Mean Shift Clustering Unsupervised

Mean shift builds upon the concept of kernel density estimation (KDE). Imagine that the data was sampled from a probability distribution. KDE is a method to estimate the underlying distribution (also called the probability density function) for a set of data.

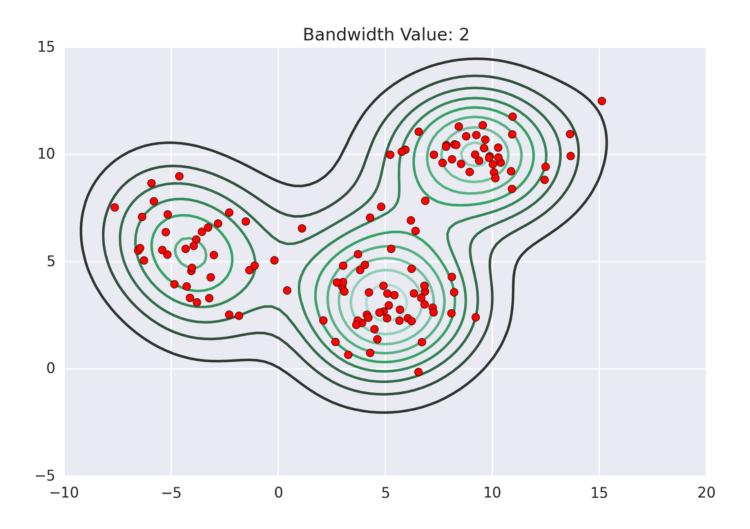
It works by placing a kernel on each point in the data set. A kernel is a fancy mathematical word for a weighting function. There are many different types of kernels, but the most popular one is the Gaussian kernel. Adding all of the individual kernels up generates a probability surface (e.g., density function). Depending on the kernel bandwidth parameter used, the resultant density function will vary.

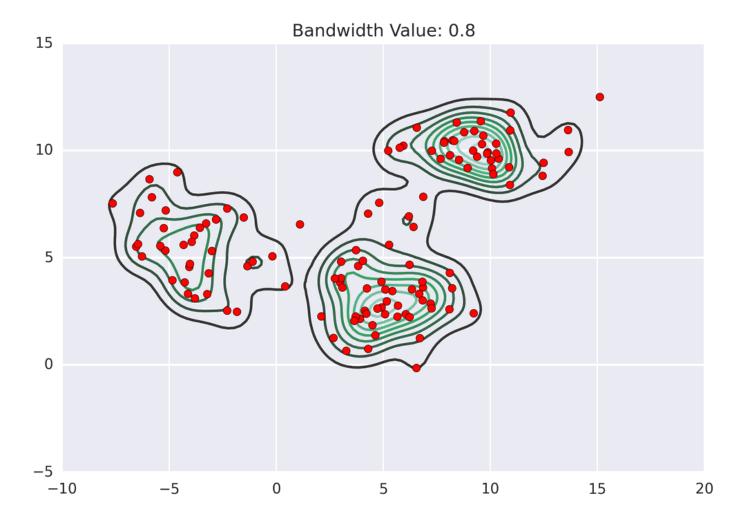
Mean shift exploits this KDE idea by imagining what the points would do if they all climbed up hill to the nearest peak on the KDE surface. It does so by iteratively shifting each point uphill until it reaches a peak.

Depending on the kernel bandwidth used, the KDE surface (and end clustering) will be different. As an extreme case, imagine that we use extremely tall skinny kernels (e.g., a small kernel bandwidth). The resultant KDE surface will have a peak for each point. This will result in each point being placed into its own cluster. On the other hand, imagine that we use an extremely short fat kernels (e.g., a large kernel bandwidth). This will result in a wide smooth KDE surface with one peak that all of the points will climb up to, resulting in one cluster. Kernels in between these two extremes will result in nicer clusterings. Below are two animations of mean shift running for different kernel bandwidth values.

