```
In [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 [2]:

```
# 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

# Read dataset
data = pd.read_csv("wholesale-customers.csv")
#print data.describe()
print '\n'
print "Dataset has {} rows, {} columns".format(*data.shape)
print '\n'
print data.head() # print the first 5 rows
print '\n'
```

Dataset has 440 rows, 6 columns

	Fresh	Milk	Grocery	Frozen	Detergents_Paper	Delicatessen
0	12669	9656	7561	214	2674	1338
1	7057	9810	9568	1762	3293	1776
2	6353	8808	7684	2405	3516	7844
3	13265	1196	4221	6404	507	1788
4	22615	5410	7198	3915	1777	5185

```
# 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:

1) List one or two ideas for what might show up as the first PCA dimensions:

PCA, as the name implies, will create "components" that best captures the variance in the data. PCA reduces the number of features, or dimensionality, while minimizing information loss. What we know about the data so far, from the assignment description and the README file, is that the wholesale distribution company caters to both high-volume customers and smaller family-run businesses. The README file gives a high-level overview of the data:

Attribute:	(Min,	Max,	Mean,	<pre>Std. Deviation)</pre>
- Fresh:	(3,	112,151,	12,000.30,	12,47.329)
- Grocery:	(3,	92,780,	7,951.28,	9,503.163)
- Milk:	(55,	73,498,	5,796.27,	7,380.377)
- Frozen:	(25,	60,869,	3,071.93,	4,854.673)
- Detergents_Paper:	(3,	40,827,	2,881.49,	4,767.854)
- Delicatessen:	(3,	47,943,	1,524.87,	2,820.106)

Given the initial information about the types of clients, high-volume customers and smaller family-run shops, and the high-level summary of the data in the README file, I sense that the first PCA dimension (which captures the most variance in the data) will relate to the "Fresh" and "Grocery" items. Both of these items have the greatest standard deviations.

2) What type of vectors will show up as ICA dimensions:

ICA, on the other hand, looks to find components that are statistically independent from each other. ICA will attempt to find clear boundaries between components. I have the sense that the ICA method is not appropriate for this data as there will be no clear division, no real independence between wholesale items. But let's find out. If ICA proves useful however, each of the ICA vectors could represent independent customers.

```
In [3]:
```

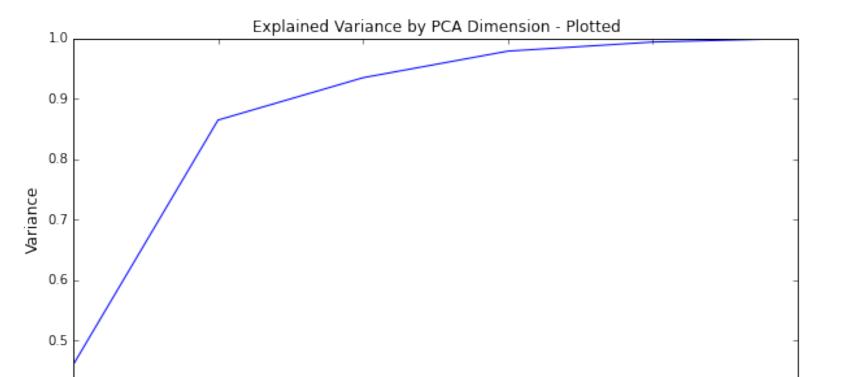
```
print 'Additional information learned after Udacity feedback...'
print 'The data.describe() function provides information on the data...'
print '\n'
print data.describe()
Additional information learned after Udacity feedback...
The data.describe() function provides information on the data...
               Fresh
                               Milk
                                          Grocery
                                                          Frozen
                                                                  \
                         440.000000
          440.000000
                                       440.000000
                                                      440.000000
count
mean
        12000.297727
                       5796.265909
                                      7951.277273
                                                     3071.931818
        12647.328865
                        7380.377175
                                      9503.162829
                                                     4854.673333
std
            3.000000
                          55.000000
                                         3.000000
                                                       25.000000
min
25%
         3127.750000
                       1533.000000
                                                      742.250000
                                      2153.000000
50%
         8504.000000
                       3627.000000
                                      4755.500000
                                                     1526.000000
        16933.750000
75%
                       7190.250000
                                     10655.750000
                                                     3554.250000
                      73498.000000
                                     92780.000000
       112151.000000
                                                    60869.000000
max
       Detergents Paper
                          Delicatessen
             440.000000
count
                            440.000000
            2881.493182
                           1524.870455
mean
std
            4767.854448
                           2820.105937
min
               3.000000
                              3.000000
25%
             256.750000
                            408.250000
50%
             816.500000
                            965.500000
75%
            3922.000000
                           1820.250000
           40827.000000
                        47943.000000
max
In [4]:
# TODO: Apply PCA with the same number of dimensions as variables in the dataset
import pandas as pd
from sklearn.decomposition import PCA
pca = PCA(n components=6)
pca.fit(data)
# for viewing
print '\n'
print 'data.head()...'
print data.head() # print the first 5 rows
print '\n'
# Print the components and the amount of variance in the data contained in each dime
print 'pca.components ...'
print pca.components_
print '\n'
print 'pca.explained variance ratio ...'
print pca.explained variance ratio
plt.figure(figsize=(10,5))
```

