# **Identify Fraud From Enron Email Dataset**

In 2000, Enron was one of the largest companies in the United States in energy trading and was named as 'America's most innovative company'. By 2002, it had collapsed into bankruptcy due to widespread corporate fraud. In the resulting Federal investigation, a significant amount of typically confidential information entered into the public record, including tens of thousands of emails and detailed financial data for top executives. In this project, i have applied my machine learning skills by building a person of interest identifier based on financial and email data made public as a result of the Enron scandal. To assist, we've combined this data with a hand-generated list of persons of interest in the fraud case, which means individuals who were indicted, reached a settlement or plea deal with the government, or testified in exchange for prosecution immunity.

There are seven major steps in my project:

- 1. Load the Dataset and Query the dataset.
- 2. Outlier Detection and Removal
- 3. Feature Pre-processing
- 4. Classifier

In [348]:

count=0

- 5. Comparison of different classifier
- 6. Parameter Tuning
- 7. Validation of Classifier

## **Load The Dataset**

```
In [347]:
"""

Starter code for exploring the Enron dataset (emails + finances);
  loads up the dataset (pickled dict of dicts).

The dataset has the form:
  enron_data["LASTNAME FIRSTNAME MIDDLEINITIAL"] = { features_dict }

  {features_dict} is a dictionary of features associated with that person
.
  and here's an example to get you started:
  enron_data["SKILLING JEFFREY K"]["bonus"] = 5600000

"""

import pickle
enron_data = pickle.load(open("../final_project/final_project_dataset.pkl",
"r"))
print "There are "+str(len(enron_data))+" executives in Enron Dataset"

There are 146 executives in Enron Dataset
```

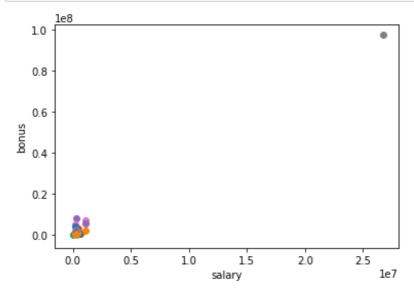
```
for i in enron data:
    if (enron data[i] ["poi"] == 1):
        count=count+1
print "There are "+str(count)+" Person of Interest(POI) and "+str((len(enro
n data))-(count))+" Non-POIs in our Dataset "
There are 18 Person of Interest (POI) and 128 Non-POIs in our Dataset
In [349]:
print "There are "+str(len(enron data["SKILLING JEFFREY K"]))+" features av
ailable for each person"
There are 21 features available for each person
In [403]:
print "The 21 features are listed below:"
features list=['poi']
for i in enron data["SKILLING JEFFREY K"]:
    print "Feature "+str(k)+": "+i
    if (i!='poi' and type(enron data["SKILLING JEFFREY K"][i]) == int):
        features list.append(i)
    k = k+1
print "Features list:"+str(features list)
The 21 features are listed below:
Feature 1: salary
Feature 2: to messages
Feature 3: deferral payments
Feature 4: total payments
Feature 5: exercised stock options
Feature 6: bonus
Feature 7: restricted stock
Feature 8: shared receipt with poi
Feature 9: restricted stock deferred
Feature 10: total stock value
Feature 11: expenses
Feature 12: loan advances
Feature 13: from messages
Feature 14: other
Feature 15: from this person to poi
Feature 16: poi
Feature 17: director fees
Feature 18: deferred income
Feature 19: long term incentive
Feature 20: email address
Feature 21: from_poi_to_this_person
Features list: ['poi', 'salary', 'to messages', 'total payments', 'exercised
_stock_options', 'bonus', 'restricted_stock', 'shared_receipt_with_poi', 't
otal_stock_value', 'expenses', 'from_messages', 'other',
'from this person to poi', 'long term incentive',
'from poi to this person']
```

## **Outlier Detection and Removal**

Just going through the Enron Dataset,I found Outlier when Bonus of people were plotted against the salary of person.

#### In [404]:

```
import pickle
import sys
import matplotlib.pyplot
sys.path.append("../tools/")
from feature format import featureFormat, targetFeatureSplit
### read in data dictionary, convert to numpy array
data_dict = pickle.load( open("../final_project/final_project_dataset.pkl",
"r") )
features = ["salary", "bonus"]
data = featureFormat(data dict, features,remove any zeroes=True)
for point in data:
    salary = point[0]
    bonus = point[1]
    matplotlib.pyplot.scatter(salary, bonus )
matplotlib.pyplot.xlabel("salary")
matplotlib.pyplot.ylabel("bonus")
matplotlib.pyplot.show()
```



## In [405]:

```
#finding the point of outlier
for key, value in data_dict.items():
    if value['bonus'] == data.max():
        print key
```

TOTAL

As it can be seen "TOTAL" is irrelavant point. Therefor is removed and grapg is replotted below.

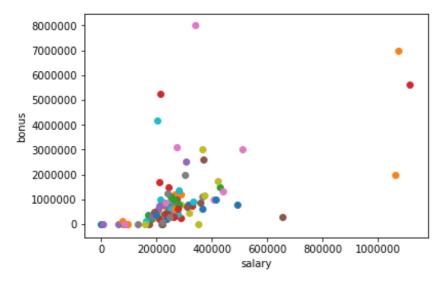
## In [406]:

```
data_dict.pop('TOTAL',0)
data = featureFormat(data_dict, features)

for point in data:
    salary = point[0]
```

```
bonus = point[1]
  matplotlib.pyplot.scatter( salary, bonus )

matplotlib.pyplot.xlabel("salary")
matplotlib.pyplot.ylabel("bonus")
matplotlib.pyplot.show()
```



### In [407]:

```
##other ouliers
outliers = []
for key in data_dict:
    val = data_dict[key]['salary']
    if val == 'NaN':
        continue

outliers.append((key,int(val)))

outliers_final = (sorted(outliers,key=lambda x:x[1],reverse=True)[:2])
outliers_final
```

```
Out[407]:
[('SKILLING JEFFREY K', 1111258), ('LAY KENNETH L', 1072321)]
```

These points cannot be removed from dataset as they are important people in Enron case and represent as the person of Interest(POI).

## Linear Regression to predict Bonus from salary

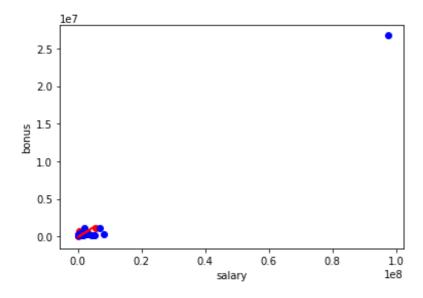
Now,To predict the bonus of an Employee when only salary of a person is only given.Linear Regression is used. In regression, you need training and testing data, just like in classification. We will see how outlier affect the Regression. Outlier Detection and Removal is a process which comprise of:

- 1. Train the dataset.
- 2. Identify the outlier and remove the points with Residual Error.
- 3. Re-Train the dataset.

```
In [408]:
```

```
## Training the data
data dict = pickle.load( open("../final project/final project dataset.pkl",
"r") )
data = featureFormat(data dict, features,remove any zeroes=True)
features = ["salary", "bonus"]
target, feature = targetFeatureSplit( data )
from sklearn.cross validation import train test split
feature train, feature test, target train, target test = train test split(f
eature, target, test_size=0.5, random_state=42)
from sklearn.linear model import LinearRegression as lr
reg=lr()
reg.fit(feature train, target train)
    matplotlib.pyplot.plot( feature test, reg.predict(feature test),color='
except NameError:
    pass
print reg.coef
print reg.score(feature test, target test)
import matplotlib.pyplot as plt
for feature, target in zip(feature test, target test):
    matplotlib.pyplot.scatter( feature, target, color="r")
for feature, target in zip(feature train, target train):
    matplotlib.pyplot.scatter( feature, target, color="b" )
matplotlib.pyplot.xlabel("salary")
matplotlib.pyplot.ylabel("bonus")
matplotlib.pyplot.show()
```

```
[ 0.27229528]
-0.877354252073
```



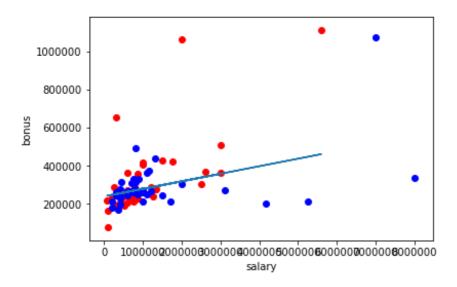
### In [409]:

```
### Identification of outlier
#!/usr/bin/python

data_dict.pop('TOTAL',0)
# data_dict.pop('LAVORATO JOHN J',0)
data = featureFormat(data_dict, features,remove_any_zeroes=True)
```

```
target, feature = targetFeatureSplit( data )
from sklearn.cross validation import train test split
feature train, feature test, target train, target test = train test split(f
eature, target, test size=0.5, random state=42)
from sklearn.linear model import LinearRegression as lr
reg=lr()
reg.fit(feature train, target train)
    matplotlib.pyplot.plot( feature test, reg.predict(feature test) )
except NameError:
    pass
print reg.coef
print reg.score(feature test , target test)
import matplotlib.pyplot as plt
for feature, target in zip(feature test, target test):
    matplotlib.pyplot.scatter( feature, target, color="r" )
for feature, target in zip(feature train, target train):
    matplotlib.pyplot.scatter( feature, target, color="b" )
matplotlib.pyplot.xlabel("salary")
matplotlib.pyplot.ylabel("bonus")
matplotlib.pyplot.show()
```

```
[ 0.03954061] 0.203020850473
```

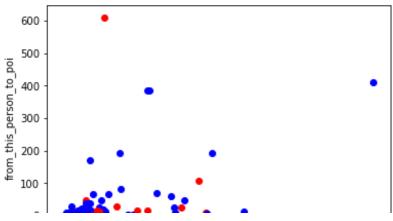


It can be observed how outlier affects the result of Regression. There is drastic difference between regression score with outlier and without outlier. Therefore outliers must be removed from dataset before any conclusions.

# **Feature Processing**

### **New Features**

```
# from sklearn.feature selection import SelectKBest, f classif
# selector = SelectKBest(f classif, k=10)
# selector.fit(features train, labels_train)
# features train transformed = selector.transform(features train).toarray()
# features test transformed = selector.transform(features test).toarray()
##New Features
def dict to list(key,normalizer):
    new list=[]
    for i in data dict:
        if data_dict[i][key] == "NaN" or data_dict[i][normalizer] == "NaN":
            new list.append(0.)
        elif data dict[i][key]>=0:
            new_list.append(float(data_dict[i][key])/float(data_dict[i][norr
alizer]))
    return new list
### create two lists of new features
fraction from poi email=dict to list("from poi to this person", "to messages
")
fraction to poi email=dict to list("from this person to poi", "from messages
### insert new features into data dict
count=0
for i in data dict:
    data dict[i]["fraction from poi email"]=fraction from poi email[count]
    data dict[i]["fraction to poi email"]=fraction to poi email[count]
    count +=1
### store to my dataset for easy export below
my dataset = data dict
for item in data dict:
    Fraction to=data dict[item]['from this person to poi']
    Fraction From=data dict[item]['from poi to this person']
    if (data dict[item]['poi']==1):
       matplotlib.pyplot.scatter( Fraction From, Fraction to,color='r')
    else:
       matplotlib.pyplot.scatter( Fraction From, Fraction to,color='b')
matplotlib.pyplot.xlabel("from poi to this person")
matplotlib.pyplot.ylabel("from this person to poi")
matplotlib.pyplot.show()
4
```

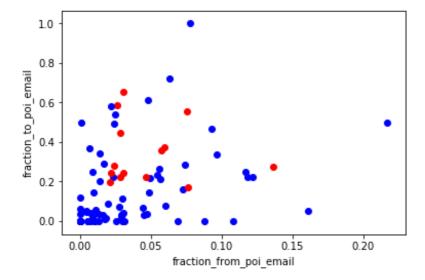


```
0 100 200 300 400 500 from poi to this person
```

When I picked 'from\_poi\_to\_this\_person' and 'from\_this\_person\_to\_poi' but there is was no strong pattern when I plotted the data so I used fractions for both features of "from/to poi messages" and "total from/to messages".

#### In [411]:

```
for item in data_dict:
    Fraction_to=data_dict[item]['fraction_to_poi_email']
    Fraction_From=data_dict[item]['fraction_from_poi_email']
    if (data_dict[item]['poi']==1):
        matplotlib.pyplot.scatter( Fraction_From, Fraction_to,color='r')
    else:
        matplotlib.pyplot.scatter( Fraction_From, Fraction_to,color='b')
matplotlib.pyplot.xlabel("fraction_from_poi_email")
matplotlib.pyplot.ylabel("fraction_to_poi_email")
matplotlib.pyplot.show()
```



Two new features were created and tested for this project. These were: ● the fraction of all emails to a person that were sent from a person of interest; ● the fraction of all emails that a person sent that were addressed to persons of interest. My assumption was that there is stronger connection between POI's via email then that between POI's and non-POI's. When we look at scatterplot we can agree that the data pattern confirms said above, e.i. there is no POI below 0.2 in "x" axis.

## **Classification Algorithm for Enron Dataset**

### In [412]:

```
##Naive Bayesian Classifier
data = featureFormat(data_dict, features_list)
labels, features = targetFeatureSplit(data)

from sklearn.cross_validation import train_test_split
features_train, features_test, labels_train, labels_test=train_test_split(feat ures, labels, test_size=0.3, random_state=42)
from sklearn.naive_bayes import GaussianNB
from time import time
```

```
TIOM CIME IMPOIL CIME
t0 = time()
clf = GaussianNB()
clf.fit(features train, labels train)
pred = clf.predict(features test)
accuracy = accuracy_score(pred,labels_test)
print "Accuracy when using Naive Bayes Classifier:"+str(accuracy)
print "Precision: " +str(precision_score(pred, labels_test))
print "Recall: "+str(recall score(pred, labels test))
print "NB algorithm time:", round(time()-t0, 3), "s"
Accuracy when using Naive Bayes Classifier: 0.590909090909
Precision: 0.5
Recall: 0.22222222222
NB algorithm time: 0.009 s
In [413]:
##Decision Tree Classifier
from sklearn.tree import DecisionTreeClassifier
clf=DecisionTreeClassifier()
t0 = time()
clf=DecisionTreeClassifier()
clf.fit(features train, labels train)
pred=clf.predict(features test)
from sklearn.metrics import accuracy score, precision score, recall score
acc=accuracy_score(pred, labels test)
print "Accuracy when using Decision Tree Classifier: " + str(acc)
print "DT algorithm time:", round(time()-t0, 3), "s"
print "Precision: " +str(precision score(pred, labels test))
print "Recall: "+str(recall score(pred, labels test))
Accuracy when using Decision Tree Classifier: 0.636363636364
DT algorithm time: 0.007 s
```

Precision: 0.25 Recall: 0.166666666667

Now accuracy is not much therefore need of Feature selection to maximize the performance. As the number of features decrease to important features the

- Dataset is reduced.Less data with more information.
- Less features Predict the label more accurately.

## **Feature Selection**

feature importances:

The feature importances. The higher, the more important the feature. The importance of a feature is computed as the (normalized) total reduction of the criterion brought by that feature. It is also known as the Gini importance. Now feature importance is calculated using decision tree

```
In [414]:
```

```
dict={}
key=0
```

```
[0.30664643327686802, 0.19287949921752737, 0.18550724637681168, 0.14814814814814, 0.08333333333333335, 0.062652006313978229, 0.020833333333333329, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0, 0.0]

New Feature List: ['poi', 'from_messages', 'total_payments', 'salary', 'total_stock_value', 'exercised_stock_options']
```

Now using above features with high gini importance is chosen for getting optimal accuracy.

```
In [415]:
```

```
data = featureFormat(data dict, feature list)
labels, features = targetFeatureSplit(data)
from sklearn.cross validation import train test split
features train, features test, labels train, labels test=train test split(feat
ures,labels,test size=0.3,random state=42)
from sklearn.naive bayes import GaussianNB
from time import time
t0 = time()
clf = GaussianNB()
clf.fit(features train, labels train)
pred = clf.predict(features test)
accuracy = accuracy_score(pred,labels_test)
print "Accuracy when using Naive Bayes Classifier:"+str(accuracy)
print "Precision: " +str(precision score(pred, labels test))
print "Recall: "+str(recall score(pred, labels test))
print "NB algorithm time:", round(time()-t0, 3), "s"
```

When there is no splitting of dataset into Training and Testing set i.e. same data is used for classification and prediction can be seen below:

#### In [416]:

```
clf=DecisionTreeClassifier()
clf.fit(features, labels)
pred=clf.predict(features)

from sklearn.metrics import accuracy_score
acc=accuracy_score(pred_labels)
```

```
acc-accuracy score (pred, tabets)
print acc
```

1.0

Accuracy Score is increased to 1 but there are two problems when there is no splitting of dataset into Training and Testing data:

- Performance of the algorithm cannot be compared.
- Overfitting of Data . Therefore splitting the dataset into Training and testing dataset as shown below:

## In [417]:

```
t0 = time()
clf=DecisionTreeClassifier()
clf.fit(features train, labels train)
pred=clf.predict(features test)
from sklearn.metrics import accuracy score,precision score,recall score
acc=accuracy score(pred, labels test)
print "Accuracy when using Decision Tree Classifier: " + str(acc)
print "DT algorithm time:", round(time()-t0, 3), "s"
print "Precision: " +str(precision score(pred, labels test))
print "Recall: "+str(recall score(pred, labels test))
Accuracy when using Decision Tree Classifier: 0.744186046512
```

DT algorithm time: 0.002 s Precision: 0.2 Recall: 0.125

Now checking the accuracy while introducing new features in the Features list.

### In [418]:

```
feature list.append('fraction to poi email')
feature list.append('fraction from poi email')
print feature list
['poi', 'from messages', 'total payments', 'salary', 'total stock value', '
exercised stock options', 'fraction to poi email',
'fraction from poi email']
In [421]:
data = featureFormat(data dict, feature list)
```

```
labels, features = targetFeatureSplit(data)
from sklearn.cross validation import train test split
features train, features test, labels train, labels test=train test split(feat
ures,labels,test size=0.3,random state=42)
t0 = time()
clf=DecisionTreeClassifier()
clf.fit(features train, labels train)
pred=clf.predict(features test)
from sklearn.metrics import accuracy score, precision score, recall score
```

```
acc=accuracy_score(pred, labels_test)

print "Accuracy when using Decision Tree Classifier: " + str(acc)
print "DT algorithm time:", round(time()-t0, 3), "s"
print "Precision: " +str(precision_score(pred, labels_test))
print "Recall: "+str(recall_score(pred, labels_test))
```

```
Accuracy when using Decision Tree Classifier: 0.767441860465 DT algorithm time: 0.001 s Precision: 0.4 Recall: 0.22222222222
```

There is no significant change when adding new features. When trying new subset of features by manually selecting different set of features as shown in below found the accuracy, precision and Recall increase significantly.

## **Manual Selection of features**

Now manually selecting features having new features into consideration which increase the accuracy from 76% to 92%.

## In [432]:

```
features list = ["poi", "fraction from poi email", "fraction to poi email",
'shared receipt with poi']
data = featureFormat(data dict, features list)
labels, features = targetFeatureSplit(data)
### use KFold for split and validate algorithm
from sklearn.cross validation import KFold
kf=KFold(len(labels),3)
for train indices, test indices in kf:
    #make training and testing sets
    features train= [features[ii] for ii in train indices]
    features test= [features[ii] for ii in test indices]
    labels train=[labels[ii] for ii in train indices]
    labels test=[labels[ii] for ii in test indices]
from sklearn.cross validation import train test split
features_train, features_test, labels train, labels test=train test split(feat
ures, labels, test size=0.3, random state=42)
t0 = time()
clf=DecisionTreeClassifier()
clf.fit(features train, labels train)
pred=clf.predict(features test)
from sklearn.metrics import accuracy score, precision score, recall score
acc=accuracy score(pred, labels test)
print "Accuracy when using Decision Tree Classifier: " + str(acc)
print "DT algorithm time:", round(time()-t0, 3), "s"
print "Precision: " +str(precision score(pred, labels test))
print "Recall: "+str(recall score(pred, labels test))
t0 = time()
clf = GaussianNB()
clf.fit(features train, labels train)
pred = clf.predict(features test)
```

```
accuracy = accuracy_score(pred,labels_test)
print "Accuracy when using Naive Bayes Classifier:"+str(accuracy)
print "Precision: " +str(precision_score(pred,labels_test))
print "Recall: "+str(recall_score(pred,labels_test))
print "NB algorithm time:", round(time()-t0, 3), "s"

Accuracy when using Decision Tree Classifier: 0.923076923077
DT algorithm time: 0.002 s
Precision: 0.5
Recall: 1.0
Accuracy when using Naive Bayes Classifier:0.807692307692
Precision: 0.0
Recall: 0.0
NB algorithm time: 0.004 s
```

## Comparison of classifier

## In [428]:

```
%%html
Naive Bayes 
>Decision Tree Classifier
\langle tr \rangle
Precision
0
 0.5 
Recall
0
1
Accuracy
0.80
 0.92
```

	Naive Bayes	<b>Decision Tree Classifier</b>
Precision	0	0.5
Recall	0	1
Accuracy	0.80	0.92

# **Parameter Tuning**

Tuning is changing values of parameters present in the classifier to get optimal accuracy matrics and comparing them to get best classifier. In this dataset I cannot use accuracy for evaluating my algorithm because there a few POI's in dataset and the best evaluator are precision and recall. There were only 18 examples of POIs in the dataset. There were 35 people who were POIs in "real life", but

for various reasons, half of those are not present in this dataset. Therefore kfold is used with the classifier.

By manually setting the min\_samples\_split parameter in Decision Tree, Precision and Recall can be compared. Parameter min\_samples\_split are used to get best classifier.

```
In [401]:
```

```
def dt min samples split(k):
    t0 = time()
    clf=DecisionTreeClassifier(min samples split=k)
    clf.fit(features train, labels train)
    pred=clf.predict(features test)
    from sklearn.metrics import accuracy score, precision score, recall score
    acc=accuracy score(pred, labels test)
    print "Accuracy when using Decision Tree Classifier: " + str(acc)
    print "DT algorithm time:", round(time()-t0, 3), "s"
    print "Precision: " +str(precision score(pred, labels test))
    print "Recall: "+str(recall score(pred, labels test))
dt min samples split(2)
dt min samples split(3)
dt min samples split(5)
dt min samples split (10)
dt min samples split(15)
dt min samples split(20)
Accuracy when using Decision Tree Classifier: 0.884615384615
DT algorithm time: 0.001 s
Precision: 0.5
Recall: 0.66666666667
Accuracy when using Decision Tree Classifier: 0.923076923077
DT algorithm time: 0.002 s
Precision: 0.5
Recall: 1.0
Accuracy when using Decision Tree Classifier: 0.884615384615
DT algorithm time: 0.001 s
Precision: 0.5
Recall: 0.66666666667
Accuracy when using Decision Tree Classifier: 0.884615384615
DT algorithm time: 0.002 s
Precision: 0.5
Recall: 0.66666666667
Accuracy when using Decision Tree Classifier: 0.846153846154
DT algorithm time: 0.002 s
Precision: 0.75
Recall: 0.5
Accuracy when using Decision Tree Classifier: 0.769230769231
DT algorithm time: 0.001 s
Precision: 0.75
Recall: 0.375
```

## Validation Of Classifier

The validation of the algorithm performance is conducted use the tester function provided (without

changing). The function uses cross validation with 1000 folds. Precision: precision is defined as the number true positive divided by the number of person labels as positive. A higher precision value means a person flag out as a POI is more likely to be a true POI. Recall: recall is defined as the number of true positive divided by the total number of positive. A higher recall value mean if a person is a POI, the algorithm is more likely to flag this person out.

First I used accuracy to evaluate my algorithm. It was a mistake because in this case we have a class imbalance problem: the number of POIs is small compared to the total number of examples in the dataset. So I had to use precision and recall for these activities instead. I was able to reach average value of precision = 0.6, recall = 0.771.

## In [429]:

```
%%html
>Precision
Recall
min sample split=2
 0.5 
.66
min sample split=3
 0.5 
1
min sample split=5
.5
.66
min sample split=10
 0.5 
.6
min sample split=15
 0.75 
.5
min sample split=20
 0.75 
 .35
```

	Precision	Recall
min_sample_split=2	0.5	.66
min_sample_split=3	0.5	1
min_sample_split=5	.5	.66

min_sample_split=10	0.5	.6
min_sample_split=15	0.75	.5
min_sample_split=20	0.75	.35

There is usually a trade off between precision and recall. An algorithm strong at one metric may be weak at the other metric. So when it comes to decide which algorithm is better, it come to how do we define risk. Is it more risky to flag out as a POI who is actually not or is it more risky to miss a true POI? That is in the eyes of the beholder.

## **Conclusion**

Firstly I tried Naive Bayes accuracy was lower than with Decision Tree Algorithm (0.80 and 0.92 respectively). I made a conclusion that that the feature set I used does not suit the distributional and interactive assumptions of Naive Bayes well. I selected Decision Tree Algorithm for the POI identifier. It gave me accuracy before tuning parameters = 0.88. No feature scaling was used, as it's not necessary when using a decision tree. After selecting features and algorithm I manually tuned parameter min\_samples\_split. After using min\_samples\_split as 3 the Decision Tree gave maximum accuracy.