ -0.09325834 \ -0.28772361 \ -0.01796047 \ \ 0.62926724 \ \ 0.71564593]]
**** PCA on cleaned data. ****
**** Cumm variance: ****
     pca
0 0.56148
1 0.89516
2 0.94353
3 0.97976
4 0.99164
5 1.00000
Out[89]: <matplotlib.text.Text at 0x106f72a50>
```

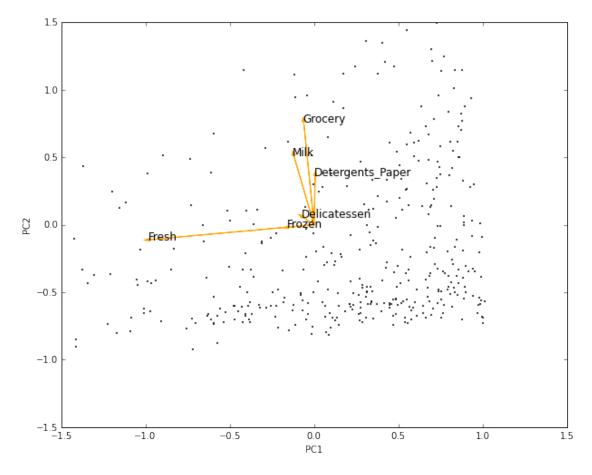


```
In [90]: ''' Following function has been taken from Udacity Forum:
         https://discussions.udacity.com/t/
         having-trouble-with-pca-and-ica-specifically-with-explaining-what-the-dimensions-mean/41890/12. \\
         def biplot12(df):
             # Fit on 2 components
             pca = PCA(n_components=2, whiten=True).fit(df)
             # Plot transformed/projected data
             ax = pd.DataFrame(
                 pca.transform(df),
                 columns=['PC1', 'PC2']
             ).plot(kind='scatter', x='PC1', y='PC2', figsize=(10, 8), s=0.8)
             # Plot arrows and labels
             for i, (pc1, pc2) in enumerate(zip(pca.components_[0], pca.components_[1])):
                 ax.arrow(0, 0, pc1, pc2, width=0.001, fc='orange', ec='orange')
                 ax.annotate(df.columns[i], (pc1, pc2), size=12)
             return ax
         print '-'*100
         print "PC1 / PC2: Bi-plot of original data"
         print '-'*100
         ax = biplot12(data)
         # Play around with the ranges for scaling the plot
         ax.set_xlim([-1.5, 1.5])
         ax.set_ylim([-1.5, 1.5])
```

```
plt.show()
print '-'*100
print "PC1 / PC2: Bi-plot of cleaned data."
print '-'*100
ax = biplot12(cleaned_data)
# Play around with the ranges for scaling the plot
ax.set_xlim([-1.5, 1.5])
```

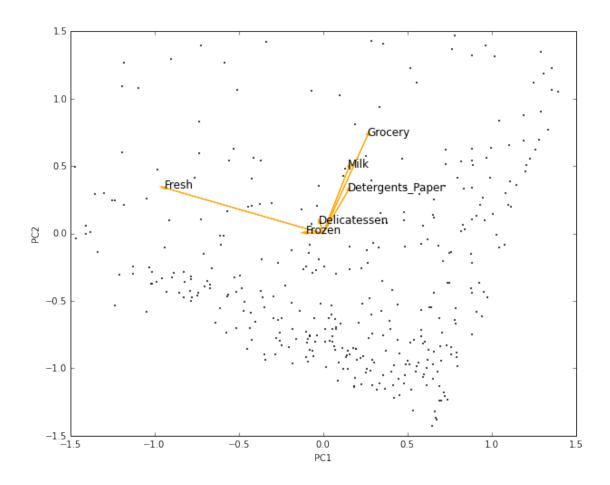
PC1 / PC2: Bi-plot of original data

 $ax.set_ylim([-1.5, 1.5])$



PC1 / PC2: Bi-plot of cleaned data.

Out[90]: (-1.5, 1.5)

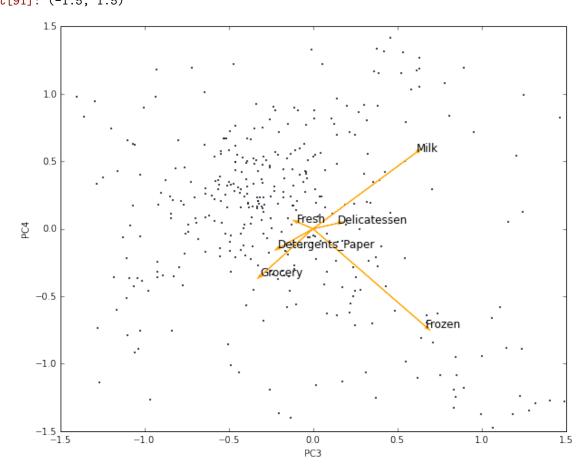