nl+ vlahel('DCA Dimensions' fontsize=12)

```
plt.wlabel('Variance', fontsize=12)
plt.title('Explained Variance by PCA Dimension - Plotted')
X = np.arange(1,7)
Y = np.cumsum(pca.explained_variance_ratio_)
plt.plot(X,Y)
print '\n'
```

data.head()...

```
Detergents_Paper
   Fresh
                                                         Delicatessen
         Milk
                 Grocery
                            Frozen
   12669
0
           9656
                     7561
                               214
                                                  2674
                                                                  1338
    7057
          9810
                     9568
                              1762
                                                  3293
                                                                  1776
1
2
    6353
           8808
                     7684
                              2405
                                                  3516
                                                                  7844
3
   13265
           1196
                     4221
                              6404
                                                   507
                                                                  1788
   22615
           5410
                     7198
                              3915
                                                  1777
                                                                  5185
pca.components ...
[[-0.97653685 -0.12118407 -0.06154039 -0.15236462]
                                                         0.00705417 - 0.06810
 [-0.11061386]
                0.51580216
                              0.76460638 - 0.01872345
                                                         0.36535076
                                                                       0.05707
921]
 [-0.17855726 \quad 0.50988675 \quad -0.27578088 \quad 0.71420037 \quad -0.20440987
747]
 [-0.04187648 - 0.64564047 \quad 0.37546049 \quad 0.64629232 \quad 0.14938013 \quad -0.02039
579]
 [ 0.015986
                0.20323566 - 0.1602915
                                           0.22018612 0.20793016 -0.91707
6591
 [-0.01576316 \quad 0.03349187 \quad 0.41093894 \quad -0.01328898 \quad -0.87128428 \quad -0.26541
687]]
pca.explained_variance_ratio_...
[ 0.45961362
               0.40517227 0.07003008 0.04402344 0.01502212
                                                                     0.006138
48]
```



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: The variance drops off significantly after the 2nd primary component. Looking at the results of the plotted PCA graph, I would choose the first two dimensions. The first two dimensions capture 86% of the variance in the data.

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

Answer:

The 1st dimension shows "Fresh" as the variable with the strongest correlation at -0.97:

Fresh Milk Grocery Frozen Deterg... Delicatessen [-0.97653685 -0.12118407 -0.06154039 -0.15236462 0.00705417 -0.06810471]

The 2nd dimension shows "Grocery" and "Milk" as the variables with the strongest correlation at 0.76 and 0.51 respectively:

Fresh Milk Grocery Frozen Deterg... Delicatessen [-0.11061386 0.51580216 0.76460638 -0.01872345 0.36535076 0.05707921]

The 1st principal component is stronly correlated with the "Fresh" variable. The 1st principal component increases as the "Fresh" variable decreases.

corrected paragraph below after feedback...

This component can be viewed as a measure of how "Fresh" orders decline with smaller customers.

Because of the high correlation for "Fresh" (-0.97), this principal component is primarily a measure of the "Fresh" variable. Therefore, high-volume customers order much more "Fresh" items (in comparison to other items) than low-volume customers.

