

```
In [1]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.cluster import KMeans
from sklearn import preprocessing
import mpl_toolkits

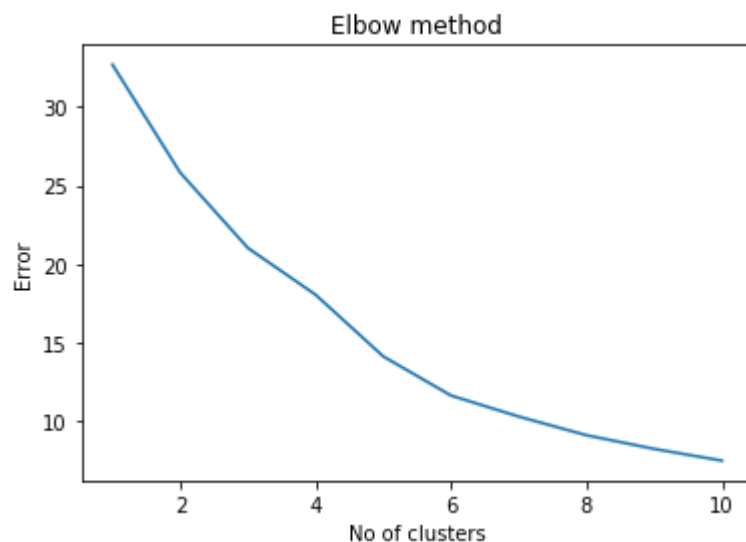
# Load data
df = pd.read_csv("cereal.csv")

# Replace all '-1' with NaN and remove
df.replace(-1, np.NaN, inplace=True)
df.dropna(inplace = True)
```

```
In [2]: X = df[["calories", "protein", "fat", "sodium", "fiber", "carbo", "sugars", "potas"]]
```

```
In [3]: min_max_scaler = preprocessing.MinMaxScaler()
norm_X = pd.DataFrame(min_max_scaler.fit_transform(X.values), columns = X.columns)
```

```
In [4]: Error = []
for i in range(1, 11):
    kmeans = KMeans(n_clusters = i).fit(norm_X)
    kmeans.fit(norm_X)
    Error.append(kmeans.inertia_)
import matplotlib.pyplot as plt
plt.plot(range(1, 11), Error)
plt.title('Elbow method')
plt.xlabel('No of clusters')
plt.ylabel('Error')
plt.show()
```



```
In [5]: kmeans2 = KMeans(n_clusters=2)
y_kmeans2 = kmeans2.fit_predict(norm_X)
print(y_kmeans2)
```

```
[0 1 0 0 1 1 1 0 0 1 0 1 1 1 0 0 1 1 1 0 1 0 1 1 0 1 1 1 1 0 0 0 1 1 1 0