```
In [91]: # TODO: draw a 3-D plot ( or triplot :))
         def biplot34(df):
             # Fit on 2 components
             pca = PCA(n_components=4, whiten=True).fit(df)
             # Plot transformed/projected data
             ax = pd.DataFrame(
                 pca.transform(df),
                 columns=['PC1', 'PC2','PC3','PC4']
             ).plot(kind='scatter', x='PC3', y='PC4', figsize=(10, 8), s=0.8)
             # Plot arrows and labels
             for i, (pc3, pc4) in enumerate(zip(pca.components_[2], pca.components_[3])):
                 ax.arrow(0, 0, pc3, pc4, width=0.001, fc='orange', ec='orange')
                 ax.annotate(df.columns[i], (pc3, pc4), size=12)
             return ax
         print '-'*100
         print "PC3 / PC4: biplot of cleaned data"
         print '-'*100
         ax = biplot34(cleaned_data)
```

```
# Play around with the ranges for scaling the plot
ax.set_xlim([-1.5, 1.5])
ax.set_ylim([-1.5, 1.5])
```

PC3 / PC4: biplot of cleaned data

Out[91]: (-1.5, 1.5)



2) How quickly does the variance drop off by dimension? If you were to use PCA on this dataset, how many dimensions would you choose for your analysis? Why?

Answer: Variance drops fast for first 2 dimentions, but then reduces slowly for remaining dimentions. Given the PCA variance graphs above, elbow is formed at 2nd PCA component, both for original data and scaled data. But since there are data points that have a some variance along multiple PCAs.

3) What do the dimensions seem to represent? How can you use this information?

Answer: PCA here can be used in 2 ways here: 1.) to identify similar customers. 2.)To find similar features. But target here is to find similar customers, and first 2 primary components seem to cover a most of variance.

Then, first PCA dimention corresponds to a segment that spends mostly on Fresh and Frozen products. Second PCA corresponds that spend mostly on Grocery, and significantly on Milk and Detergent_Paper in that order.

We can use this information in many ways: 1.) To transform the data along these 2 PCA, and then find cluster of users using transformed data. But this may not be good approach, since PCA-transformed data might loose some information which could impact be useful for un-biased clustering.

- 2.) To do clustering independently, and then compare the results with those from PCA, to see if both these results are convergent of divergent.
- 3.) We can use the results of PCA further components for supervised learning analysis regression or classification.
- 4.) We could also use K=2 and K=3 for k-mens clustering. Although value of K could depend on elbow in sum-of-square vs k plot.

1.1.2 ICA