The 2nd principal component is showing a rather steady decline in variance starting with "Grocery" at 0.76 down to "Fresh" at -0.11 (0.76 -> 0.51 -> 0.36 -> 0.05 -> -0.018 -> -0.11). Looking at the data, I will (subjectively) deem important all correlation values above 0.5. This criteria leaves me with two variables, "Grocery" at 0.76 and "Milk" at 0.51. This would indicate that these two variables vary together. I would say that the 2nd principal component is strongly correlated with "Grocery" and "Milk". The high-volume customers order more "Grocery" and "Milk" items than low-volume customers.

In [6]:

print '\n'

print 'The graph below shows, among other things, how high-volume customers order mu

```
#import pandas as pd
from sklearn.decomposition import PCA
def biplot(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=['Primary Component 1', 'Primary Component 2']
    ).plot(kind='scatter', x='Primary Component 1', y='Primary Component 2', figsize
    # 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='blue', ec='blue')
        ax.annotate(df.columns[i], (pc1, pc2), size=12)
    return ax
ax = biplot(data)
ax.set xlim([-1.1, .5])
ax.set_ylim([-.25, 1])
The graph below shows, among other things, how high-volume customers o
rder much more "Fresh" items (in comparison to other items) than low-v
olume customers.
Out[6]:
(-0.25, 1)
   1.0
```

Grocery

Delicatessen

Detergents Paper

Milk

0.8

0.6

0.4

0.2

0.0

Primary Component 2

```
-0.2 - -1.0 -0.8 -0.6 -0.4 -0.2 0.0 0.2 0.4

Primary Component 1
```

```
### ICA
In [7]:
# TODO: Fit an ICA model to the data
# Note: Adjust the data to have center at the origin first!
from sklearn.decomposition import FastICA
#import pandas as pd
data mean = data.mean(axis=0)
data_centered = data - data_mean
ica = FastICA(n components=6, random state=12) # added random state after answering
ica.fit transform(data centered)
# Print the independent components
print "Independent Components:"
print ica.components_
print '\n'
print 'Generated the bar graph below after Udacity review.'
print 'This makes evident the anti-correlation between Grocery and Detergents Paper
print pd.DataFrame(ica.components_, columns = data.columns).plot(kind = 'bar')
Independent Components:
```

```
[[ -3.00060334e-07
                    2.28232144e-06
                                      1.20918798e-05
                                                      -1.46070309e-06
  -2.82157485e-05
                   -5.72571727e-06]
[ 1.52206749e-07
                    9.85239783e-06
                                     -5.79548167e-06
                                                      -3.67037530e-07
   3.26239877e-06 -6.06614791e-06]
[ -2.11582533e-07
                  1.88002073e-06
                                     -6.37841461e-06
                                                      -4.16791731e-07
   7.08979178e-07
                    1.44281972e-06]
   3.97588207e-06 -8.57329842e-07
                                                      -6.77708744e-07
                                     -6.23625425e-07
   2.05778889e-06
                   -1.04492227e-061
[ -8.65226341e-07
                  -1.40291145e-07
                                      7.73545338e-07
                                                       1.11461095e-05
  -5.55265521e-07
                   -5.95215276e-06]
                   2.19351274e-07
                                      6.00186230e-07
                                                       5.22177278e-07
3.86486545e-07
  -5.08096211e-07
                   -1.80922752e-05]]
```

```
Generated the bar graph below after Udacity review.

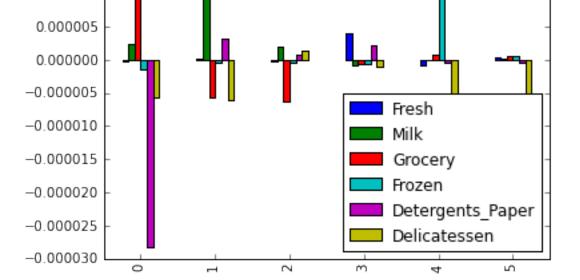
This makes evident the anti-correlation between Grocery and Detergents

_Paper in the first vector...

Axes(0.125,0.125;0.775x0.775)
```

0.000015

0.000010



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:

ICA assumes the independence of each variable. In other words, it assumes that knowing the value of one variable will not tell you anything about other variables. ICA is a good technique to decompose multivariate signals into independent non-gaussian signals. For example, ICA can take a voice recording from a busy cocktail party and decompose the sounds to each of the speakers.