 1 0 0 1 0 1 1 1 0 1 1 0 1 1 0 0 0 0 1 1 0 0 0 0 0 0 1 0 0 0 1 0 0 1 0 0 1]
```

```
In [6]: d = {'Cereal':df['name'], 'Cluster': y_kmeans2, 'Rating': df['rating'], 'Calories': df['calories'],
test = pd.DataFrame(data=d)
```

```
In [32]: test.head()
```

```
Out[32]:
```

	Cereal	Cluster	Rating	Calories	Protein	Fat	Sodium	Fiber	Carbohydrates	Sugars	Pro
0	100% Bran	0	68.402973	70	4	1	130	10.0	5.0	6.0	
1	100% Natural Bran	1	33.983679	120	3	5	15	2.0	8.0	8.0	
2	All-Bran	0	59.425505	70	4	1	260	9.0	7.0	5.0	
3	All-Bran with Extra Fiber	0	93.704912	50	4	0	140	14.0	8.0	0.0	
5	Apple Cinnamon Cheerios	1	29.509541	110	2	2	180	1.5	10.5	10.0	

```
In [8]: df['rating'].median()
```

```
Out[8]: 40.2530865
```

```
In [9]: cluster_1 = test[test['Cluster'] == 0]
med_1 = cluster_1['Rating'].median()
mean_1 = cluster_1['Rating'].mean()

print("Median of cluster 1: %f" %med_1)
print("Mean of cluster 1: %f" %mean_1)
```

```
Median of cluster 1: 50.764999
Mean of cluster 1: 52.573632
```

```
In [10]: cluster_2 = test[test['Cluster'] == 1]
med_2 = cluster_2['Rating'].median()
mean_2 = cluster_2['Rating'].mean()