```
In [92]: # TODO: Fit an ICA model to the data
        # Note: Adjust the data to have center at the origin first!
        from sklearn.decomposition import FastICA
        from sklearn import preprocessing
        scaled_data=data.copy(deep=True)
        #from sklearn import preprocessing
        scaled_data[['Fresh','Milk','Grocery','Frozen','Detergents_Paper','Delicatessen']] = scaled_da
        ica = FastICA(whiten=True,random_state=0)
        transformed_data=ica.fit_transform(scaled_data)
        # Print the independent components
        print "\n"
        print scaled_data.columns.values
        print ica.components_
        print "\n"
        print preprocessing.StandardScaler().fit_transform(ica.components_)
        #print "\n"
        #print ica.mixing_
['Fresh' 'Milk' 'Grocery' 'Frozen' 'Detergents_Paper' 'Delicatessen']
[[ 0.00259749 -0.01304261  0.06424104  0.00176503 -0.00789576 -0.00472804]
 [ \ 0.0036662 \ \ -0.01675528 \ \ -0.11301178 \ \ \ 0.00711535 \ \ \ 0.13424464 \ \ \ 0.01592772]
 [-0.00189529 \ -0.07279239 \ \ 0.05444162 \ \ \ 0.00183269 \ \ -0.01463357 \ \ \ 0.01719393]
  \begin{bmatrix} -0.05024607 & 0.00639506 & 0.00647498 & 0.00325086 & -0.0104146 & 0.00291214 \end{bmatrix} 
 [-0.00485887 -0.00161266 -0.00552872 -0.00242502 0.0023066
                                                           0.05090388]
 [ 0.51224275 -0.02365303 -1.95603618  0.67015972  2.21983183 -0.03356863]
 [ 0.23572202 -2.13321102  0.9452022
                                    0.42088438 -0.6157666 0.03913673]
  \begin{bmatrix} -2.16830789 & 0.84785851 & 0.11414921 & 0.48780439 & -0.53541034 & -0.78091823 \end{bmatrix} 
 [ 0.08837092  0.54640212  -0.09382264  0.21997389  -0.29311692  1.97475038]
```

4) For each vector in the ICA decomposition, write a sentence or two explaining what sort of object or property it corresponds to. What could these components be used for?

Answer:

```
['Fresh' 'Milk' 'Grocery' 'Frozen' 'Detergents_Paper' 'Delicatessen']
```

A. [0.45910575 0.11611315 1.11498351 0.41769164 -0.48743545 -1.21961489] – Delicacy, Grocery, small qty of everything.

- B. [0.51224275 0.02365303 1.95603618 0.67015972 2.21983183 0.03356863] Detergents_Paper, Grocery,
- C. $[0.23572202 2.13321102 \ 0.9452022 \ 0.42088438 0.6157666 \ 0.03913673]$ Milk and Grocery, small qty of everything except delicacy.
 - D. [-2.16830789 0.84785851 0.11414921 0.48780439 -0.53541034 -0.78091823] Fresh, Milk, Delicacy
 - E. $[0.08837092\ 0.54640212\ -0.09382264\ 0.21997389\ -0.29311692\ 1.97475038]$ Delicacy, Milk
 - F. [0.87286647 0.64649027 -0.12447609 -2.21651401 -0.28810251 0.02021463] Frozen, fresh, some milk
- -> Store of type A and C buy primarily Groceries, but also little bit of everything. -> Store B buys Detergent, Paper, and Groceries. -> D,E, and F dont buy much grocery, and seem to be specialized store, like bakery or chocolatier. -> So this could mean the purchasing habbits of consumers. For example, consumers purchase milk from different sources. -> There seems to be 3 types of stores. Those buying Grocery and Detergent, those buying milk, fresh, delicacies, and those buy everything.

1.2 Clustering

In this section you will choose either K Means clustering or Gaussian Mixed Models clustering, which implements expectation-maximization. Then you will sample elements from the clusters to understand their significance.

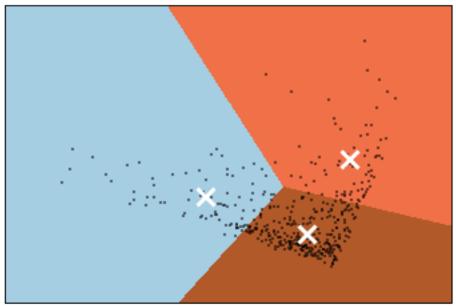
1.2.1 Choose a Cluster Type

- 5) What are the advantages of using K Means clustering or Gaussian Mixture Models? Answer:
 - 1. k-means is intuitive, and fast.
 - 2. k-means can be computed and stored, for later application. This would allow quickly finding similarity.
 - 3. But k-means is strict.
 - 4. GMM is more soft, and allows more realistic / fuzzy interpretation of distributions.
 - 5. GMM is fast, but uses all available features. SO it is important to reduce features before applying GMM.
- 6) Below is some starter code to help you visualize some cluster data. The visualization is based on this demo from the sklearn documentation.

