Summarize for us the goal of this project and how machine learning is useful in trying to accomplish it. As part of your answer, give some background on the dataset and how it can be used to answer the project question. Were there any outliers in the data when you got it, and how did you handle those? [relevant rubric items: "data exploration", "outlier investigation"]

Enron Corporation was U.S. energy-trading and utilities company founded in 1985. It achieved spectacular success in a very short period of time. This company claimed revenues of around 101 billion dollars in 2000. In only 15 years, this company was named America's Most innovative Company for six consecutive years by Fortune 500. Company stock price rose to 90 dollars. However, this success of Enron was short-lived. It started falling off more rapidly and bankruptcy was filed on Dec.2, 2001. Federal Energy Regulatory Commission investigated after the company's collapse and approximately 600,000 emails generated by employees of the Enron Corporation was made public which is called enron corpus.

In this project, main goal is to build a classification algorithm to predict a person of interest identifier (POI) based on email and financial features in the enron dataset involving 146 enron executives. A POI is anyone who is indicted, settled without admitting the guilt and testified in exchange for immunity. Predicted POI from this analysis and actual POI in the dataset will be compared to validate my prediction. Such model could be used to find additional suspects who were not indicted during the original investigation, or to find persons of interest during fraud investigations at other businesses. This study involves: -Exploration of Enron dataset -Selection of Features -Selection of Algorithms -Validation and Evaluation

```
import matplotlib.pyplot as plt
In [1]:
        from IPython.display import Image
        import numpy as np
        import pandas as pd
        import sys
        import pickle
        from sklearn.naive bayes import GaussianNB
        from sklearn.metrics import accuracy score
        from sklearn.metrics import precision score
        from sklearn.metrics import recall score
        from sklearn.metrics import recall score, precision score
        from time import time
        from sklearn.grid search import GridSearchCV
        sys.path.append("../tools/")
        from feature format import featureFormat, targetFeatureSplit
```

/anaconda/lib/python2.7/site-packages/sklearn/cross_validation.py:44 : DeprecationWarning: This module was deprecated in version 0.18 in favor of the model_selection module into which all the refactored cl asses and functions are moved. Also note that the interface of the n ew CV iterators are different from that of this module. This module will be removed in 0.20.

"This module will be removed in 0.20.", DeprecationWarning) /anaconda/lib/python2.7/site-packages/sklearn/grid_search.py:43: DeprecationWarning: This module was deprecated in version 0.18 in favor of the model_selection module into which all the refactored classes and functions are moved. This module will be removed in 0.20.

DeprecationWarning)

Exploration of Enron Dataset

Dataset was investigated. First structure of the dataset was examined.

In [4]: enron_data.head()

Out[4]:

	salary	to_messages	deferral_payments	total_payments	exercised_s
ALLEN PHILLIP K	201955	2902	2869717	4484442	1729541
BADUM JAMES P	NaN	NaN	178980	182466	257817
BANNANTINE JAMES M	477	566	NaN	916197	4046157
BAXTER JOHN C	267102	NaN	1295738	5634343	6680544
BAY FRANKLIN R	239671	NaN	260455	827696	NaN

5 rows × 21 columns

In [5]: #structure of enron_data set
print "There were", len(enron_data), "persons involved."

There were 146 persons involved.

In [6]: print data_dict.keys()

['METTS MARK', 'BAXTER JOHN C', 'ELLIOTT STEVEN', 'CORDES WILLIAM R' , 'HANNON KEVIN P', 'MORDAUNT KRISTINA M', 'MEYER ROCKFORD G', 'MCMA HON JEFFREY', 'HORTON STANLEY C', 'PIPER GREGORY F', 'HUMPHREY GENE E', 'UMANOFF ADAM S', 'BLACHMAN JEREMY M', 'SUNDE MARTIN', 'GIBBS DA NA R', 'LOWRY CHARLES P', 'COLWELL WESLEY', 'MULLER MARK S', 'JACKSO N CHARLENE R', 'WESTFAHL RICHARD K', 'WALTERS GARETH W', 'WALLS JR R OBERT H', 'KITCHEN LOUISE', 'CHAN RONNIE', 'BELFER ROBERT', 'SHANKMA N JEFFREY A', 'WODRASKA JOHN', 'BERGSIEKER RICHARD P', 'URQUHART JOH N A', 'BIBI PHILIPPE A', 'RIEKER PAULA H', 'WHALEY DAVID A', 'BECK S ALLY W', 'HAUG DAVID L', 'ECHOLS JOHN B', 'MENDELSOHN JOHN', 'HICKER SON GARY J', 'CLINE KENNETH W', 'LEWIS RICHARD', 'HAYES ROBERT E', ' MCCARTY DANNY J', 'KOPPER MICHAEL J', 'LEFF DANIEL P', 'LAVORATO JOH N J', 'BERBERIAN DAVID', 'DETMERING TIMOTHY J', 'WAKEHAM JOHN', 'POW ERS WILLIAM', 'GOLD JOSEPH', 'BANNANTINE JAMES M', 'DUNCAN JOHN H', 'SHAPIRO RICHARD S', 'SHERRIFF JOHN R', 'SHELBY REX', 'LEMAISTRE CHA RLES', 'DEFFNER JOSEPH M', 'KISHKILL JOSEPH G', 'WHALLEY LAWRENCE G' , 'MCCONNELL MICHAEL S', 'PIRO JIM', 'DELAINEY DAVID W', 'SULLIVAN-S HAKLOVITZ COLLEEN', 'WROBEL BRUCE', 'LINDHOLM TOD A', 'MEYER JEROME J', 'LAY KENNETH L', 'BUTTS ROBERT H', 'OLSON CINDY K', 'MCDONALD RE BECCA', 'CUMBERLAND MICHAEL S', 'GAHN ROBERT S', 'MCCLELLAN GEORGE', 'HERMANN ROBERT J', 'SCRIMSHAW MATTHEW', 'GATHMANN WILLIAM D', 'HAED ICKE MARK E', 'BOWEN JR RAYMOND M', 'GILLIS JOHN', 'FITZGERALD JAY L ', 'MORAN MICHAEL P', 'REDMOND BRIAN L', 'BAZELIDES PHILIP J', 'BELD EN TIMOTHY N', 'DURAN WILLIAM D', 'THORN TERENCE H', 'FASTOW ANDREW S', 'FOY JOE', 'CALGER CHRISTOPHER F', 'RICE KENNETH D', 'KAMINSKI W INCENTY J', 'LOCKHART EUGENE E', 'COX DAVID', 'OVERDYKE JR JERE C', 'PEREIRA PAULO V. FERRAZ', 'STABLER FRANK', 'SKILLING JEFFREY K', 'B LAKE JR. NORMAN P', 'SHERRICK JEFFREY B', 'PRENTICE JAMES', 'GRAY RO DNEY', 'PICKERING MARK R', 'THE TRAVEL AGENCY IN THE PARK', 'NOLES J AMES L', 'KEAN STEVEN J', 'TOTAL', 'FOWLER PEGGY', 'WASAFF GEORGE', 'WHITE JR THOMAS E', 'CHRISTODOULOU DIOMEDES', 'ALLEN PHILLIP K', 'S HARP VICTORIA T', 'JAEDICKE ROBERT', 'WINOKUR JR. HERBERT S', 'BROWN MICHAEL', 'BADUM JAMES P', 'HUGHES JAMES A', 'REYNOLDS LAWRENCE', 'D IMICHELE RICHARD G', 'BHATNAGAR SANJAY', 'CARTER REBECCA C', 'BUCHAN AN HAROLD G', 'YEAP SOON', 'MURRAY JULIA H', 'GARLAND C KEVIN', 'DOD SON KEITH', 'YEAGER F SCOTT', 'HIRKO JOSEPH', 'DIETRICH JANET R', 'D ERRICK JR. JAMES V', 'FREVERT MARK A', 'PAI LOU L', 'BAY FRANKLIN R' , 'HAYSLETT RODERICK J', 'FUGH JOHN L', 'FALLON JAMES B', 'KOENIG MA RK E', 'SAVAGE FRANK', 'IZZO LAWRENCE L', 'TILNEY ELIZABETH A', 'MAR TIN AMANDA K', 'BUY RICHARD B', 'GRAMM WENDY L', 'CAUSEY RICHARD A', 'TAYLOR MITCHELL S', 'DONAHUE JR JEFFREY M', 'GLISAN JR BEN F']

```
{'salary': 274975, 'to_messages': 873, 'deferral_payments': 'NaN', '
total_payments': 1272284, 'exercised_stock_options': 384728, 'bonus'
: 600000, 'restricted_stock': 393818, 'shared_receipt_with_poi': 874
, 'restricted_stock_deferred': 'NaN', 'total_stock_value': 778546, '
expenses': 125978, 'loan_advances': 'NaN', 'from_messages': 16, 'oth
er': 200308, 'from_this_person_to_poi': 6, 'poi': True, 'director_fe
es': 'NaN', 'deferred_income': 'NaN', 'long_term_incentive': 71023,
'email_address': 'ben.glisan@enron.com', 'from_poi_to_this_person':
52}
```

```
In [8]: (enron_data['poi'].value_counts()[True])
```

Out[8]: 18

```
In [9]: enron_data.describe().transpose()
```

In [7]: print data dict["GLISAN JR BEN F"]

Out[9]:

	count	unique	top	freq
salary	146	95	NaN	51
to_messages	146	87	NaN	60
deferral_payments	146	40	NaN	107
total_payments	146	126	NaN	21
exercised_stock_options	146	102	NaN	44
bonus	146	42	NaN	64
restricted_stock	146	98	NaN	36
shared_receipt_with_poi	146	84	NaN	60
restricted_stock_deferred	146	19	NaN	128
total_stock_value	146	125	NaN	20
expenses	146	95	NaN	51
loan_advances	146	5	NaN	142
from_messages	146	65	NaN	60
other	146	93	NaN	53
from_this_person_to_poi	146	42	NaN	60
poi	146	2	False	128
director_fees	146	18	NaN	129
deferred_income	146	45	NaN	97
long_term_incentive	146	53	NaN	80
email_address	146	112	NaN	35
from_poi_to_this_person	146	58	NaN	60

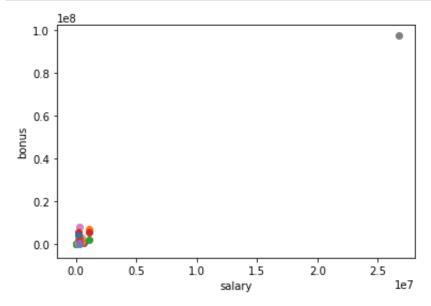
Enron dataset consists of lot of missing values (NaN). All the features have at least one missing values. Some features have more than 50% of their values missing, as shown by the table above. NaNs are replaced by 0 for training our algorithm later.

```
In [10]: enron_data.replace(to_replace='NaN', value=0.0, inplace=True)
```

Scatter plots are great tools for finding outliers. So first I plotted salary against bonus in a scatter plot and looked for the data distribution.

```
In [11]: from IPython.display import Image
    features = ["salary", "bonus"]
    #data_dict.pop('TOTAL', 0)
    data = featureFormat(data_dict, features)
### plot features
    for point in data:
        salary = point[0]
        bonus = point[1]
        plt.scatter( salary, bonus )

plt.xlabel("salary")
    plt.ylabel("bonus")
    plt.show()
```



The above plot allowed to see an outlier named 'Total'which was sum of all the data points in the given plot(artifact in scatterplot). So this outlier was excluded from our analysis manually. There were several other high number values (outliers) which could possibly represent POIs, thus were included in the dataset. Scatter plot was drawn after removing outliers as below.

```
In [12]: ##2 removing outliers
    features = ["salary", "bonus"]
    data_dict.pop('TOTAL', 0)
    data = featureFormat(data_dict, features)

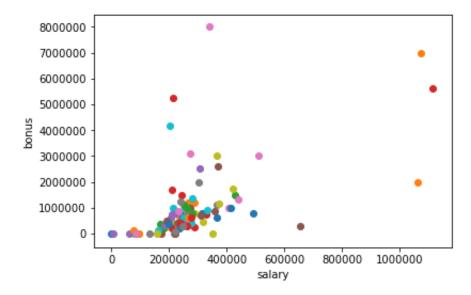
#remove NAN from dataset
    outliers = []
    for key in data_dict:
        val = data_dict[key]['salary']
        if val == 'NaN':
            continue
        outliers.append((key, int(val)))
    outliers_final = (sorted(outliers, key=lambda x:x[1], reverse = True)[
    :10])
    print outliers_final
```

[('SKILLING JEFFREY K', 1111258), ('LAY KENNETH L', 1072321), ('FREV ERT MARK A', 1060932), ('PICKERING MARK R', 655037), ('WHALLEY LAWRE NCE G', 510364), ('DERRICK JR. JAMES V', 492375), ('FASTOW ANDREW S', 440698), ('SHERRIFF JOHN R', 428780), ('RICE KENNETH D', 420636), ('CAUSEY RICHARD A', 415189)]

```
In [13]: #scatterplot after outlier removal

features = ["salary", "bonus"]
  #data_dict.pop('TOTAL', 0)
  data = featureFormat(data_dict, features)
  ### plot features
  for point in data:
      salary = point[0]
      bonus = point[1]
      plt.scatter( salary, bonus )

plt.xlabel("salary")
  plt.ylabel("bonus")
  plt.show()
```



Selection of Feature/Feature engineering

What features did you end up using in your POI identifier, and what selection process did you use to pick them? Did you have to do any scaling? Why or why not? As part of the assignment, you should attempt to engineer your own feature that does not come ready-made in the dataset -- explain what feature you tried to make, and the rationale behind it. (You do not necessarily have to use it in the final analysis, only engineer and test it.) In your feature selection step, if you used an algorithm like a decision tree, please also give the feature importances of the features that you use, and if you used an automated feature selection function like SelectKBest, please report the feature scores and reasons for your choice of parameter values. [relevant rubric items: "create new features", "intelligently select features", "properly scale features"]

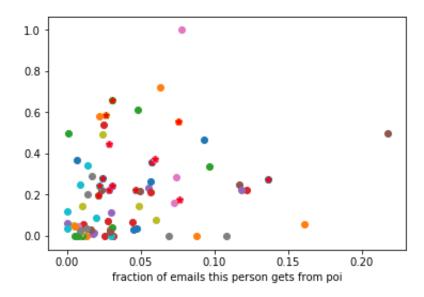
Feature selection in machine learning is selection of a feature which potentially would show some sort of pattern in the prediction analysis. Enron dataset consists two set of features - financial(salary, bonus, stock etc) and communication(to and from emails). Here I am more interested in emails those involve POIs. The assumption is that "Communication between POI and POI would be more frequent than between POI and non-POIs". I attempted to create two features, fractions of emails this person gets from POI(fraction_from_poi) and fraction of emails this person sends to POI(fraction_to_poi).

fraction from poi = number of emails this person gets from POI/total number of from messages

fraction to poi = number of emails this person sends to POI/total number sent messages

If email data is 'NaN', ratio is set to 0.

```
In [14]: #3 create new features
         #new features: fraction to poi = fraction of emails sent to POIs, frac
         tion from poi = fraction of emails received from POI
         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
         ][normalizer]))
             return new list
         ### create two lists of new features
         fraction from poi = dict to list("from poi to this person", "to message
         s")
         fraction_to_poi = dict_to_list("from_this_person_to_poi", "from_message
         s")
         ### insert new features into data dict
         count=0
         for i in data dict:
             data_dict[i]["fraction_from_poi"]=fraction_from_poi[count]
             data dict[i]["fraction to poi"]=fraction to poi[count]
             count +=1
         features list = ["poi", "fraction from poi", "fraction to poi"]
             ### store to my dataset for easy export below
         my dataset = data dict
         ### these two lines extract the features specified in features list
         ### and extract them from data dict, returning a numpy array
         data = featureFormat(my dataset, features list)
         ### plot new features
         for point in data:
             from poi = point[1]
             to poi = point[2]
             plt.scatter( from poi, to poi )
             if point[0] == 1:
                 plt.scatter(from_poi, to_poi, color="r", marker="*")
         plt.xlabel("fraction of emails this person gets from poi")
         plt.show()
```



```
In [15]: print features_list
    ['poi', 'fraction_from_poi', 'fraction_to_poi']
```

To select more impactful features for classification, 'feature_importances' attribute of "Decision Tree" was used. Features were ranked and selection process was half manual process. First all the possible features were included in features_list and then selection was done on the basis of feature ranking.

```
In [16]: features list = ["poi", "salary", "bonus", "fraction from poi", "fract
         ion_to_poi",
                          'deferral payments', 'total payments', 'loan advances
         ', 'restricted stock deferred',
                           'deferred income', 'total stock value', 'expenses', '
         exercised stock options',
                          'long term incentive', 'shared receipt with poi', 're
         stricted stock', 'director fees']
         data = featureFormat(my dataset, features list)
         ### split into labels and features (this line assumes that the first
         ### feature in the array is the label, which is why "poi" must always
         ### be first in features list
         labels, features = targetFeatureSplit(data)
         ### split data into training and testing datasets
         #deploying feature selection
         from sklearn import cross_validation
         from sklearn.cross validation import train test split
         features train, features test, labels train, labels test = cross valid
         ation.train test split(features, labels, test size=0.1, random state=4
         2)
         ##try Decision tree
         from sklearn.tree import DecisionTreeClassifier
         t0 = time()
         clf = DecisionTreeClassifier()
         clf.fit(features train, labels train)
         score = clf.score(features test, labels test)
         print'Decision Tree'
         print 'accuracy before tuning', score
         print "Decision tree algorithm time:", round(time()-t0, 3), "s"
         importances = clf.feature importances
         import numpy as np
         indices = np.argsort(importances)[::-1]
         print 'Feature Ranking: '
         for i in range(16):
             print "{} feature {} ({})".format(i+1,features list[i+1],importanc
         es[indices[i]])
```

```
Decision Tree
accuracy before tuning 0.666666666667
Decision tree algorithm time: 0.007 s
Feature Ranking:
1 feature salary (0.174317879326)
2 feature bonus (0.158290984378)
3 feature fraction from poi (0.14622972935)
4 feature fraction_to_poi (0.128198757764)
5 feature deferral payments (0.118337314859)
6 feature total payments (0.0955181169023)
7 feature loan advances (0.0879795396419)
8 feature restricted stock deferred (0.0534161490683)
9 feature deferred income (0.0377115287109)
10 feature total stock value (0.0)
11 feature expenses (0.0)
12 feature exercised stock options (0.0)
13 feature long term incentive (0.0)
14 feature shared receipt with poi (0.0)
15 feature restricted stock (0.0)
16 feature director fees (0.0)
```

We select and keep least number of features which hold maximum information and show pattern and trends in data. From here I selected 9 features: ["salary", "bonus", "fraction_from_poi", "fraction_to_poi", "deferral_payments", "total_payments", "loan_advances", "restricted_stock_deferred", "deferred_income"] Accuracy for this feature set was 0.73. Recall and precision scores were too low, so I manually chose features which gave recall and precision score higher than 0.3. My final feature selections were: ["fraction_from_poi", "fraction_to_poi", "shared_receipt_with_poi"]

Algorithm selection and tuning

```
features list = ["poi", "fraction from poi", "fraction to poi", "shared
In [17]:
         _receipt_with_poi"]
         ##try Naive Bayes
         t0 = time()
         clf = GaussianNB()
         clf.fit(features train, labels train)
         pred = clf.predict(features test)
         print "Naive Bayes recall score", (recall score(labels test,pred))
         print "Naive Bayes precision score", (precision_score(labels_test,pred
         ))
         #print accuracy #(clf.score(features test, labels test))
         accuracy = accuracy score(pred, labels test)
         print 'accuracy', accuracy
         print "NB algorithm time:", round(time()-t0, 3), 's'
         ###Adaboost
         from sklearn.ensemble import AdaBoostClassifier
         clf = AdaBoostClassifier(DecisionTreeClassifier(min samples split = 40
         ),
                                  algorithm="SAMME",
                                  n estimators=200)
         clf.fit(features_train,labels_train)
         pred = clf.predict(features test)
         acc = accuracy score(pred, labels test)
         print "ADABOOST:"
         print acc
         print "AB Recall Score" + str(recall score(labels test, pred))
         print "AB Precision Score" + str(precision score(labels test, pred))
         Naive Bayes recall score 0.5
         Naive Bayes precision score 0.181818181818
         accuracy 0.26666666667
         NB algorithm time: 0.01 s
         ADABOOST:
         0.8
         AB Recall Score0.25
         AB Precision Score1.0
```

What algorithm did you end up using? What other one(s) did you try? How did model performance differ between algorithms?

What does it mean to tune the parameters of an algorithm, and what can happen if you don't do this well? How did you tune the parameters of your particular algorithm? What parameters did you tune?

Naive Bayes (accuracy = 0.27), decision tree (accuracy = 0.67) and adaboost(accuracy= 0.73) algorithms were applied. Accuracy was lower with Naive Bayes, could probably due to distribution pattern of features in the dataset. Adaboost showed decent accuracy but had recall score of less than 0.3. Here, Decision tree would be more appropriate algorithm for POI identifier, since as it has accuracy of 0.67 before any tuning. Besides it is more efficient in finding irregular decision boundaries and does not need feature scaling. To optimize its performance, parameters like min_sample_split could be varied.

According to the enron dataset there were only 18 POIs. Since there were few POI subjects in the dataset, precision and recall score were considered better evaluater. I chose decision tree as a final algorithm. After final algorithm was chosen, algorithm parameter min_samples_split was tuned manually. We varied the min_samples_split from 2 to 7 and compared the recall and precision score. We found recall and precision score did not change between numbers 3 to 7.

```
In [18]: | ### use manual tuning parameter min samples split
         t0 = time()
         clf = DecisionTreeClassifier(min samples split=5)
         clf = clf.fit(features_train,labels_train)
         pred= clf.predict(features test)
         print("done in %0.3fs" % (time() - t0))
         acc=accuracy score(labels test, pred)
         print "Validating algorithm:"
         print "accuracy after tuning = ", acc
         # function for calculation ratio of true positives
         # out of all positives (true + false)
         print 'precision = ', precision score(labels test,pred)
         # function for calculation ratio of true positives
         # out of true positives and false negatives
         print 'recall = ', recall_score(labels_test,pred)
         ### dump your classifier, dataset and features list so
         ### anyone can run/check your results
         pickle.dump(clf, open("my classifier.pkl", "w") )
         pickle.dump(data_dict, open("my_dataset.pkl", "w") )
         pickle.dump(features_list, open("my_feature_list.pkl", "w") )
         done in 0.006s
         Validating algorithm:
         accuracy after tuning = 0.733333333333
         precision = 0.5
         recall = 0.5
In [19]: \#t0 = time()
         #clf = DecisionTreeClassifier(min samples split=2)
         #clf = clf.fit(features train, labels train)
         #pred= clf.predict(features test)
         #print("done in %0.3fs" % (time() - t0))
         #acc=accuracy score(labels_test, pred)
         #print "Validating algorithm:"
         #print "accuracy after tuning = ", acc
         #function for calculation ratio of true positives
         #out of all positives (true + false)
         #print 'precision = ', precision score(labels test,pred)
         #function for calculation ratio of true positives
         #out of true positives and false negatives
         #print 'recall = ', recall_score(labels_test,pred)
```

min_samples_split precision recall 2 0.4 0.67 3 0.5 0.67 4 0.5 0.67 5 0.67 0.67 6 0.67 0.67 7 0.67 0.67

Analysis Validation and Performance

What is validation, and what's a classic mistake you can make if you do it wrong? How did you validate your analysis?

Validation is the process where we determine the robustness of our predictive models. In our case, analysis was validated using K-fold validation. Such validation process enhances the likelihood that our algorithm will be reliable and robust. A classic mistake in validation process is called over-fitting, where the model is trained and it performs very well on the training dataset, but is actually worse on the cross-validation and test datasets.

```
In [20]: ### features list is a list of strings, each of which is a feature nam
         ### first feature must be "poi", as this will be singled out as the la
         features_list = ["poi", "fraction_from_poi", "fraction to poi", "share
         d receipt with poi"]
         ### store to my_dataset for easy export below
         my dataset = data dict
         ### these two lines extract the features specified in features list
         ### and extract them from data dict, returning a numpy array
         data = featureFormat(my dataset, features list)
         ### split into labels and features (this line assumes that the first
         ### feature in the array is the label, which is why "poi" must always
         ### be first in features list
         labels, features = targetFeatureSplit(data)
         ### machine learning goes here!
         ### please name your classifier clf for easy export below
         ### deploying feature selection
         from sklearn import cross validation
         features train, features test, labels train, labels test = cross valid
         ation.train test split(features, labels, test size=0.1, random state=4
         2)
```

```
### 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.tree import DecisionTreeClassifier
t0 = time()
clf = DecisionTreeClassifier()
clf.fit(features train, labels train)
score = clf.score(features test, labels test)
print 'accuracy before tuning ', score
print "Decision tree algorithm time:", round(time()-t0, 3), "s"
### use manual tuning parameter min samples split
t0 = time()
clf = DecisionTreeClassifier(min samples split=5)
clf = clf.fit(features train, labels train)
pred= clf.predict(features_test)
print("done in %0.3fs" % (time() - t0))
acc=accuracy score(labels test, pred)
print "Validating algorithm:"
print "accuracy after tuning = ", acc
# function for calculation ratio of true positives
# out of all positives (true + false)
print 'precision = ', precision score(labels test,pred)
# function for calculation ratio of true positives
# out of true positives and false negatives
print 'recall = ', recall score(labels test,pred)
### dump your classifier, dataset and features list so
### anyone can run/check your results
pickle.dump(clf, open("my classifier.pkl", "w") )
pickle.dump(data_dict, open("my_dataset.pkl", "w") )
pickle.dump(features list, open("my feature list.pkl", "w") )
```

accuracy before tuning 0.857142857143
Decision tree algorithm time: 0.003 s
done in 0.001s
Validating algorithm:
accuracy after tuning = 0.892857142857
precision = 0.5
recall = 0.666666666667

Discussion and Conclusions

As this dataset was small and imbalanced dataset, accuracy was not a good metric for evaluating the algorithm. So, we used different scoring metric; precision and recall score. Decision Tree classifier showed the precision score and recall score higher than 0.3. Comparatively, Adaboost and Naive Bayes showed low accuracy and low precision score.

Give at least 2 evaluation metrics and your average performance for each of them. Explain an interpretation of your metrics that says something humanunderstandable about your algorithm's performance.

I chose Decision Tree as a final algorithm to predict whether POI identified through this test is indeed POI. Feature scaling was not done as it is not required while using Decision Tree. The precision score is the probability that the person identified as POI is infact POI. In our case, precision = 0.67 means 67 percent of the time POI identified in the test would be the real POI whereas 33% of the time this test could flag a wrong person. These numbers could be increased by changing or exploring more into email information. Finally, Our prediction was validated by using K-fold validation.

Thus in this project, I used machine learning algorithms to identify POI in the Enron dataset. Naive Bayes, Adaboost and Decision Tree algorithms were compared. Decision Tree showed best precision and recall score among them showing 0.67 and 0.67 respectively. Small size of data set and even smaller number of POI made this analysis more challanging.

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