Vector 1

Fresh Milk Grocery Frozen

Detergents_P.. Delicatessen

[-3.86493444e-07, -2.19535813e-07, -6.00059586e-07, -5.22109683e-07, 5.09251107e-07, 1.80923299e-05]

- 1. There looks to be an inverse relationship between Detergents_paper and Grocery.
- 2. The same can be said for Detergents paper and Frozen.
- 3. If I could interpret this vector in terms of a specific customer, I would say that this customer orders items related to "delicatessen" and "detergents_paper". # Additional comment added after Udacity review
- 4. Independent of other effects, there is an anti-correlation between Detergents_paper and grocery. Possibly this could come from a distinction between something like grocery stores and pharmacies which carry some food but far more paper products. Meaning that these customers would purchase either detegents or grocery but not both.

There looks to be an inverse relationship between Grocery and Detergents_Paper.
 If I could interpret this vector in terms of a specific customer, I would say

that this customer orders items related in the "Grocery" category.

Vector 3				
Fresh	Milk	Grocery	Frozen	Detergents_P
Delicatessen				
[-3.97600172e-06,	8.59682742e-07,	6.29261564e-07,	6.77057210e-07,	-2.07148819e-06,
1.04089660e-06]				

1. If I could interpret this vector in terms of a specific customer, I would say that this customer orders items related to "Milk", "Grocery", and "Frozen".

- 1. There looks to be an inverse relationship between Fresh and Grocery
 2. If I could interpret this vector in terms of a specific customer, I would say that this customer orders items related to "Delicatessen", "Fresh" and "Detergents paper".
- Vector 5
 Fresh Milk Grocery Frozen Detergents_P..
 Delicatessen
 [-2.98917871e-07, 2.30935710e-06, 1.20484157e-05, -1.46342068e-06, -2.82047555e-05, -5.72930585e-06]
- 1. If I could interpret this vector in terms of a specific customer, I would say that this customer orders items related to "Grocery" and "Milk".
- 1. There looks to be an inverse relationship between Milk and Delicatessen.
- 2. The same can be said for Milk and Grocery.
- 3. If I could interpret this vector in terms of a specific customer, I would say that this customer orders items related to "Delicatessen" and "Grocery".

What could these components be used for?

The information obtained from the above 6 ICA vectors can be used to understand the types of customers that are purchasing items related to the six categories (fresh, Milk, ..., Delicatessen). You can see that some of the items are highly correlated to each other, such as Grocery and Frozen in vector 1. Equipped with this information, grocery stores could shelve these items closer together for a quicker sale or separate them on opposite parts of the store to have customers walk-by many other products for additional sales opportunities. The success of these strategies could be measured by an A/B test (more on A/B test on question

9).

##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.

Choose a Cluster Type

5) What are the advantages of using K Means clustering or Gaussian Mixture Models?

Answer:

New comments after Udacity review...

The two main differences have to do with speed and structural information of each: Speed

- -> K-Means is much faster and much more scalable
- -> GMM is slower since it has to incorporate information about the distributions of the data, thus it has to deal with co-variance, mean, variance, and prior probabilities of the data, and also has to assign probabilities to belonging to each of the clusters.

Structure

- -> K-Means has straight boundaries (hard clustering)
- -> with GMM, you get much more structural information. This means that you can measure how wide each cluster is, since it works on probabilities (soft clustering)

--- Original answers START---

K-means advantages:

- Fast, scalable
- Gives best results when data is distinct & well seperated

K-means disadvantages:

- not good for unbalaned clusters (when 1 cluster is bigger than the other).
- Non-optimal clustering can occur, depending on where the initial centroids are placed.
- Inconsistent end-results. This algorithm converges to a local optimum. In other words, running the algorithm multiple times will give a different end-result (depending the where the initial centroids are placed).
- K-means is a hill-climbing algorithm. So it's important to place the initial centroids properly

Gaussian Mixture Model (GMM) advantages:

- GMM generalizes k-means clustering. It includes the covariance structure in the data as well as the centers of the latent gaussians.

Gaussian Mixture Model (GMM) disadvantages:

- Computational cost increases dramatically with increase in dimensionality --- Original answers END---

Which algorithm is best-suited for this particular data?

The data does not seem to have a clear divide between the known types of customers. For this reason, I believe that GMM is a better technique than k-means. K-means is also not the best choice for elongated clusters or irregular shapes as it doesn't respond well to them. Also, I believe that clustering the data between 2 groups with GMM will help visualize and relate the data back to the two main customer groups (high-volume and smaller family-run shops).

In addition...

K-means is a special case of GMM where it performs "hard" clustering. In other words, every point is assigned specifically to one cluster instead of each point assigned a probability of belonging to a specific cluster. Also, K-means doesn't work well with clusters of different sizes and density. I think GMM will be a better algorithm as it will describe the data, and the complexities within the data, in ways that K-means cannot capture (such as incorporating covariance structure).

6) Below is some starter code to help you visualize some cluster data. The visualization is based on this demo (http://scikit-learn.org/stable/auto examples/cluster/plot kmeans digits.html) from the sklearn documentation.

```
In [8]:
```

```
# Import clustering modules
from sklearn.cluster import KMeans
from sklearn.mixture import GMM
```

In [9]:

TODO: First we reduce the data to two dimensions using PCA to capture variation reduced_data = PCA(n_components=2).fit_transform(data_centered)

print reduced_data[:10] # print upto 10 elements

```
-650.02212207
                    1585.51909007]
] ]
    4426.80497937
                    4042.45150884]
 4841.9987068
ſ
                    2578.762176
   -990.34643689
                   -6279.80599663]
 [-10657.99873116
                   -2159.72581518]
   2765.96159271
                    -959.87072713]
    715.55089221
                   -2013.00226567]
    4474.58366697
                    1429.49697204]
 [
    6712.09539718
                   -2205.90915598]
    4823.63435407
                   13480.55920489]]
```

```
In [10]:
# TODO: Implement your clustering algorithm here, and fit it to the reduced data for
# The visualizer below assumes your clustering object is named 'clusters'
#GMM - testing/comparing
clusters GMM components 1 = GMM(n components=1).fit(reduced data)
clusters GMM components 2 = GMM(n components=2).fit(reduced data)
clusters_GMM_components_3 = GMM(n_components=3).fit(reduced_data)
clusters GMM components 4 = GMM(n components=4).fit(reduced data)
# The chosen one ...
clusters = GMM(n components=2).fit(reduced data)
#KMeans - testing/comparing
clusters kmeans 1 = KMeans(n clusters=1).fit(reduced data)
clusters kmeans 2 = KMeans(n clusters=2).fit(reduced data)
clusters kmeans 3 = KMeans(n clusters=3).fit(reduced data)
clusters kmeans 4 = KMeans(n clusters=4).fit(reduced data)
# The chosen cluster -> GMM 2...
print 'The chosen cluster variable -> GMM 2...'
print clusters
The chosen cluster variable -> GMM 2...
GMM(covariance type='diag', init params='wmc', min covar=0.001,
  n components=2, n init=1, n iter=100, params='wmc', random state=Non
 thresh=None, tol=0.001)
In [11]:
# Plot the decision boundary by building a mesh grid to populate a graph.
x_{min}, x_{max} = reduced_data[:, 0].min() - 1, <math>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.

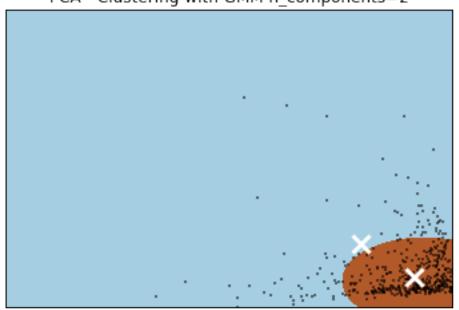
Z = clusters.predict(np.c [xx.ravel(), yy.ravel()])