```
In [93]: # Import clustering modules
         from sklearn.cluster import KMeans
         from sklearn.mixture import GMM
In [94]: # TODO: First we reduce the data to two dimensions using PCA to capture variation
         pca = PCA(n_components=2, whiten=True)
         reduced_data = pca.fit_transform(cleaned_data)
         print reduced_data[:10] # print upto 10 elements
[[-0.02449642 0.35143511]
 [ 0.49258938  0.33859559]
 [ 0.48026585  0.15697005]
 [-0.35621581 -0.47218347]
 [-0.98151052 0.47176439]
 [ 0.15642939 -0.09731991]
 [-0.07681722 -0.0824213 ]
 [ 0.37789019  0.07512968]
 [ 0.41110717 -0.41820212]
 [ 0.88192697   1.32444758]]
```

```
In [95]: # TODO: Implement your clustering algorithm here, and fit it to the reduced data for visualiza
         # The visualizer below assumes your clustering object is named 'clusters'
         from scipy.spatial.distance import cdist
         clusters = KMeans(init='k-means++', n_clusters=3, n_init=5).fit(reduced_data)
         print clusters
         # Plot the decision boundary by building a mesh grid to populate a graph.
         x_min, x_max = reduced_data[:, 0].min() - 1, reduced_data[:, 0].max() + 1
         y_min, y_max = reduced_data[:, 1].min() - 1, reduced_data[:, 1].max() + 1
         hx = (x_max-x_min)/1000.
         hy = (y_max-y_min)/1000.
         xx, yy = np.meshgrid(np.arange(x_min, x_max, hx), np.arange(y_min, y_max, hy))
         # Obtain labels for each point in mesh. Use last trained model.
         Z = clusters.predict(np.c_[xx.ravel(), yy.ravel()])
KMeans(copy_x=True, init='k-means++', max_iter=300, n_clusters=3, n_init=5,
    n_jobs=1, precompute_distances='auto', random_state=None, tol=0.0001,
    verbose=0)
In [96]: # TODO: Find the centroids for KMeans or the cluster means for GMM
         centroids = clusters.cluster_centers_
         print centroids
[[-1.56091044 0.46290062]
 [ 0.96914205  1.50025771]
 [ 0.21062931 -0.55903733]]
In [97]: # Put the result into a color plot
         Z = Z.reshape(xx.shape)
         plt.figure(1)
         plt.clf()
         plt.imshow(Z, interpolation='nearest',
                    extent=(xx.min(), xx.max(), yy.min(), yy.max()),
                    cmap=plt.cm.Paired,
                    aspect='auto', origin='lower')
         plt.plot(reduced_data[:, 0], reduced_data[:, 1], 'k.', markersize=2)
         plt.scatter(centroids[:, 0], centroids[:, 1],
                     marker='x', s=169, linewidths=3,
                     color='w', zorder=10)
         plt.title('Clustering on the wholesale grocery dataset (PCA-reduced data)\n'
                   'Centroids are marked with white cross')
         plt.xlim(x_min, x_max)
         plt.ylim(y_min, y_max)
         plt.xticks(())
         plt.yticks(())
         plt.show()
         print cleaned_data.columns.values
         print centroids
         print pca.inverse_transform(centroids)
```

Clustering on the wholesale grocery dataset (PCA-reduced data) Centroids are marked with white cross

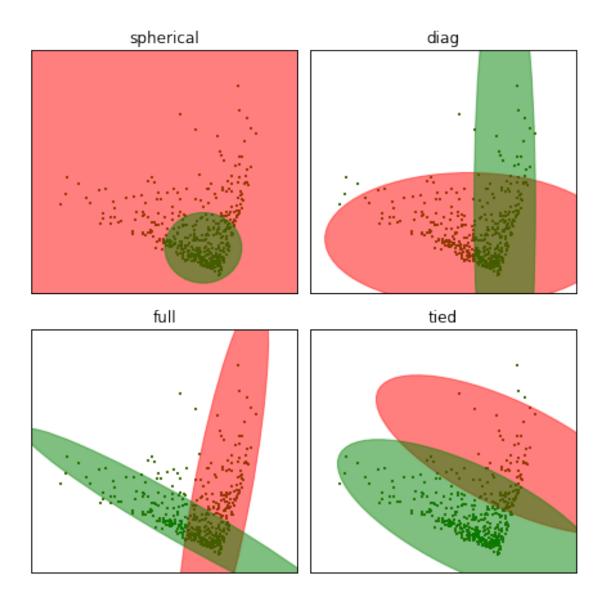


```
['Fresh' 'Milk' 'Grocery' 'Frozen' 'Detergents_Paper' 'Delicatessen']
[[-1.56091044 0.46290062]
[ 0.96914205 1.50025771]
[ 0.21062931 -0.55903733]]
5480.99960792
                                                 4434.28007521
   1216.65602684 2007.14909155]
[ 5525.33801205 13146.09913836 19386.44803806
                                                 1736.19490052
   8272.88602236 2187.37115107]
 [ 7451.44491529 3164.84605956
                                  4306.47936921
                                                 2465.87828229
   1349.06332628 952.61403359]]
In [98]: from matplotlib.colors import LogNorm
        import matplotlib as mpl
        n_classes=2
        def make_ellipses(gmm, ax):
            for n, color in enumerate('rg'):
                v, w = np.linalg.eigh(gmm._get_covars()[n][:2, :2])
                u = w[0] / np.linalg.norm(w[0])
                angle = np.arctan2(u[1], u[0])
                angle = 180 * angle / np.pi # convert to degrees
                v *= 9
                ell = mpl.patches.Ellipse(gmm.means_[n, :2], v[0], v[1],
                                         180 + angle, color=color)
                ell.set_clip_box(ax.bbox)
                ell.set_alpha(0.5)
                ax.add_artist(ell)
```

```
covariance_type=covar_type, init_params='wc', n_iter=20))
                           for covar_type in ['spherical', 'diag', 'tied', 'full'])
         #clf=GMM(n_components=3, covariance_type='full').fit(reduced_data)
        n classifiers = len(classifiers)
        plt.figure(figsize=(3 * n_classifiers / 2, 6))
        plt.subplots_adjust(bottom=.01, top=0.95, hspace=.15, wspace=.05,
                            left=.01, right=.99)
        for index, (name, classifier) in enumerate(classifiers.items()):
             # Since we have class labels for the training data, we can
             # initialize the GMM parameters in a supervised manner.
             # classifier.means_ = np.array([X_train[y_train == i].mean(axis=0)
                                           for i in xrange(n_classes)])
            # Train the other parameters using the EM algorithm.
            classifier.fit(reduced_data)
            h = plt.subplot(2, n_classifiers / 2, index + 1)
            make_ellipses(classifier, h)
            for n, color in enumerate('rg'):
                data = reduced data
                plt.scatter(data[:, 0], data[:, 1], 0.8, color=color)
            plt.xticks(())
            plt.yticks(())
            plt.title(name)
            print cleaned_data.columns.values
            print classifier.sample
            print pca.inverse_transform(classifier.means_)
        plt.legend(loc='lower right', prop=dict(size=12))
        plt.show()
['Fresh' 'Milk' 'Grocery' 'Frozen' 'Detergents_Paper' 'Delicatessen']
<bound method GMM.sample of GMM(covariance_type='spherical', init_params='wc', min_covar=0.001,</pre>
 n_components=2, n_init=1, n_iter=20, params='wmc', random_state=None,
 thresh=None, tol=0.001, verbose=0)>
3069.52928174
   4370.54192259 2000.1008162 ]
 [ 7858.39938446 3247.55825763
                                  4416.06139148
                                                   2500.29482239
   1386.20390951
                   980.38215285]]
['Fresh' 'Milk' 'Grocery' 'Frozen' 'Detergents_Paper' 'Delicatessen']
<bound method GMM.sample of GMM(covariance_type='diag', init_params='wc', min_covar=0.001, n_components=</pre>
 n_init=1, n_iter=20, params='wmc', random_state=None, thresh=None,
```

classifiers = dict((covar_type, GMM(n_components=2,

```
tol=0.001, verbose=0)>
2962.75900897
   1500.71903364 1262.57444695]
 [ \quad 4796.73931178 \quad 10308.36558469 \quad 15145.53444246 \quad 1821.54538804
   6365.46239208 1783.40473597]]
['Fresh' 'Milk' 'Grocery' 'Frozen' 'Detergents_Paper' 'Delicatessen']
<bound method GMM.sample of GMM(covariance_type='full', init_params='wc', min_covar=0.001, n_components='</pre>
 n_init=1, n_iter=20, params='wmc', random_state=None, thresh=None,
 tol=0.001, verbose=0)>
[[ 5217.14891445 8890.91236022 12998.66249246 1939.22189258
   5375.17175343 1614.14527774]
 3907.0170284
                                              3158.0958139
    942.48785762 1238.32459163]]
['Fresh' 'Milk' 'Grocery' 'Frozen' 'Detergents_Paper' 'Delicatessen']
<bound method GMM.sample of GMM(covariance_type='tied', init_params='wc', min_covar=0.001, n_components='</pre>
 n_init=1, n_iter=20, params='wmc', random_state=None, thresh=None,
 tol=0.001, verbose=0)>
[[ 6024.280768
               14741.19838442 21767.02792998
                                              1696.77215436
   9340.7777129 2418.14870719]
 2862.24701105
   1588.78015829 1227.95203683]]
```



7) What are the central objects in each cluster? Describe them as customers.

Answer: We will consider the "full" one correctly explains the soft clustering. Based inverse transformming the centroids from k-means: [0.1663451 - 0.5683769][0.95776694 1.39994104][-1.64821881 0.52313298][7883.6470263 3050.04393497 4118.11897994 2513.50309189 1249.80730062 955.41459363][5354.00667651 12703.41122854 18726.96082769 1743.9919777 7978.08628612 2121.9609403][29844.22811199 4566.91163844 5600.46624755 4531.73759903 1237.26861427 2065.78464029]]

, and also the inverse transform of means values in GMMS, we have following observations:

Centroids in k-means coincide with means of 3 clusters of GMMs.

Cluster 1 (overlap area) has a consumption of everything. Cluster 2 (green)has highest consumption of Milk, Grocery, and Detergents. Cluster 3 (pink) mostly consumes Fresh produce.

1.2.2 Conclusions

** 8)** Which of these techniques did you feel gave you the most insight into the data? Answer:

Tried multiple number of clusters for GMM and k-means, and 3 seemed best for k-means and 2 for GMMs. Although both point to 3 types of clusters, including the overlap in GMMs.

All 3 techniques gave different information, and collating all 3 techniques gives confidence in solution. PCA gave more direct info on primary conponents, while ICA and Clustering gave insight into source and unlabeled similarity in data.

But GMMs seem to be best suited for such a problem, due to overlapping of types. Both PCA and k-means showed that there is no clear separation between the clusters in data.

- 9) How would you use that technique to help the company design new experiments?
- Answer: 1. Company could test and record results using additional feature set, like time of delivery / sales, combine certain features like Fresh and Frozen, or by separating detergent and Paper.
 - 2. Company can also breakup sales for each feature by week / day.
 - 3. Perform classification on each cluster, to identify better components.
 - 10) How would you use that data to help you predict future customer needs?

Answer: 1. We can use day / week data to help identify cyclical patterns 2. Given data can be used to predict sames of one type of product for a customer given its sales of other types of products. 3. Primary components are orthogonal. We can further use k-nearest neighbours to predict how likely is a person to buy Milk, if he already buys fresh produce.