Iteratively playing around with the bandwith I was able to get two clusters when the bandwidth was 83. This bandwidth parameter will vary depending on the dataset. Therefore for prediction, entire datasets need to be run from the beginning and fitted with meanshift clustering.

This algorithm is parameter sensitive. it is one of the main drawbacks of using it





The top animation results in three KDE surface peaks, and thus three clusters. The second animation uses a smaller kernel bandwidth, and results in more than three clusters. As with all clustering problems, there is no correct clustering. Rather, correct is usually defined by what seems reasonable given the problem domain and application. Mean shift provides one nice knob (the kernel bandwidth parameter) that can easily be tuned appropriately for different applications.

```
In [1]: | from sklearn import datasets
        from sklearn import preprocessing
        from sklearn.model selection import train test split
        from sklearn import cluster
        from sklearn.cluster import KMeans
        from sklearn.ensemble import IsolationForest
        from sklearn.neighbors import LocalOutlierFactor
        from sklearn import svm, neighbors
        from sklearn.neighbors import NearestNeighbors
        from sklearn.cluster import MeanShift, estimate_bandwidth
        from sklearn.metrics import classification_report
        from sklearn.metrics import confusion matrix
        from sklearn.metrics import recall score
        from sklearn.metrics import roc auc score
        from sklearn.model selection import GridSearchCV
        from sklearn.metrics import make scorer
        from sklearn.metrics import accuracy score
        import pandas as pd
        import numpy as np
        import itertools
        import matplotlib.pyplot as plt
        import datetime
        %matplotlib inline
```

Out[2]: (237024, 58)

```
In [3]: column_list=df.columns.tolist()
```

In [4]: df.head()

Out[4]:

	originalloanamount	originalloanterm	originalinterestratepercentage	graceperiodnuml
0	66711.84	60	3.29	1
1	16258.45	60	0.90	0
2	31930.41	72	2.90	1
3	26065.02	65	0.90	0
4	42091.00	72	3.90	0

5 rows × 58 columns

In [5]: # prepare label for scikit-learn
Y=df.label.values
Y.shape

Out[5]: (237024,)

In [6]: # prepare input data for scikit-learn
input=df.values
input.shape

Out[6]: (237024, 58)

In [7]: # calculate train/test split
 len_train = int(len(input)*train_split)
 print(len_train)

189619

```
In [8]: # apply train/test split to labels
    y_train = Y[0:len_train]
    y_test = Y[len_train:]
    x_train = input[0:len_train]
    x_test = input[len_train:]
    x_train.shape
```

Out[8]: (189619, 58)

In [9]: export_x_test = pd.DataFrame(data=x_test)

```
In [10]: export_x_test.columns=column_list
    export_x_test.rename(columns={'label':'True Label'}, inplace=True)
    export_x_test.head()
```

Out[10]:

	originalloanamount	originalloanterm	originalinterestratepercentage	graceperiodnuml
0	36863.24	72.0	1.00	1.0
1	23811.32	60.0	1.90	0.0
2	30669.00	48.0	1.00	1.0
3	54083.21	72.0	1.00	0.0
4	31557.75	72.0	3.89	1.0

5 rows × 58 columns

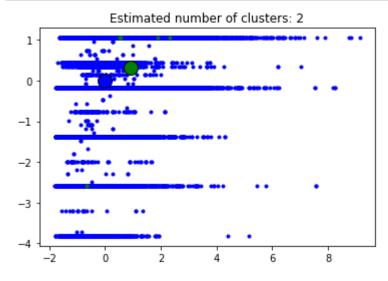
```
In [11]: #from sklearn.preprocessing import MinMaxScaler
    # from sklearn.preprocessing import minmax_scale
    # from sklearn.preprocessing import MaxAbsScaler
    from sklearn.preprocessing import StandardScaler
    # from sklearn.preprocessing import RobustScaler
    # from sklearn.preprocessing import Normalizer
    # from sklearn.preprocessing import QuantileTransformer
    # from sklearn.preprocessing import PowerTransformer
```

```
In [12]: x_scaler=StandardScaler()
    x_train = x_scaler.fit_transform(x_train)
    x_test = x_scaler.fit_transform(x_test)
```