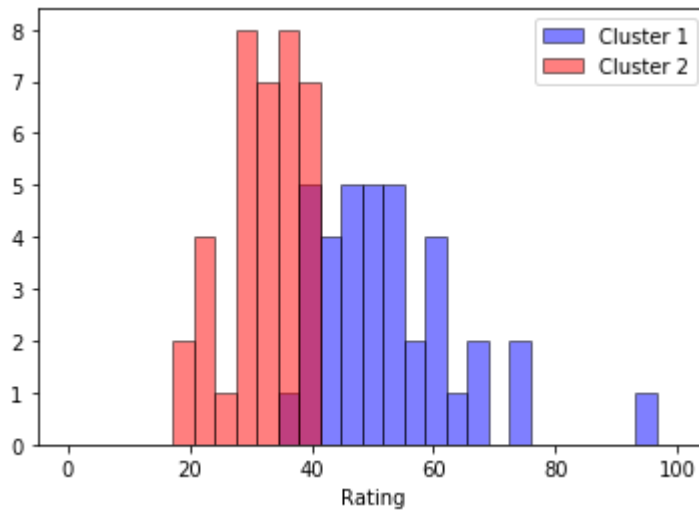
print("Median of cluster 2: %f" %med_2)
print("Mean of cluster 2: %f" %mean_2)
```

```
Median of cluster 2: 32.207582
Mean of cluster 2: 32.169942
```

```
In [12]: from matplotlib import pyplot

bins = np.linspace(0, 100, 30)

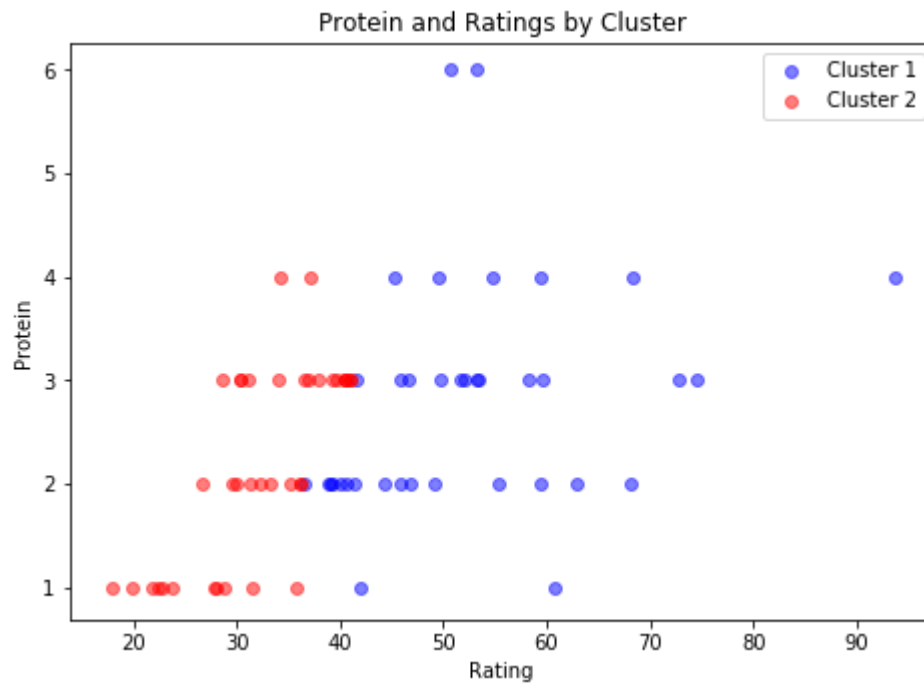
pyplot.hist(cluster_1['Rating'], bins, alpha=0.5, edgecolor='black', color = 'blue')
pyplot.hist(cluster_2['Rating'], bins, alpha=0.5, edgecolor='black', color = 'red')
pyplot.legend(loc='upper right')
pyplot.xlabel('Rating')
pyplot.show()
```



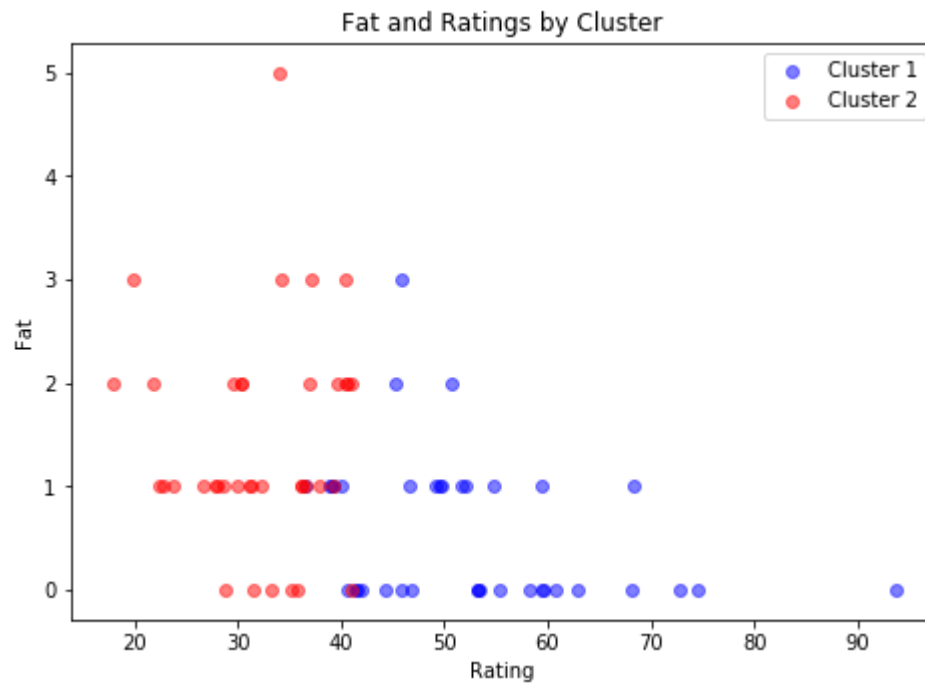
```
In [14]: fig=plt.figure()
ax=fig.add_axes([0,0,1,1])
ax.scatter(cluster_1['Rating'], cluster_1['Calories'], color='blue', alpha=0.5, label='Cluster 1')
ax.scatter(cluster_2['Rating'], cluster_2['Calories'], color='red', alpha=0.5, label='Cluster 2')
ax.set_xlabel('Rating')
ax.set_ylabel('Calories')
ax.set_title('Calories and Ratings by Cluster')
pyplot.legend(loc='upper right')
plt.show()
```



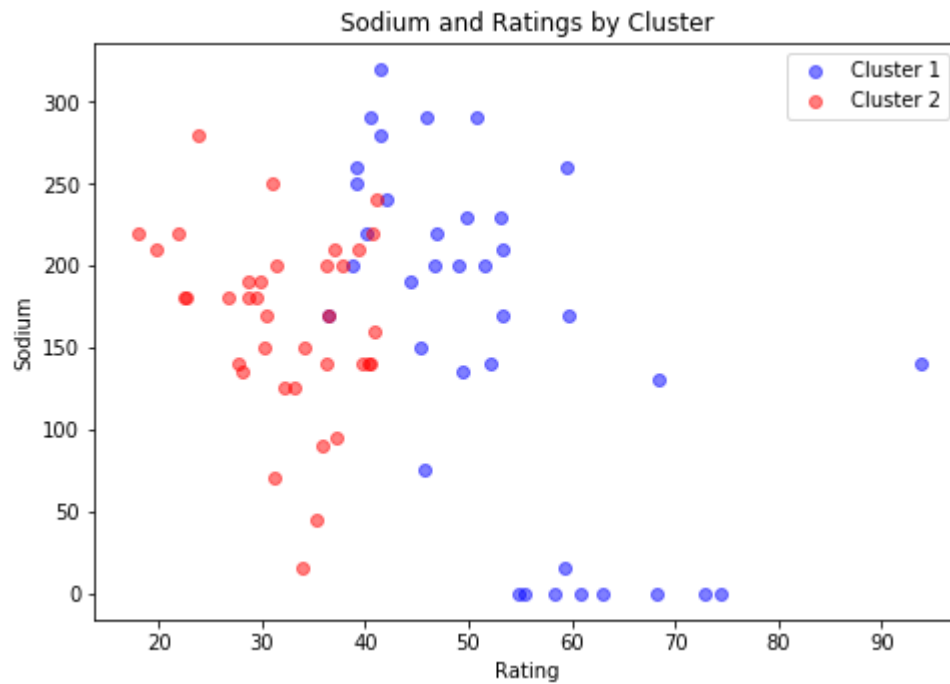
```
In [18]: fig=plt.figure()
ax=fig.add_axes([0,0,1,1])
ax.scatter(cluster_1['Rating'], cluster_1['Protein'], color='blue', alpha=0.5, label='Cluster 1')
ax.scatter(cluster_2['Rating'], cluster_2['Protein'], color='red', alpha=0.5, label='Cluster 2')
ax.set_xlabel('Rating')
ax.set_ylabel('Protein')
ax.set_title('Protein and Ratings by Cluster')
pyplot.legend(loc='upper right')
plt.show()
```



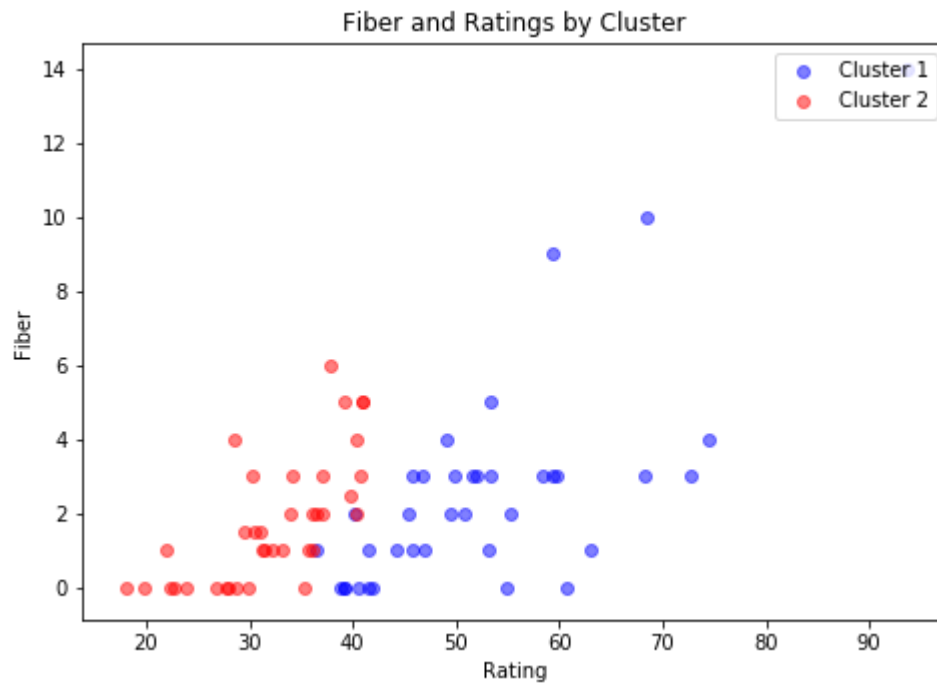
```
In [23]: fig=plt.figure()
ax=fig.add_axes([0,0,1,1])
ax.scatter(cluster_1['Rating'], cluster_1['Fat'], color='blue', alpha=0.5, label='Cluster 1')
ax.scatter(cluster_2['Rating'], cluster_2['Fat'], color='red', alpha=0.5, label='Cluster 2')
ax.set_xlabel('Rating')
ax.set_ylabel('Fat')
ax.set_title('Fat and Ratings by Cluster')
pyplot.legend(loc='upper right')
plt.show()
```



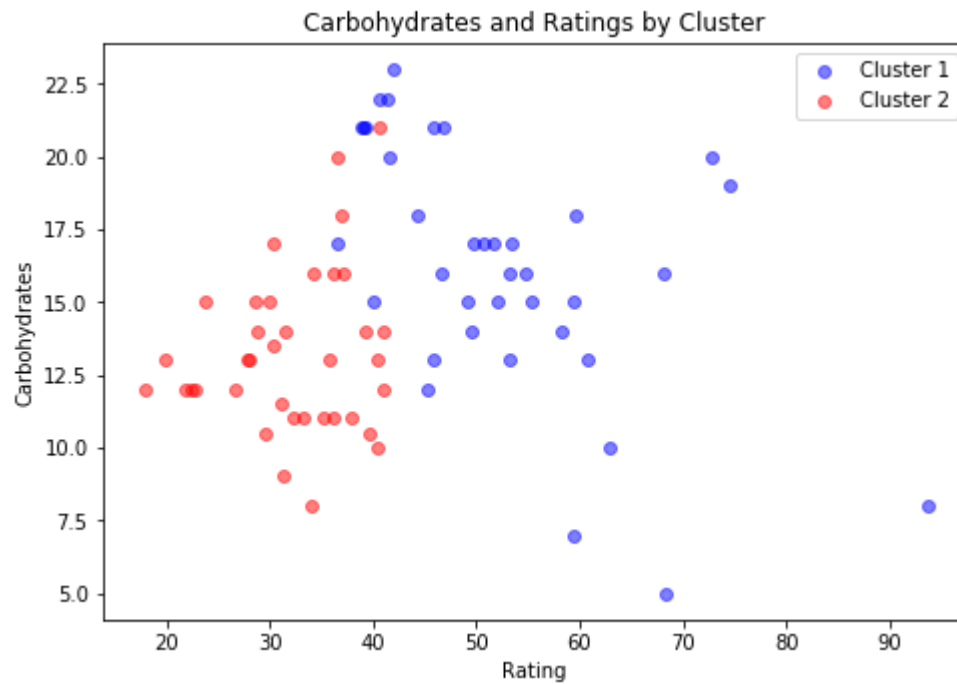
```
In [25]: fig=plt.figure()
ax=fig.add_axes([0,0,1,1])
ax.scatter(cluster_1['Rating'], cluster_1['Sodium'], color='blue', alpha=0.5, label='Cluster 1')
ax.scatter(cluster_2['Rating'], cluster_2['Sodium'], color='red', alpha=0.5, label='Cluster 2')
ax.set_xlabel('Rating')
ax.set_ylabel('Sodium')
ax.set_title('Sodium and Ratings by Cluster')
pyplot.legend(loc='upper right')
plt.show()
```



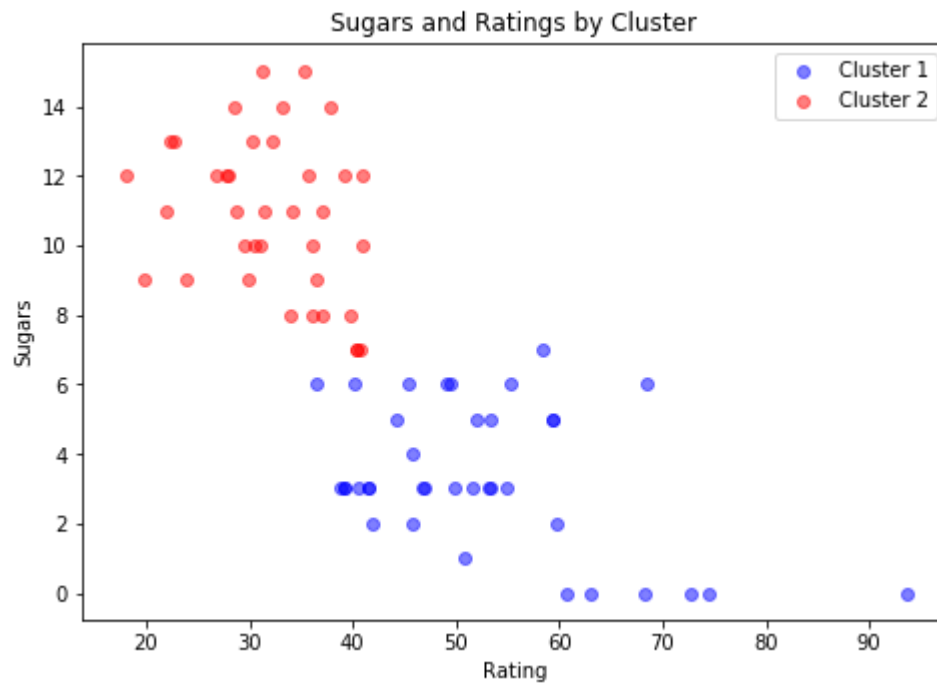
```
In [26]: fig=plt.figure()
ax=fig.add_axes([0,0,1,1])
ax.scatter(cluster_1['Rating'], cluster_1['Fiber'], color='blue', alpha=0.5, label='Cluster 1')
ax.scatter(cluster_2['Rating'], cluster_2['Fiber'], color='red', alpha=0.5, label='Cluster 2')
ax.set_xlabel('Rating')
ax.set_ylabel('Fiber')
ax.set_title('Fiber and Ratings by Cluster')
pyplot.legend(loc='upper right')
plt.show()
```



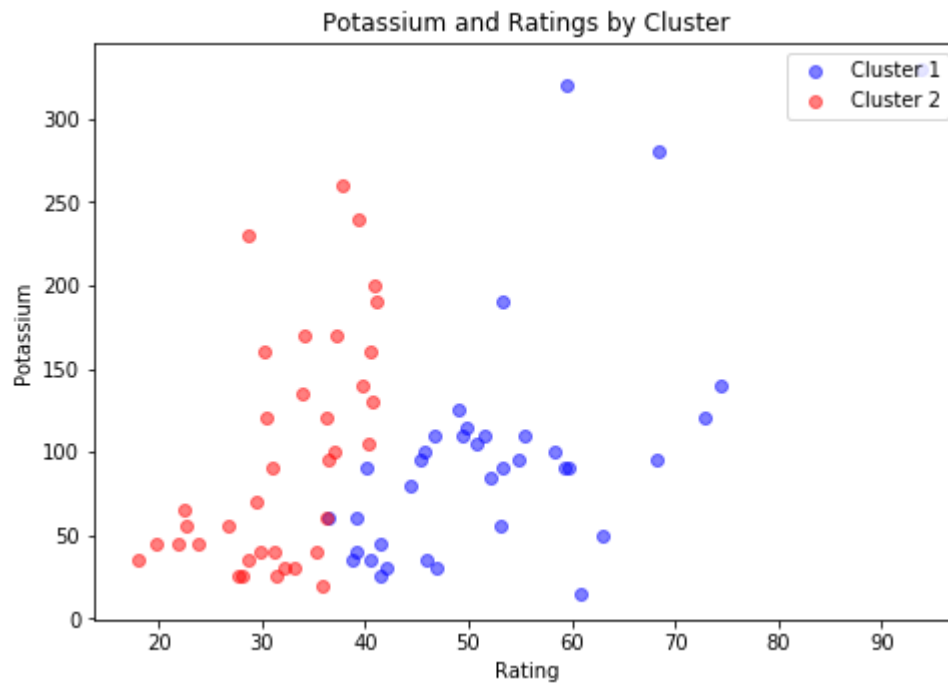

```
In [27]: fig=plt.figure()
ax=fig.add_axes([0,0,1,1])
ax.scatter(cluster_1['Rating'], cluster_1['Carbohydrates'], color='blue', alpha=0.5)
ax.scatter(cluster_2['Rating'], cluster_2['Carbohydrates'], color='red', alpha=0.5)
ax.set_xlabel('Rating')
ax.set_ylabel('Carbohydrates')
ax.set_title('Carbohydrates and Ratings by Cluster')
pyplot.legend(loc='upper right')
plt.show()
```



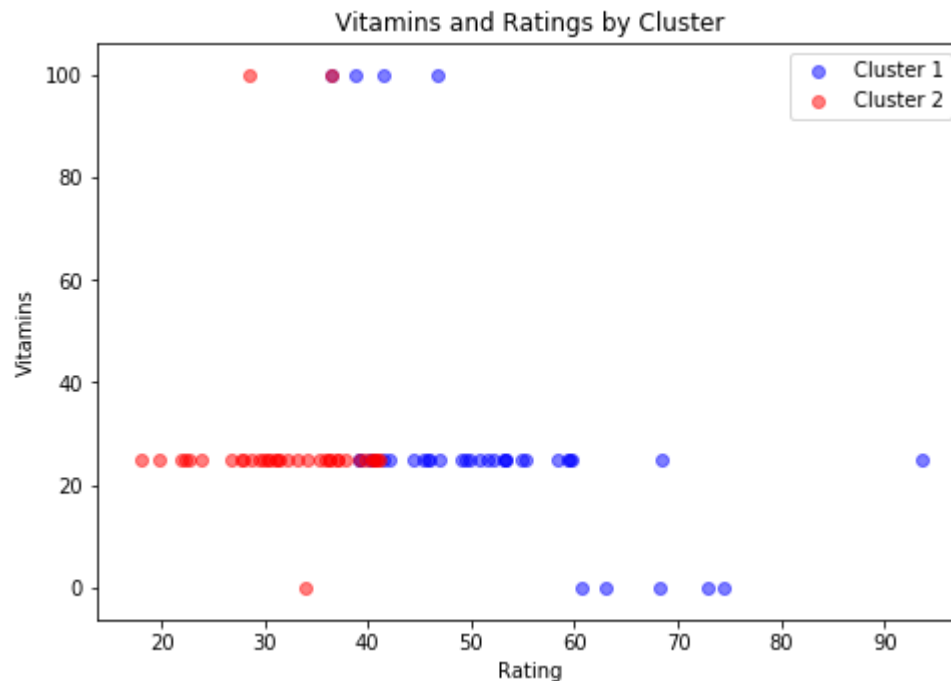
```
In [28]: fig=plt.figure()
ax=fig.add_axes([0,0,1,1])
ax.scatter(cluster_1['Rating'], cluster_1['Sugars'], color='blue', alpha=0.5, label='Cluster 1')
ax.scatter(cluster_2['Rating'], cluster_2['Sugars'], color='red', alpha=0.5, label='Cluster 2')
ax.set_xlabel('Rating')
ax.set_ylabel('Sugars')
ax.set_title('Sugars and Ratings by Cluster')
pyplot.legend(loc='upper right')
plt.show()
```



```
In [29]: fig=plt.figure()
ax=fig.add_axes([0,0,1,1])
ax.scatter(cluster_1['Rating'], cluster_1['Potassium'], color='blue', alpha=0.5, )
ax.scatter(cluster_2['Rating'], cluster_2['Potassium'], color='red', alpha=0.5, )
ax.set_xlabel('Rating')
ax.set_ylabel('Potassium')
ax.set_title('Potassium and Ratings by Cluster')
pyplot.legend(loc='upper right')
plt.show()
```



```
In [30]: fig=plt.figure()
ax=fig.add_axes([0,0,1,1])
ax.scatter(cluster_1['Rating'], cluster_1['Vitamins'], color='blue', alpha=0.5, label='Cluster 1')
ax.scatter(cluster_2['Rating'], cluster_2['Vitamins'], color='red', alpha=0.5, label='Cluster 2')
ax.set_xlabel('Rating')
ax.set_ylabel('Vitamins')
ax.set_title('Vitamins and Ratings by Cluster')
pyplot.legend(loc='upper right')
plt.show()
```



```
In [34]: test_good = test.loc[test['Cluster'] == 0]
test_bad = test.loc[test['Cluster'] == 1]
```

```
In [58]: # Sort by highest rating first
test_good = test_good.sort_values('Rating', ascending=False)
# Sort by lowest rating first
test_bad = test_bad.sort_values('Rating')
```

```
In [61]: print("TOP 10 GOOD FOR YOU\n", test_good['Cereal'][0:10])
print("\n")
print("TOP 10 BAD FOR YOU\n", test_bad['Cereal'][0:10])
```

TOP 10 GOOD FOR YOU

```
3      All-Bran with Extra Fiber
64     Shredded Wheat 'n'Bran
65     Shredded Wheat spoon size
0      100% Bran
63     Shredded Wheat
55     Puffed Wheat
54     Puffed Rice
50     Nutri-grain Wheat
2      All-Bran
68     Strawberry Fruit Wheats
Name: Cereal, dtype: object
```

TOP 10 BAD FOR YOU

```
10     Cap'n'Crunch
12     Cinnamon Toast Crunch
35     Honey Graham Ohs
18     Count Chocula
14     Cocoa Puffs
31     Golden Grahams
42     Lucky Charms
73     Trix
29     Fruity Pebbles
70     Total Raisin Bran
Name: Cereal, dtype: object
```

```
In [ ]:
```