```
In [12]:
# TODO: Find the centroids for KMeans or the cluster means for GMM
# GMM - testing/comparing...
centroids GMM components 1 = clusters GMM components 1.means
centroids GMM components 2 = clusters GMM components 2.means
centroids GMM components 3 = clusters GMM components 3.means
centroids_GMM_components_4 = clusters_GMM_components_4.means_
centroids = clusters.means
# KMeans - testing/comparing...
centroids KMeans clusters_1 = clusters_kmeans_1.cluster_centers_
centroids KMeans clusters 2 = clusters kmeans 2.cluster centers
centroids KMeans clusters 3 = clusters kmeans 3.cluster centers
centroids KMeans clusters 4 = clusters kmeans 4.cluster centers
#centroids = clusters.cluster centers
# The chosen centroid variable -> GMM 2...
print 'The chosen centroid variable -> GMM 2...'
print centroids
The chosen centroid variable -> GMM 2...
   3308.39301792 -3017.017396981
[-10810.23008886 9858.15532401]]
In [13]:
### testing GMM with various n components ###
# test GMM with n components = 2
clusters = clusters_GMM_components_2
Z = clusters.predict(np.c [xx.ravel(), yy.ravel()])
centroids = centroids GMM components 2
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('PCA - Clustering with GMM n components=2')
plt.xlim(x min, x max)
plt.ylim(y min, y max)
plt.xticks(())
plt.yticks(())
plt.show()
nrint 'centroids GMM components ?
```

```
Centrotab_dim_Components_
print centroids_GMM_components_2
print '\n'
# test GMM with n components = 3
clusters = clusters GMM components 3
Z = clusters.predict(np.c [xx.ravel(), yy.ravel()])
centroids = centroids_GMM_components_3
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('PCA - Clustering with GMM n components=3')
plt.xlim(x_min, x_max)
plt.ylim(y min, y max)
plt.xticks(())
plt.yticks(())
plt.show()
print 'centroids GMM components 3'
print centroids_GMM_components_3
print '\n'
# test GMM with n components = 4
clusters = clusters GMM components 4
Z = clusters.predict(np.c_[xx.ravel(), yy.ravel()])
centroids = centroids_GMM_components_4
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('PCA - Clustering with GMM n components=4')
plt.xlim(x_min, x_max)
plt.ylim(y min, y max)
plt.xticks(())
plt.yticks(())
plt.show()
nwint 'controids CMM components A
```

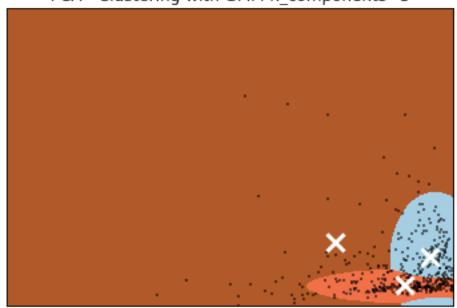
```
print centroids_GMM_components_4
print '\n'
```

PCA - Clustering with GMM n_components=2



```
centroids_GMM_components_2...
[[-10810.23008886 9858.15532401]
[ 3308.39301792 -3017.01739698]]
```

PCA - Clustering with GMM n_components=3



PCA - Clustering with GMM n_components=4

```
× × ×
```

```
centroids_GMM_components_4
[[ 7182.42527042    5428.27138593]
  [-15313.48033345    -3338.2980532 ]
  [ -9418.46902103    34454.41705257]
  [ 2336.74338421    -6721.26945836]]
```

In [14]:

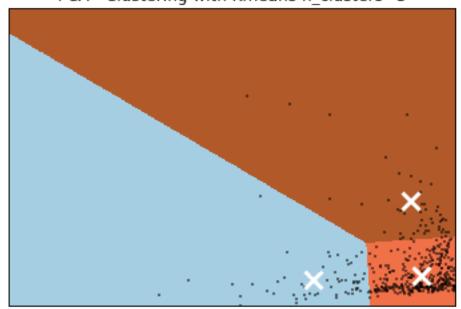
```
### testing K-means with various n clusters ###
# test K-means with n clusters = 2
clusters = clusters kmeans 2
Z = clusters.predict(np.c [xx.ravel(), yy.ravel()])
centroids = centroids_KMeans_clusters_2
Z = Z.reshape(xx.shape)
plt.figure(1)
plt.clf()
plt.imshow(Z, interpolation='nearest',
           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('PCA - Clustering with Kmeans n clusters=2')
plt.xlim(x_min, x_max)
plt.ylim(y min, y max)
plt.xticks(())
plt.yticks(())
plt.show()
print 'centroids Kmeans clusters 2...'
print centroids_KMeans_clusters_2
print '\n'
# test K-means with n clusters = 3
clusters = clusters kmeans 3
Z = clusters.predict(np.c [xx.ravel(), yy.ravel()])
centroids = centroids_KMeans_clusters_3
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('PCA - Clustering with Kmeans n clusters=3')
plt.xlim(x_min, x_max)
plt.ylim(y min, y max)
plt.xticks(())
plt.yticks(())
plt.show()
print 'centroids Kmeans clusters 3...'
print centroids KMeans clusters 3
print '\n'
# test K-means with n clusters = 4
clusters = clusters kmeans 4
Z = clusters.predict(np.c [xx.ravel(), yy.ravel()])
centroids = centroids KMeans clusters 4
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('PCA - Clustering with Kmeans n clusters=4')
plt.xlim(x min, x max)
plt.ylim(y_min, y_max)
plt.xticks(())
plt.yticks(())
plt.show()
print 'centroids Kmeans clusters 4...'
print centroids KMeans clusters 4
print '\n'
```



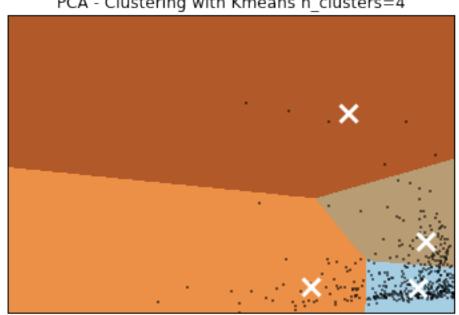
```
centroids Kmeans clusters 2...
[[-24088.33276689
                   1218.17938291]
    4175.31101293
                   -211.15109304]]
```

PCA - Clustering with Kmeans n_clusters=3



```
centroids_Kmeans_clusters_3...
[[-23978.86566553 -4445.56611772]
    4165.1217824
                  -3105.15811456]
    1341.31124554 25261.39189714]]
```

PCA - Clustering with Kmeans n_clusters=4



```
centroids_Kmeans_clusters_4...
    3542.08605212 -4936.7212132 ]
] ]
```

```
In [15]:
# The chosen centroid variable -> GMM 2...
#centroids = clusters.means
centroids = centroids GMM components 2
print 'The chosen centroid variable -> GMM 2...'
print centroids
# The chosen one...
clusters = GMM(n components=2).fit(reduced_data)
Z = clusters.predict(np.c [xx.ravel(), yy.ravel()])
centroids = clusters.means
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('PCA - Clustering with GMM n components=2')
plt.xlim(x_min, x_max)
plt.ylim(y_min, y_max)
plt.xticks(())
plt.yticks(())
plt.show()
print 'The above graph seems to represent data and the two different types of custor
The chosen centroid variable -> GMM 2...
```

```
[ 3308.39301792 -3017.01739698]]

PCA - Clustering with GMM n_components=2
```

9858.15532401]

[[-10810.23008886

5710.98964991 12661.45687292]

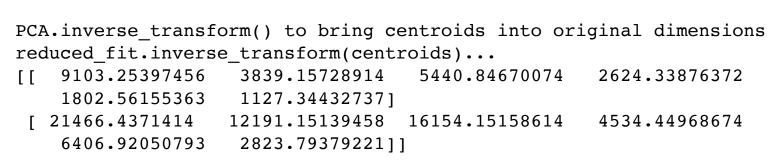
[-24220.71188261 -4364.45560022] [-14537.71774395 61715.67085248]]

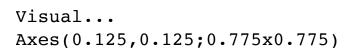


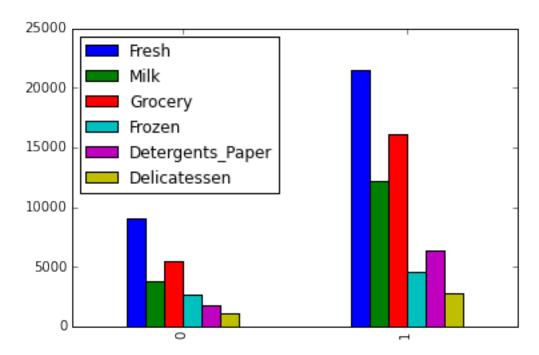
The above graph seems to represent data and the two different types of customers the most appropriately.

```
In [16]:
```

```
# PCA.inverse transform() to bring centroids into original dimensions...
reduced_fit = PCA(n_components=2).fit(data)
reduced data = reduced fit.transform(data)
print 'Chosen centroids...'
print centroids
print '\n'
print 'PCA.inverse transform() to bring centroids into original dimensions'
print 'reduced fit.inverse transform(centroids)...'
print reduced fit.inverse transform(centroids)
print '\n'
print 'Visual...'
print pd.DataFrame(reduced fit.inverse transform(centroids),columns=data.columns).pl
Chosen centroids...
    3308.39301792 -3017.01739698]
 [-10810.23008886]
                    9858.15532401]]
```







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

Answer:

The central objects represent the average customer in each of the clusters. Given the initial information about the types of clients in the assignment briefing, the first cluster represents the high-volume customers and the second cluster represents the smaller family-run shops. The tighter cluster located at the bottom right corner of the graph represents the smaller family-run shops. The "higher-variation" cluster that occupies the rest of the graph represents the high-volume customers.

Using the bar plot above, the high-volume customers buy a lot more "Fresh" products than the smaller-run family shops. Overall, "Fresh", "Milk", and "Grocery" products visibly show the greatest difference between high-volume and smaller family-run shops. The other three products, "Frozen", Detergents_paper, and "Delicatessen", are ordered in larger volumes by the high-volume customers but the difference is not as great.

###Conclusions

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

Answer:

I feel that PCA coupled with GMM gave the most insight into the data.

First, the PCA algorithm was able to create latent features and, in turn, we were able to reduce the amount of data into two principal components with minimal data loss. Reducing the data to two principal components enabled better visualization.

side note: PCA enables better visualization but not better clustering. Similar clusters would be generated with the original data. And with the original data, the less prevalent features that are not part of the first two PCA's would also be taken into account.

Because the data didn't seem to have a clear divide between to the known types of customers, GMM was a better technique than k-means. Clustering the data between 2 groups with GMM helped visualize and relate the data back to the two main customer groups (high-volume and smaller family-run shops).

9) How would you use that technique to help the company design new experiments?

Answer:

	*		High-Volume		Small Family-run	*		
	*		Customers		Businesses	*		
	*					*		
	* Morning Delivery (Variant A)		[Data]		[Data]	*	<	
_	Control groups							
	*					*		
	* Evening Delivery (Variant B)		[Data]		[Data]	*	<	
_	Experiment groups							
	********	* * *	*****	***	******	**		

note: only one variable can be tested at a time (to prevent bias in the experiment)

Now that we have the data seperated into two defined customer segments, new experiments can be implemented within a subset of customers within each of the segments for validation. For example, in the above graph, the control groups continue to receive morning deliveries while a subset of each customer segment will receive evening deliveries. If the change shows promise, for one or both customers, the change can then be implemented. Given the initial information about the types of clients, high-volume customers and smaller family run shops, I would predict that this exercise would result in no significant decline in satisfaction for high-volume customers yet a signicant decline in satisfaction from the smaller family-run shops.

10) How would you use that data to help you predict future customer needs?

Answer:

With customer segments defined, there is now the ability to apply supervised learning techniques to better understand the behaviours and financial results between the customer segments. For example, the data can be used drive strategic decisions to focus on one of the customer segments over the other. Overall, the data can assist in strategic decision-making with the support of calculated predictions. Coupled with the A/B test technique, the calculated decisions can be validated for implementation.

With customer segments defined, supervised learning techniques, such as classification and regression, can be applied. Classification learning algorithms can be used to relate a new customer, and their corresponding consumption elements, to one of the clusters that was generated from the unsupervised learning technique. For regression, the clusters can be used to predict purchase orders of future wholesale customers.