- In [13]: #bandwidth = estimate_bandwidth(x_test, quantile=0.2, n_samples=1000)
 bandwidth=83
 ms = cluster.MeanShift(bandwidth=bandwidth, bin_seeding=True)
 clf_ms= ms.fit(x_test)
- In [14]: x_pred = x_test
- In [15]: prediction_ms = clf_ms.predict(x_pred)
- In [16]: prediction_ms = ms.labels_
 cluster_centers = ms.cluster_centers_

 labels_unique = np.unique(prediction_ms)
 n_clusters_ = len(np.unique(prediction_ms))
- In [17]: print('Number of clusters :', n_clusters_)

Number of clusters: 2



```
In [19]: np.unique(prediction_ms)
Out[19]: array([0, 1], dtype=int64)
In [20]: np.bincount(np.array(prediction_ms).reshape(1,prediction_ms.size)[0])
Out[20]: array([47400, 5], dtype=int64)
In [21]: export_x_test['Predicted Label']=prediction_ms
```

In [22]: export_x_test.head()

Out[22]:

	originalloanamount	originalloanterm	originalinterestratepercentage	graceperiodnuml
0	36863.24	72.0	1.00	1.0
1	23811.32	60.0	1.90	0.0
2	30669.00	48.0	1.00	1.0
3	54083.21	72.0	1.00	0.0
4	31557.75	72.0	3.89	1.0

5 rows × 59 columns

```
In [23]: export x test.shape
Out[23]: (47405, 59)
In [24]: export x test.to csv(path+"prediction/ms/predicated ms abs loans "+str(nrows)+
         ".csv", chunksize=10000)
In [28]: def plot confusion matrix(cm, title, classes=['Current', 'Non Current'],
                                    cmap=plt.cm.Blues, save=False, saveas="MyFigure.png"
         ):
             # print Confusion matrix with blue gradient colours
             cm = cm.astype('float') / cm.sum(axis=1)[:, np.newaxis]
             plt.imshow(cm, interpolation='nearest', cmap=cmap)
             plt.title(title)
             plt.colorbar()
             tick marks = np.arange(len(classes))
             plt.xticks(tick marks, classes, rotation=90)
             plt.yticks(tick_marks, classes)
             fmt = '.1%'
             thresh = cm.max() / 2.
             for i, j in itertools.product(range(cm.shape[0]), range(cm.shape[1])):
                 plt.text(j, i, format(cm[i, j], fmt),
                          horizontalalignment="center",
                          color="white" if cm[i, j] > thresh else "black")
             plt.tight layout()
             plt.ylabel('True label')
             plt.xlabel('Predicted label')
```

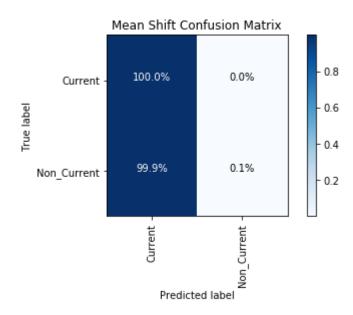
plt.savefig(saveas, dpi=100)

if save:

```
In [29]: def plot gridsearch cv(results, estimator, x min, x max, y min, y max, save=F
         alse, saveas="MyFigure.png"):
             # print GridSearch cross-validation for parameters
             plt.figure(figsize=(10,8))
             plt.title("GridSearchCV for "+estimator, fontsize=24)
             plt.xlabel(estimator)
             plt.ylabel("Score")
             plt.grid()
             ax = plt.axes()
             ax.set xlim(x min, x max)
             ax.set ylim(y min, y max)
             pad = 0.005
             X axis = np.array(results["param "+estimator].data, dtype=float)
             for scorer, color in zip(sorted(scoring), ['b', 'k']):
                 for sample, style in (('train', '--'), ('test', '-')):
                     sample_score_mean = results['mean_%s_%s' % (sample, scorer)]
                     sample score std = results['std %s %s' % (sample, scorer)]
                     ax.fill between(X axis, sample score mean - sample score std,
                                  sample score mean + sample score std,
                                  alpha=0.1 if sample == 'test' else 0, color=color)
                     ax.plot(X axis, sample score mean, style, color=color,
                         alpha=1 if sample == 'test' else 0.7,
                         label="%s (%s)" % (scorer, sample))
                 best_index = np.nonzero(results['rank_test_%s' % scorer] == 1)[0][0]
                 best_score = results['mean_test_%s' % scorer][best index]
                 # Plot a dotted vertical line at the best score for that scorer mark
         ed by x
                 ax.plot([X_axis[best_index], ] * 2, [0, best_score],
                     linestyle='-.', color=color, marker='x', markeredgewidth=3, ms=8
                 # Annotate the best score for that scorer
                 ax.annotate("%0.2f" % best_score,
                          (X axis[best index], best score+pad))
             plt.legend(loc="best")
             plt.grid('off')
             plt.tight layout()
             if save:
                 plt.savefig(saveas, dpi=100)
             plt.show()
```

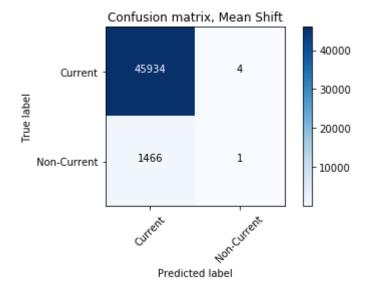
	precision	recall	f1-score	support
Current Non_Current	0.97 0.20	1.00 0.00	0.98 0.00	45938 1467
avg / total	0.95	0.97	0.95	47405

AUC: 50.0%



```
In [32]: | class_names = ['Current', 'Non-Current']
         def plot confusion matrix(cm, classes,
                                    normalize=False,
                                    title='Confusion matrix',
                                    cmap=plt.cm.Blues):
              .. .. ..
             This function prints and plots the confusion matrix.
             Normalization can be applied by setting `normalize=True`.
             if normalize:
                  cm = cm.astype('float') / cm.sum(axis=1)[:, np.newaxis]
                 print("Normalized confusion matrix")
             else:
                  print('Confusion matrix, without normalization')
             print(cm)
             plt.imshow(cm, interpolation='nearest', cmap=cmap)
             plt.title(title)
             plt.colorbar()
             tick_marks = np.arange(len(classes))
             plt.xticks(tick marks, classes, rotation=45)
             plt.yticks(tick_marks, classes)
             fmt = '.2f' if normalize else 'd'
             thresh = cm.max() / 2.
             for i, j in itertools.product(range(cm.shape[0]), range(cm.shape[1])):
                  plt.text(j, i, format(cm[i, j], fmt),
                           horizontalalignment="center",
                           color="white" if cm[i, j] > thresh else "black")
             plt.ylabel('True label')
             plt.xlabel('Predicted label')
             plt.tight_layout()
         print('ROC_AUC_SCORE ; ', roc_auc_score(y_test, prediction_ms))
         # Compute confusion matrix
         cnf matrix = confusion matrix(y test, prediction ms)
         np.set_printoptions(precision=2)
         # Plot non-normalized confusion matrix
         plt.figure()
         plot_confusion_matrix(cnf_matrix, classes=class_names, title= 'Confusion matri
         x, Mean Shift')
         plt.savefig('prediction/ms/cm'+str(' ms-')+str(nrows)+'.jpg')
         plt.show()
```

ROC_AUC_SCORE ; 0.500297294688
Confusion matrix, without normalization
[[45934 4]
 [1466 1]]



In []: