

Microsoft Malware Detection

November 26, 2019

1 Microsoft Malware detection

1. Business/Real-world Problem

1.1. What is Malware?

The term malware is a contraction of malicious software. Put simply, malware is any piece of software that was written with the intent of doing harm to data, devices or to people. Source: <https://www.avg.com/en/signal/what-is-malware>

1.2. Problem Statement

In the past few years, the malware industry has grown very rapidly that, the syndicates invest heavily in technologies to evade traditional protection, forcing the anti-malware groups/communities to build more robust softwares to detect and terminate these attacks. The major part of protecting a computer system from a malware attack is to identify whether a given piece of file/software is a malware.

1.3 Source/Useful Links

Microsoft has been very active in building anti-malware products over the years and it runs its anti-malware utilities over 150 million computers around the world. This generates tens of millions of daily data points to be analyzed as potential malware. In order to be effective in analyzing and classifying such large amounts of data, we need to be able to group them into groups and identify their respective families. This dataset provided by Microsoft contains about 9 classes of malware. ,

Source: <https://www.kaggle.com/c/malware-classification>

1.4. Real-world/Business objectives and constraints.

1. Minimize multi-class error.
2. Multi-class probability estimates.
3. Malware detection should not take hours and block the user's computer. It should finish in a few seconds or a minute.

2. Machine Learning Problem

2.1. Data

2.1.1. Data Overview

Source : <https://www.kaggle.com/c/malware-classification/data>

For every malware, we have two files

.asm file (read more: <https://www.reviversoft.com/file-extensions/asm>)

.bytes file (the raw data contains the hexadecimal representation of the file's binary content, without the PE header)

Total train dataset consist of 200GB data out of which 50Gb of data is .bytes files and 150GB of data is .asm files:

Lots of Data for a single-box/computer.

There are total 10,868 .bytes files and 10,868 asm files total 21,736 files

There are 9 types of malwares (9 classes) in our give data

Types of Malware:

Ramnit

Lollipop

Kelihos_ver3

Vundo

Simda

Tracur

Kelihos_ver1

Obfuscator.ACY

Gatak

2.1.2. Example Data Point

.asm file

.bytes file

2.2. Mapping the real-world problem to an ML problem

2.2.1. Type of Machine Learning Problem

There are nine different classes of malware that we need to classify a given a data point :

2.2.2. Performance Metric

Source: <https://www.kaggle.com/c/malware-classification#evaluation>

Metric(s): * Multi class log-loss * Confusion matrix

2.2.3. Machine Learning Objectives and Constraints

Objective: Predict the probability of each data-point belonging to each of the nine classes.

Constraints:

- Class probabilities are needed.
- Penalize the errors in class probabilities => Metric is Log-loss.
- Some Latency constraints.

2.3. Train and Test Dataset

Split the dataset randomly into three parts train, cross validation and test with 64%,16%, 20% of data respectively

2.4. Useful blogs, videos and reference papers

<http://blog.kaggle.com/2015/05/26/microsoft-malware-winners-interview-1st-place-no-to-overfitting/> <https://arxiv.org/pdf/1511.04317.pdf> First place solution in Kaggle competition: <https://www.youtube.com/watch?v=VLQTRILGz5Y>
<https://github.com/dchad/malware-detection> <http://vizsec.org/files/2011/Nataraj.pdf>
https://www.dropbox.com/sh/gfqzv0ckgs4l1bf/AAB6EelnEjvvuQg2nu_pIB6ua?dl=0 " Cross validation is more trustworthy than domain knowledge."

3. Exploratory Data Analysis

```
[1]: import warnings
warnings.filterwarnings("ignore")
import shutil
import os
import pandas as pd
import matplotlib
matplotlib.use(u'nbAgg')
import matplotlib.pyplot as plt
import seaborn as sns
import numpy as np
import pickle
from sklearn.manifold import TSNE
from sklearn import preprocessing
import pandas as pd
from multiprocessing import Process# this is used for multithreading
import multiprocessing
import codecs# this is used for file operations
import random as r
from xgboost import XGBClassifier
from sklearn.model_selection import RandomizedSearchCV
from sklearn.tree import DecisionTreeClassifier
from sklearn.calibration import CalibratedClassifierCV
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import log_loss
from sklearn.metrics import confusion_matrix
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression, SGDClassifier
from sklearn.ensemble import RandomForestClassifier
import nltk
from nltk.util import everygrams

[2]: #separating byte files and asm files

source = 'train'
destination_1 = 'byteFiles'
destination_2 = 'asmFiles'

# we will check if the folder 'byteFiles' exists if it not there we will create
→a folder with the same name
if not os.path.isdir(destination_1):
    os.makedirs(destination_1)
if not os.path.isdir(destination_2):
    os.makedirs(destination_2)

# if we have folder called 'train' (train folder contains both .asm files and .
→bytes files) we will rename it 'asmFiles'
```

```

# for every file that we have in our 'asmFiles' directory we check if it is
→ending with .bytes, if yes we will move it to
# 'byteFiles' folder

# so by the end of this snippet we will separate all the .byte files and .asm
→files
if os.path.isdir(source):
    data_files = os.listdir(source)
    for file in data_files:
        print(file)
        if (file.endswith("bytes")):
            shutil.move(source+'\\'+file,destination_1)
        if (file.endswith("asm")):
            shutil.move(source+'\\'+file,destination_2)

```

train

3.1. Distribution of malware classes in whole data set

```

[3]: Y=pd.read_csv("trainLabels.csv")
total = len(Y)*1.
ax=sns.countplot(x="Class", data=Y)
for p in ax.patches:
    ax.annotate('{:.1f}%'.format(100*p.get_height()/total), (p.get_x()+0.1,
→p.get_height()+5))

#put 11 ticks (therefore 10 steps), from 0 to the total number of rows in the
→dataframe
ax.yaxis.set_ticks(np.linspace(0, total, 11))

#adjust the ticklabel to the desired format, without changing the position of
→the ticks.
ax.set_yticklabels(map('{:.1f}%'.format, 100*ax.yaxis.get_majorticklocs()/
→total))
plt.show()

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

3.2. Feature extraction

3.2.1 File size of byte files as a feature

[4]: #file sizes of byte files

```

files=os.listdir('byteFiles')
filenames=Y['Id'].tolist()

```

```

class_y=Y['Class'].tolist()
class_bytes=[]
sizebytes=[]
fnames=[]
for file in files:
    # print(os.stat('byteFiles/0A32eTdBKayjCWhZqDOQ.txt'))
    # os.stat_result(st_mode=33206, st_ino=1125899906874507, st_dev=3561571700,
→st_nlink=1, st_uid=0, st_gid=0,
    # st_size=3680109, st_atime=1519638522, st_mtime=1519638522,
→st_ctime=1519638522)
    # read more about os.stat: here https://www.tutorialspoint.com/python/os\_stat.htm
→os_stat.htm
    statinfo=os.stat('byteFiles/'+file)
    # split the file name at '.' and take the first part of it i.e the file
→name
    file=file.split('.')[0]
    if any(file == filename for filename in filenames):
        i=filenames.index(file)
        class_bytes.append(class_y[i])
        # converting into Mb's
        sizebytes.append(statinfo.st_size/(1024.0*1024.0))
        fnames.append(file)
data_size_byte=pd.DataFrame({'ID':fnames,'size':sizebytes,'Class':class_bytes})
print (data_size_byte.head())

```

	ID	size	Class
0	01azqd4InC7m9JpocGv5	4.234863	9
1	01IsoiSMh5gxyDYTl4CB	5.538818	2
2	01jsnpXSA1gw6aPeDxrU	3.887939	9
3	01kcPWA9K2B0xQeS5Rju	0.574219	1
4	01SuzwMJEIXsK7A8dQbl	0.370850	8

3.2.2 box plots of file size (.byte files) feature

```

[5]: #boxplot of byte files
ax = sns.boxplot(x="Class", y="size", data=data_size_byte)
plt.title("boxplot of .bytes file sizes")
plt.show()

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

3.2.3 feature extraction from byte files

[6]:

```
# #removal of addres from byte files
# # contents of .byte files
# # -----
# #00401000 56 8D 44 24 08 50 8B F1 E8 1C 1B 00 00 C7 06 08
# #-----
# #we remove the starting address 00401000

# files = os.listdir('byteFiles')
# filenames=[]
# array=[]
# for file in files:
#     if(file.endswith("bytes")):
#         file=file.split('.')[0]
#         text_file = open('byteFiles/'+file+".txt", 'w+')
#         with open('byteFiles/'+file+".bytes", "r") as fp:
#             lines=""
#             for line in fp:
#                 a=line.rstrip().split(" ")[1:]
#                 b=' '.join(a)
#                 b=b+"\n"
#                 text_file.write(b)
#             fp.close()
#             os.remove('byteFiles/'+file+".bytes")
#         text_file.close()

# files = os.listdir('byteFiles')
# filenames2=[]
# feature_matrix = np.zeros((len(files),257),dtype=int)
# k=0

# #program to convert into bag of words of bytefiles
# #this is custom-built bag of words this is unigram bag of words
# byte_feature_file=open('result.csv', 'w+')
# byte_feature_file.
#     →write("ID,0,1,2,3,4,5,6,7,8,9,0a,0b,0c,0d,0e,0f,10,11,12,13,14,15,16,17,18,19,1a,1b,1c,1d,1
#     →?")
# byte_feature_file.write("\n")
# for file in files:
#     filenames2.append(file)
#     byte_feature_file.write(file+",")
#     if(file.endswith("txt")):
#         with open('byteFiles/'+file,"r") as byte_flie:
#             for lines in byte_flie:
#                 line=line.rstrip().split(" ")
#                 for hex_code in line:
#                     if hex_code=='??':
```

```

#             feature_matrix[k][256]+=1
#             else:
#                 feature_matrix[k][int(hex_code,16)]+=1
#             byte_file.close()
#         for i, row in enumerate(feature_matrix[k]):
#             if i!=len(feature_matrix[k])-1:
#                 byte_feature_file.write(str(row)+",")
#             else:
#                 byte_feature_file.write(str(row))
#         byte_feature_file.write("\n")

#     k += 1

# byte_feature_file.close()

```

[7]: #removal of addres from byte files
contents of .byte files

#00401000 56 8D 44 24 08 50 8B F1 E8 1C 1B 00 00 C7 06 08
#-----
#we remove the starting address 00401000

```

files = os.listdir('byteFiles')
filenames=[]
array=[]
for file in files:
    if(file.endswith("bytes")):
        file=file.split('.')[0]
        text_file = open('byteFiles/'+file+".txt", 'w+')
        with open('byteFiles/'+file+".bytes","r") as fp:
            lines=""
            for line in fp:
                a=line.rstrip().split(" ")[1:]
                b=' '.join(a)
                b=b+"\n"
                text_file.write(b)
            fp.close()
            os.remove('byteFiles/'+file+".bytes")
        text_file.close()

files = os.listdir('byteFiles')

```

[8]:

```

if not os.path.isfile('result_bigram.csv'):
    dataframe = pd.DataFrame()
    count=0
    bigrams_present =True
    if(not bigrams_present):
        for file in os.listdir('byteFiles'):

```

```

count+=1
with open("byteFiles/"+file,"r") as byte_flie:
    temp_dict = {}
    for lines in byte_flie:
        lines = lines.rstrip()
        tokenize = lines.split(" ")
        bigrams = everygrams(tokenize,1,2)
        bigram_fd = nltk.FreqDist(bigrams)
        for i,j in bigram_fd.items():
            key = str(i)[1:-1].replace("\'", "").replace(",","").
→replace(" ", "")

            if(key in temp_dict):
                temp_dict[key]+=j
            else:
                temp_dict[key]=j
        #print(temp_dict)
temp_fd = pd.DataFrame(temp_dict,index =[file.split('.')[0]])
#print(temp_fd)
dataframe = pd.concat([dataframe,temp_fd],axis=0)
if(len(dataframe)%1000 == 0):
    print("processed", count)
    dataframe.to_csv("file_"+str(count)+".csv")
    dataframe = pd.DataFrame()

```

```

[9]: if not os.path.isfile('result_bigram.csv'):
    all_bigram_features = pd.DataFrame()
    for f in os.listdir('./'):
        if 'file_' in f:
            temp = pd.read_csv(f)
            all_bigram_features = pd.concat([all_bigram_features,temp],axis=0)
    all_bigram_features = all_bigram_features.rename(columns= {'Unnamed: 0':
→'ID'})
    all_bigram_features.to_csv('result_bigram.csv')

```

```

[10]: byte_features=pd.read_csv("result_bigram.csv")
#byte_features['ID'] = byte_features['ID'].str.split('.').str[0]
byte_features.fillna(0, inplace=True)
byte_features.head(2)

```

```

[10]: Unnamed: 0      0      00      0000      0001      0002      0003      0004      0005  \
0          0  601905.0  273053.0  1002.0  801.0  1170.0  943.0  840.0
1          1  39755.0   19852.0   719.0   64.0   43.0  159.0   10.0

      0006      0007  ...  FFF7  FFF8  FFF9  FFFA  FFFB  FFFC  FFFD  FFFE  \
0  1125.0  1003.0  ...   10.0    9.0    7.0    5.0    7.0   11.0    9.0    6.0
1    6.0    10.0  ...   68.0   23.0   72.0   45.0   65.0   15.0   101.0  125.0

      FFFF
      ID

```



```
0 829.0 01azqd4InC7m9JpocGv5
1 4686.0 01IsoiSMh5gxyDYTl4CB
```

[2 rows x 66034 columns]

[11]: data_size_byte.head(2)

```
[11]:          ID      size  Class
0 01azqd4InC7m9JpocGv5  4.234863      9
1 01IsoiSMh5gxyDYTl4CB  5.538818      2
```

```
[12]: byte_features_with_size = byte_features.merge(data_size_byte, on='ID')
if not os.path.isfile('result_bigram.csv'):
    byte_features_with_size.to_csv("result_with_size.csv")
byte_features_with_size.head(2)
```

```
[12]: Unnamed: 0      00      0000      0001      0002      0003      0004      0005  \
0          0 601905.0 273053.0 1002.0 801.0 1170.0 943.0 840.0
1          1 39755.0 19852.0 719.0 64.0 43.0 159.0 10.0
```

```
      0006      0007  ...  FFF9  FFFA  FFFB  FFFC  FFFD  FFFE  FFFF  \
0 1125.0 1003.0  ...   7.0   5.0   7.0  11.0   9.0   6.0  829.0
1   6.0  10.0  ...  72.0  45.0  65.0  15.0 101.0 125.0 4686.0
```

```
          ID      size  Class
0 01azqd4InC7m9JpocGv5  4.234863      9
1 01IsoiSMh5gxyDYTl4CB  5.538818      2
```

[2 rows x 66036 columns]

```
[13]: # https://stackoverflow.com/a/29651514
def normalize(df):
    result1 = df.copy()
    for feature_name in df.columns:
        if (str(feature_name) != str('ID') and str(feature_name) != str('Class')):
            max_value = df[feature_name].max()
            min_value = df[feature_name].min()
            result1[feature_name] = (df[feature_name] - min_value) / (max_value -
↪ min_value)
    return result1
result = normalize(byte_features_with_size)
```

[14]: result.head(2)

```
[14]: Unnamed: 0      00      0000      0001      0002      0003      0004  \
0  0.000000  0.262806  0.127389  0.079943  0.054323  0.08898  0.064972
1  0.001001  0.017358  0.009262  0.057364  0.004340  0.00327  0.010955

      0005      0006      0007  ...  FFF9  FFFA  FFFB  FFFC  \
0  0.090303  0.109255  0.121901  ...  0.001933  0.003526  0.001031  0.001188
```

```
1 0.001075 0.000583 0.001215 ... 0.019884 0.031735 0.009574 0.001619
```

	FFFD	FFFE	FFFF	ID	size	Class
0	0.001294	0.000759	0.001227	01azqd4InC7m9JpocGv5	0.092219	9
1	0.014518	0.015811	0.006936	01IsoiSMh5gxyDYTl4CB	0.121236	2

[2 rows x 66036 columns]

```
[15]: data_y = result['Class']
result.head()
```

```
[15]: Unnamed: 0      00      0000      0001      0002      0003      0004 \
0      0.000000  0.262806  0.127389  0.079943  0.054323  0.088980  0.064972
1      0.001001  0.017358  0.009262  0.057364  0.004340  0.003270  0.010955
2      0.002002  0.040827  0.007479  0.047232  0.010648  0.010951  0.035070
3      0.003003  0.009209  0.004620  0.016276  0.004001  0.005248  0.007097
4      0.004004  0.008629  0.007132  0.004627  0.001356  0.008366  0.000551

      0005      0006      0007 ...      FFF9      FFFA      FFFB      FFFC \
0  0.090303  0.109255  0.121901 ...  0.001933  0.003526  0.001031  0.001188
1  0.001075  0.000583  0.001215 ...  0.019884  0.031735  0.009574  0.001619
2  0.063427  0.053511  0.017744 ...  0.022370  0.076164  0.017381  0.007125
3  0.003655  0.001845  0.002552 ...  0.002486  0.009168  0.002504  0.009284
4  0.001183  0.000291  0.000608 ...  0.006628  0.001410  0.000147  0.000000
```

	FFFD	FFFE	FFFF	ID	size	Class
0	0.001294	0.000759	0.001227	01azqd4InC7m9JpocGv5	0.092219	9
1	0.014518	0.015811	0.006936	01IsoiSMh5gxyDYTl4CB	0.121236	2
2	0.013943	0.010625	0.003393	01jsnpXSAlgW6aPeDxrU	0.084499	9
3	0.003450	0.007969	0.001954	01kcPWA9K2B0xQeS5Rju	0.010759	1
4	0.000575	0.000379	0.000108	01SuzwMJEIXsK7A8dQbl	0.006233	8

[5 rows x 66036 columns]

3.2.4 Multivariate Analysis

```
[16]: # #multivariate analysis on byte files
# #this is with perplexity 50
# xtsne=TSNE(perplexity=50)
# results=xtsne.fit_transform(result.drop(['ID','Class'], axis=1).sample(1000))
# vis_x = results[:, 0]
# vis_y = results[:, 1]
# plt.scatter(vis_x, vis_y, c=data_y, cmap=plt.cm.get_cmap("jet", 9))
# plt.colorbar(ticks=range(10))
# plt.clim(0.5, 9)
# plt.show()
```

```
[17]: # #this is with perplexity 30
# xtsne=TSNE(perplexity=30)
# results=xtsne.fit_transform(result.drop(['ID','Class'], axis=1).sample(1000))
```

```
# vis_x = results[:, 0]
# vis_y = results[:, 1]
# plt.scatter(vis_x, vis_y, c=data_y, cmap=plt.cm.get_cmap("jet", 9))
# plt.colorbar(ticks=range(10))
# plt.clim(0.5, 9)
# plt.show()
```

2 Train Test split

```
[18]: data_y = result['Class']
# split the data into test and train by maintaining same distribution of output
→variable 'y_true' [stratify=y_true]
X_train, X_test, y_train, y_test = train_test_split(result.drop(['ID', 'Class'],
→axis=1), data_y, stratify=data_y, test_size=0.20)
# split the train data into train and cross validation by maintaining same
→distribution of output variable 'y_train' [stratify=y_train]
X_train, X_cv, y_train, y_cv = train_test_split(X_train,
→y_train, stratify=y_train, test_size=0.20)

[19]: print('Number of data points in train data:', X_train.shape[0])
print('Number of data points in test data:', X_test.shape[0])
print('Number of data points in cross validation data:', X_cv.shape[0])
```

Number of data points in train data: 6955
Number of data points in test data: 2174
Number of data points in cross validation data: 1739

```
[20]: # it returns a dict, keys as class labels and values as the number of data
→points in that class
train_class_distribution = y_train.value_counts()
test_class_distribution = y_test.value_counts()
cv_class_distribution = y_cv.value_counts()

my_colors = list('rgbkymc')
train_class_distribution.plot(kind='bar', color=my_colors)
plt.xlabel('Class')
plt.ylabel('Data points per Class')
plt.title('Distribution of yi in train data')
plt.grid()
plt.show()

# ref: argsort https://docs.scipy.org/doc/numpy/reference/generated/numpy.
→argsort.html
# -(train_class_distribution.values): the minus sign will give us in decreasing
→order
sorted_yi = np.argsort(-train_class_distribution.values)
```

```

for i in sorted_yi:
    print('Number of data points in class', i+1, ':', train_class_distribution.
    ↪ values[i], '(', np.round((train_class_distribution.values[i]/y_train.
    ↪ shape[0]*100), 3), '%)')

print('-'*80)
my_colors = list('rbkymc')
test_class_distribution.plot(kind='bar', color=my_colors)
plt.xlabel('Class')
plt.ylabel('Data points per Class')
plt.title('Distribution of yi in test data')
plt.grid()
plt.show()

# ref: argsort https://docs.scipy.org/doc/numpy/reference/generated/numpy.
    ↪ argsort.html
# -(train_class_distribution.values): the minus sign will give us in decreasing
    ↪ order
sorted_yi = np.argsort(-test_class_distribution.values)
for i in sorted_yi:
    print('Number of data points in class', i+1, ':', test_class_distribution.
    ↪ values[i], '(', np.round((test_class_distribution.values[i]/y_test.
    ↪ shape[0]*100), 3), '%)')

print('-'*80)
my_colors = list('rbkymc')
cv_class_distribution.plot(kind='bar', color=my_colors)
plt.xlabel('Class')
plt.ylabel('Data points per Class')
plt.title('Distribution of yi in cross validation data')
plt.grid()
plt.show()

# ref: argsort https://docs.scipy.org/doc/numpy/reference/generated/numpy.
    ↪ argsort.html
# -(train_class_distribution.values): the minus sign will give us in decreasing
    ↪ order
sorted_yi = np.argsort(-train_class_distribution.values)
for i in sorted_yi:
    print('Number of data points in class', i+1, ':', cv_class_distribution.
    ↪ values[i], '(', np.round((cv_class_distribution.values[i]/y_cv.
    ↪ shape[0]*100), 3), '%)')

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Number of data points in class 1 : 1883 (27.074 %)
Number of data points in class 2 : 1586 (22.804 %)
Number of data points in class 3 : 986 (14.177 %)
Number of data points in class 4 : 786 (11.301 %)
Number of data points in class 5 : 648 (9.317 %)
Number of data points in class 6 : 481 (6.916 %)
Number of data points in class 7 : 304 (4.371 %)
Number of data points in class 8 : 254 (3.652 %)
Number of data points in class 9 : 27 (0.388 %)

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Number of data points in class 1 : 588 (27.047 %)
Number of data points in class 2 : 496 (22.815 %)
Number of data points in class 3 : 308 (14.167 %)
Number of data points in class 4 : 246 (11.316 %)
Number of data points in class 5 : 203 (9.338 %)
Number of data points in class 6 : 150 (6.9 %)
Number of data points in class 7 : 95 (4.37 %)
Number of data points in class 8 : 80 (3.68 %)
Number of data points in class 9 : 8 (0.368 %)

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Number of data points in class 1 : 471 (27.085 %)
Number of data points in class 2 : 396 (22.772 %)
Number of data points in class 3 : 247 (14.204 %)
Number of data points in class 4 : 196 (11.271 %)
Number of data points in class 5 : 162 (9.316 %)
Number of data points in class 6 : 120 (6.901 %)
Number of data points in class 7 : 76 (4.37 %)
Number of data points in class 8 : 64 (3.68 %)
Number of data points in class 9 : 7 (0.403 %)

```

[21]: def plot_confusion_matrix(test_y, predict_y):
    C = confusion_matrix(test_y, predict_y)
    print("Number of misclassified points ", (len(test_y)-np.trace(C))/
    → len(test_y)*100)
    # C = 9,9 matrix, each cell (i,j) represents number of points of class i
    → are predicted class j

    A = (((C.T)/(C.sum(axis=1))).T)
    #divid each element of the confusion matrix with the sum of elements in
    → that column

    # C = [[1, 2],
    #      [3, 4]]
    # C.T = [[1, 3],
    #        [2, 4]]
    # C.sum(axis = 1) axis=0 corresponds to columns and axis=1 corresponds to
    → rows in two dimensional array
    # C.sum(axis = 1) = [[3, 7]]
    # ((C.T)/(C.sum(axis=1))) = [[1/3, 3/7]
    #                             [2/3, 4/7]]

    # ((C.T)/(C.sum(axis=1))).T = [[1/3, 2/3]
    #                               [3/7, 4/7]]
    # sum of row elements = 1

    B = (C/C.sum(axis=0))
    #divid each element of the confusion matrix with the sum of elements in
    → that row
    # C = [[1, 2],
    #      [3, 4]]
    # C.sum(axis = 0) axis=0 corresponds to columns and axis=1 corresponds to
    → rows in two dimensional array
    # C.sum(axis = 0) = [[4, 6]]
    # (C/C.sum(axis=0)) = [[1/4, 2/6],
    #                       [3/4, 4/6]]

    labels = [1,2,3,4,5,6,7,8,9]
    cmap=sns.light_palette("green")
    # representing A in heatmap format
    print("-"*50, "Confusion matrix", "-"*50)
    plt.figure(figsize=(10,5))
    sns.heatmap(C, annot=True, cmap=cmap, fmt=".3f", xticklabels=labels,
    → yticklabels=labels)
    plt.xlabel('Predicted Class')
    plt.ylabel('Original Class')
    plt.show()

```

```

print("-"*50, "Precision matrix", "-"*50)
plt.figure(figsize=(10,5))
sns.heatmap(B, annot=True, cmap=cmap, fmt=".3f", xticklabels=labels,
→yticklabels=labels)
plt.xlabel('Predicted Class')
plt.ylabel('Original Class')
plt.show()
print("Sum of columns in precision matrix",B.sum(axis=0))

# representing B in heatmap format
print("-"*50, "Recall matrix", "-"*50)
plt.figure(figsize=(10,5))
sns.heatmap(A, annot=True, cmap=cmap, fmt=".3f", xticklabels=labels,
→yticklabels=labels)
plt.xlabel('Predicted Class')
plt.ylabel('Original Class')
plt.show()
print("Sum of rows in precision matrix",A.sum(axis=1))

```

4. Machine Learning Models

4.1. Machine Learning Models on bytes files

4.1.1. Random Model

[22]: *# we need to generate 9 numbers and the sum of numbers should be 1*
one solution is to generate 9 numbers and divide each of the numbers by their
→sum
ref: <https://stackoverflow.com/a/18662466/4084039>

```

test_data_len = X_test.shape[0]
cv_data_len = X_cv.shape[0]

# we create a output array that has exactly same size as the CV data
cv_predicted_y = np.zeros((cv_data_len,9))
for i in range(cv_data_len):
    rand_probs = np.random.rand(1,9)
    cv_predicted_y[i] = ((rand_probs/sum(sum(rand_probs))))[0])
print("Log loss on Cross Validation Data using Random
→Model",log_loss(y_cv,cv_predicted_y, eps=1e-15))

# Test-Set error.
#we create a output array that has exactly same as the test data
test_predicted_y = np.zeros((test_data_len,9))
for i in range(test_data_len):
    rand_probs = np.random.rand(1,9)
    test_predicted_y[i] = ((rand_probs/sum(sum(rand_probs))))[0])

```

```

print("Log loss on Test Data using Random_
→Model", log_loss(y_test, test_predicted_y, eps=1e-15))

predicted_y = np.argmax(test_predicted_y, axis=1)
plot_confusion_matrix(y_test, predicted_y+1)

```

Log loss on Cross Validation Data using Random Model 2.443796312798294

Log loss on Test Data using Random Model 2.4940427524996958

Number of misclassified points 89.0524379024839

----- Confusion matrix

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

----- Precision matrix

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Sum of columns in precision matrix [1. 1. 1. 1. 1. 1. 1. 1. 1.]

----- Recall matrix

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Sum of rows in precision matrix [1. 1. 1. 1. 1. 1. 1. 1. 1.]

4.1.2. Logistic Regression

[23]: *# read more about SGDClassifier() at http://scikit-learn.org/stable/modules/generated/sklearn.linear_model.SGDClassifier.html*
-----
default parameters
SGDClassifier(loss=hinge, penalty=l2, alpha=0.0001, l1_ratio=0.15,
→fit_intercept=True, max_iter=None, tol=None,
shuffle=True, verbose=0, epsilon=0.1, n_jobs=1, random_state=None,
→learning_rate=optimal, eta0=0.0, power_t=0.5,


```

# class_weight=None, warm_start=False, average=False, n_iter=None)

# some of methods
# fit(X, y[, coef_init, intercept_init, ])          Fit linear model with
→Stochastic Gradient Descent.
# predict(X)          Predict class labels for samples in X.

#-----
# video link: https://www.appliedaicourse.com/course/applied-ai-course-online/
→lessons/geometric-intuition-1/
#-----

alpha = [10 ** x for x in range(-5, 4)]
cv_log_error_array=[]
for i in alpha:
    print("for alpha :", i)
    logisticR=LogisticRegression(penalty='l2',C=i,class_weight='balanced')
    #logisticR = SGDClassifier(loss='log', penalty='l2', alpha=i,
→class_weight='balanced')
    logisticR.fit(X_train,y_train)
    sig_clf = CalibratedClassifierCV(logisticR, method="sigmoid")
    sig_clf.fit(X_train, y_train)
    predict_y = sig_clf.predict_proba(X_cv)
    cv_log_error_array.append(log_loss(y_cv, predict_y, labels=logisticR.
→classes_, eps=1e-15))

for i in range(len(cv_log_error_array)):
    print ('log_loss for c = ',alpha[i],'is',cv_log_error_array[i])

best_alpha = np.argmin(cv_log_error_array)

fig, ax = plt.subplots()
ax.plot(alpha, cv_log_error_array,c='g')
for i, txt in enumerate(np.round(cv_log_error_array,3)):
    ax.annotate((alpha[i],np.round(txt,3)), (alpha[i],cv_log_error_array[i]))
plt.grid()
plt.title("Cross Validation Error for each alpha")
plt.xlabel("Alpha i's")
plt.ylabel("Error measure")
plt.show()

```

```

for alpha : 1e-05
for alpha : 0.0001
for alpha : 0.001
for alpha : 0.01
for alpha : 0.1
for alpha : 1

```

```

for alpha : 10
for alpha : 100
for alpha : 1000
log_loss for c = 1e-05 is 1.5091639470652556
log_loss for c = 0.0001 is 1.1860813213733203
log_loss for c = 0.001 is 0.9049340046491288
log_loss for c = 0.01 is 0.4712626337114123
log_loss for c = 0.1 is 0.2602354198766801
log_loss for c = 1 is 0.1659372318509424
log_loss for c = 10 is 0.15520639357525837
log_loss for c = 100 is 0.17001171748331728
log_loss for c = 1000 is 0.17844179270952115

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

[24]: alpha[best_alpha]

[24]: 10

```

[25]: logisticR=LogisticRegression(penalty='l2',C=alpha[best_alpha],class_weight='balanced')
logisticR.fit(X_train,y_train)
sig_clf = CalibratedClassifierCV(logisticR, method="sigmoid")
sig_clf.fit(X_train, y_train)
pred_y=sig_clf.predict(X_test)

predict_y = sig_clf.predict_proba(X_train)
print ('log loss for train data',log_loss(y_train, predict_y, labels=logisticR.
    ↳classes_, eps=1e-15))
predict_y = sig_clf.predict_proba(X_cv)
print ('log loss for cv data',log_loss(y_cv, predict_y, labels=logisticR.
    ↳classes_, eps=1e-15))
predict_y = sig_clf.predict_proba(X_test)
print ('log loss for test data',log_loss(y_test, predict_y, labels=logisticR.
    ↳classes_, eps=1e-15))
plot_confusion_matrix(y_test, sig_clf.predict(X_test))

```

```

log loss for train data 0.06654660321053883
log loss for cv data 0.15520639357525837
log loss for test data 0.21466039403254375
Number of misclassified points 2.391904323827047

```

```

----- Confusion matrix
-----

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

----- Precision matrix

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Sum of columns in precision matrix [1. 1. 1. 1. 1. 1. 1. 1. 1.]
----- Recall matrix

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Sum of rows in precision matrix [1. 1. 1. 1. 1. 1. 1. 1. 1.]

4.2 Modeling with .asm files

4.2.1 Feature extraction from asm files

To extract the unigram features from the .asm files we need to process ~150GB of data

Note: Below two cells will take lot of time (over 48 hours to complete)

We will provide you the output file of these two cells, which you can directly use it

```
[26]: # #initially create five folders
# #first
# #second
# #thrid
# #fourth
# #fifth
# #this code tells us about random split of files into five folders
# folder_1='first'
# folder_2='second'
# folder_3='third'
# folder_4='fourth'
# folder_5='fifth'
# folder_6='output'
# for i in [folder_1,folder_2,folder_3,folder_4,folder_5,folder_6]:
#     if not os.path.isdir(i):
#         os.makedirs(i)

# source='train/'
# files = os.listdir('train')
# ID=df['Id'].tolist()
```

```

# data=range(0,10868)
# r.shuffle(data)
# count=0
# for i in range(0,10868):
#     if i % 5==0:
#         shutil.move(source+files[data[i]], 'first')
#     elif i%5==1:
#         shutil.move(source+files[data[i]], 'second')
#     elif i%5 ==2:
#         shutil.move(source+files[data[i]], 'thrid')
#     elif i%5 ==3:
#         shutil.move(source+files[data[i]], 'fourth')
#     elif i%5==4:
#         shutil.move(source+files[data[i]], 'fifth')

```

[27]: <http://flint.cs.yale.edu/cs421/papers/x86-asm/asm.html>

```

# def firstprocess():
#     #The prefixes tells about the segments that are present in the asm files
#     #There are 450 segments(approx) present in all asm files.
#     #this prefixes are best segments that gives us best values.
#     #https://en.wikipedia.org/wiki/Data_segment

#     prefixes = ['HEADER:', '.text:', '.Pav:', '.idata:', '.data:', '.bss:', '.rdata:',
→ '.edata:', '.rsrc:', '.tls:', '.reloc:', '.BSS:', '.CODE']
#     #this are opcodes that are used to get best results
#     #https://en.wikipedia.org/wiki/X86_instruction_listings

#     opcodes = ['jmp', 'mov', 'retf', 'push', 'pop', 'xor', 'retn', 'nop',
→ 'sub', 'inc', 'dec', 'add', 'imul', 'xchg', 'or', 'shr', 'cmp', 'call',
→ 'shl', 'ror', 'rol', 'jnb', 'jz', 'rtn', 'lea', 'movzx']
#     #best keywords that are taken from different blogs
#     keywords = ['.dll', 'std:', ':dword']
#     #Below taken registers are general purpose registers and special
→ registers
#     #All the registers which are taken are best
#     registers=['edx', 'esi', 'eax', 'ebx', 'ecx', 'edi', 'ebp', 'esp', 'eip']
#     file1=open("output\asm\smallfile.txt", "w+")
#     files = os.listdir('first')
#     for f in files:
#         #filling the values with zeros into the arrays
#         prefixescount=np.zeros(len(prefixes), dtype=int)
#         opcodecount=np.zeros(len(opcodes), dtype=int)
#         keywordcount=np.zeros(len(keywords), dtype=int)
#         registerscount=np.zeros(len(registers), dtype=int)
#         features=[]
#         f2=f.split('.')[0]

```

```

#         file1.write(f2+",")
#         opcodefile.write(f2+" ")
#         # https://docs.python.org/3/library/codecs.html#codecs.ignore_errors
#         # https://docs.python.org/3/library/codecs.html#codecs.Codec.encode
#         with codecs.open('first/'+f,encoding='cp1252',errors='replace') as f:
#             fli:
#                 for lines in fli:
#                     # https://www.tutorialspoint.com/python3/string_rstrip.htm
#                     line=lines.rstrip().split()
#                     l=line[0]
#                     #counting the prefixes in each and every line
#                     for i in range(len(prefixes)):
#                         if prefixes[i] in line[0]:
#                             prefixescount[i]+=1
#                     line=line[1:]
#                     #counting the opcodes in each and every line
#                     for i in range(len(opcodes)):
#                         if any(opcodes[i]==li for li in line):
#                             features.append(opcodes[i])
#                             opcodescount[i]+=1
#                     #counting registers in the line
#                     for i in range(len(registers)):
#                         for li in line:
#                             # we will use registers only in 'text' and 'CODE'
#             segments
#                 if registers[i] in li and ('text' in l or 'CODE' in
#             l):
#                 registerscount[i]+=1
#                 #counting keywords in the line
#                 for i in range(len(keywords)):
#                     for li in line:
#                         if keywords[i] in li:
#                             keywordcount[i]+=1
#                 #pushing the values into the file after reading whole file
#                 for prefix in prefixescount:
#                     file1.write(str(prefix)+",")
#                 for opcode in opcodescount:
#                     file1.write(str(opcode)+",")
#                 for register in registerscount:
#                     file1.write(str(register)+",")
#                 for key in keywordcount:
#                     file1.write(str(key)+",")
#                 file1.write("\n")
#             file1.close()

# #same as above

```

```

# def secondprocess():
#     prefixes = ['HEADER:', '.text:', '.Pav:', '.idata:', '.data:', '.bss:', '.rdata:',
→ '.edata:', '.rsrc:', '.tls:', '.reloc:', '.BSS:', '.CODE']
#     opcodes = ['jmp', 'mov', 'retf', 'push', 'pop', 'xor', 'retn', 'nop',
→ 'sub', 'inc', 'dec', 'add', 'imul', 'xchg', 'or', 'shr', 'cmp', 'call',
→ 'shl', 'ror', 'rol', 'jnb', 'jz', 'rtn', 'lea', 'movzx']
#     keywords = ['.dll', 'std:', ':dword']
#     registers=['edx', 'esi', 'eax', 'ebx', 'ecx', 'edi', 'ebp', 'esp', 'eip']
#     file1=open("output\mediumasmfile.txt", "w+")
#     files = os.listdir('second')
#     for f in files:
#         prefixescount=np.zeros(len(prefixes), dtype=int)
#         opcodescount=np.zeros(len(opcodes), dtype=int)
#         keywordcount=np.zeros(len(keywords), dtype=int)
#         registerscount=np.zeros(len(registers), dtype=int)
#         features=[]
#         f2=f.split('.')[0]
#         file1.write(f2+",")
#         opcodefile.write(f2+" ")
#         with codecs.open('second/'+f, encoding='cp1252', errors='replace') as
→ fli:
#             for lines in fli:
#                 line=lines.rstrip().split()
#                 l=line[0]
#                 for i in range(len(prefixes)):
#                     if prefixes[i] in line[0]:
#                         prefixescount[i]+=1
#                 line=line[1:]
#                 for i in range(len(opcodes)):
#                     if any(opcodes[i]==li for li in line):
#                         features.append(opcodes[i])
#                         opcodescount[i]+=1
#                 for i in range(len(registers)):
#                     for li in line:
#                         if registers[i] in li and ('text' in l or 'CODE' in
→ l):
#                             registerscount[i]+=1
#                 for i in range(len(keywords)):
#                     for li in line:
#                         if keywords[i] in li:
#                             keywordcount[i]+=1
#                 for prefix in prefixescount:
#                     file1.write(str(prefix)+",")
#                 for opcode in opcodescount:
#                     file1.write(str(opcode)+",")
#                 for register in registerscount:
#                     file1.write(str(register)+",")

```

```

#         for key in keywordcount:
#             file1.write(str(key)+",")
#         file1.write("\n")
#     file1.close()

# # same as smallprocess() functions
# def thirdprocess():
#     prefixes = ['HEADER:', '.text:', '.Pav:', '.idata:', '.data:', '.bss:', '.rdata:',
# → '.edata:', '.rsrc:', '.tls:', '.reloc:', '.BSS:', '.CODE']
#     opcodes = ['jmp', 'mov', 'retf', 'push', 'pop', 'xor', 'retn', 'nop',
→ 'sub', 'inc', 'dec', 'add', 'imul', 'xchg', 'or', 'shr', 'cmp', 'call',
→ 'shl', 'ror', 'rol', 'jnb', 'jz', 'rtn', 'lea', 'movzx']
#     keywords = ['.dll', 'std:', ':dword']
#     registers=['edx', 'esi', 'eax', 'ebx', 'ecx', 'edi', 'ebp', 'esp', 'eip']
#     file1=open("output\largeasmfile.txt", "w+")
#     files = os.listdir('thrid')
#     for f in files:
#         prefixescount=np.zeros(len(prefixes), dtype=int)
#         opcodescount=np.zeros(len(opcodes), dtype=int)
#         keywordcount=np.zeros(len(keywords), dtype=int)
#         registerscount=np.zeros(len(registers), dtype=int)
#         features=[]
#         f2=f.split('.')[0]
#         file1.write(f2+",")
#         opcodefile.write(f2+" ")
#         with codecs.open('thrid/'+f, encoding='cp1252', errors='replace') as
→ fli:
#             for lines in fli:
#                 line=line.rstrip().split()
#                 l=line[0]
#                 for i in range(len(prefixes)):
#                     if prefixes[i] in line[0]:
#                         prefixescount[i]+=1
#                 line=line[1:]
#                 for i in range(len(opcodes)):
#                     if any(opcodes[i]==li for li in line):
#                         features.append(opcodes[i])
#                         opcodescount[i]+=1
#                 for i in range(len(registers)):
#                     for li in line:
#                         if registers[i] in li and ('text' in l or 'CODE' in
→ l):
#                             registerscount[i]+=1
#                 for i in range(len(keywords)):
#                     for li in line:
#                         if keywords[i] in li:
#                             keywordcount[i]+=1

```

```

#         for prefix in prefixescount:
#             file1.write(str(prefix)+",")
#         for opcode in opcodescount:
#             file1.write(str(opcode)+",")
#         for register in registerscount:
#             file1.write(str(register)+",")
#         for key in keywordcount:
#             file1.write(str(key)+",")
#         file1.write("\n")
#     file1.close()

# def fourthprocess():
#     prefixes = ['HEADER:', '.text:', '.Pau:', '.idata:', '.data:', '.bss:', '.rdata:',
→ '.edata:', '.rsrc:', '.tls:', '.reloc:', '.BSS:', '.CODE']
#     opcodes = ['jmp', 'mov', 'retf', 'push', 'pop', 'xor', 'retn', 'nop',
→ 'sub', 'inc', 'dec', 'add', 'imul', 'xchg', 'or', 'shr', 'cmp', 'call',
→ 'shl', 'ror', 'rol', 'jnb', 'jz', 'rtn', 'lea', 'movzx']
#     keywords = ['.dll', 'std:', ':dword']
#     registers=['edx', 'esi', 'eax', 'ebx', 'ecx', 'edi', 'ebp', 'esp', 'eip']
#     file1=open("output\hugeasmfile.txt", "w+")
#     files = os.listdir('fourth/')
#     for f in files:
#         prefixescount=np.zeros(len(prefixes), dtype=int)
#         opcodescount=np.zeros(len(opcodes), dtype=int)
#         keywordcount=np.zeros(len(keywords), dtype=int)
#         registerscount=np.zeros(len(registers), dtype=int)
#         features=[]
#         f2=f.split('.')[0]
#         file1.write(f2+",")
#         opcodefile.write(f2+" ")
#         with codecs.open('fourth/'+f, encoding='cp1252', errors = 'replace') as
→ fli:
#             for lines in fli:
#                 line=lines.rstrip().split()
#                 l=line[0]
#                 for i in range(len(prefixes)):
#                     if prefixes[i] in line[0]:
#                         prefixescount[i]+=1
#                 line=line[1:]
#                 for i in range(len(opcodes)):
#                     if any(opcodes[i]==li for li in line):
#                         features.append(opcodes[i])
#                         opcodescount[i]+=1
#                 for i in range(len(registers)):
#                     for li in line:

```



```

#                                     if registers[i] in li and ('text' in l or 'CODE' in
→l):
#                                     registerscount[i]+=1
#                                     for i in range(len(keywords)):
#                                     for li in line:
#                                     if keywords[i] in li:
#                                     keywordcount[i]+=1
#                                     for prefix in prefixescount:
#                                     file1.write(str(prefix)+",")
#                                     for opcode in opcodescount:
#                                     file1.write(str(opcode)+",")
#                                     for register in registerscount:
#                                     file1.write(str(register)+",")
#                                     for key in keywordcount:
#                                     file1.write(str(key)+",")
#                                     file1.write("\n")
#                                     file1.close()

# def fifthprocess():
#     prefixes = ['HEADER:', '.text:', '.Pau:', '.idata:', '.data:', '.bss:', '.rdata:
→', '.edata:', '.rsrc:', '.tls:', '.reloc:', '.BSS:', '.CODE']
#     opcodes = ['jmp', 'mov', 'retf', 'push', 'pop', 'xor', 'retn', 'nop',
→'sub', 'inc', 'dec', 'add', 'imul', 'xchg', 'or', 'shr', 'cmp', 'call',
→'shl', 'ror', 'rol', 'jnb', 'jz', 'rtn', 'lea', 'movzx']
#     keywords = ['.dll', 'std:', ':dword']
#     registers=['edx', 'esi', 'eax', 'ebx', 'ecx', 'edi', 'ebp', 'esp', 'eip']
#     file1=open("output\trainasmfile.txt", "w+")
#     files = os.listdir('fifth/')
#     for f in files:
#         prefixescount=np.zeros(len(prefixes), dtype=int)
#         opcodescount=np.zeros(len(opcodes), dtype=int)
#         keywordcount=np.zeros(len(keywords), dtype=int)
#         registerscount=np.zeros(len(registers), dtype=int)
#         features=[]
#         f2=f.split('.')[0]
#         file1.write(f2+",")
#         opcodefile.write(f2+" ")
#         with codecs.open('fifth/'+f, encoding='cp1252', errors = 'replace') as
→fli:
#             for lines in fli:
#                 line=lines.rstrip().split()
#                 l=line[0]
#                 for i in range(len(prefixes)):
#                     if prefixes[i] in line[0]:
#                         prefixescount[i]+=1
#                 line=line[1:]

```

```

#         for i in range(len(opcodes)):
#             if any(opcodes[i]==li for li in line):
#                 features.append(opcodes[i])
#                 opcodescount[i]+=1
#         for i in range(len(registers)):
#             for li in line:
#                 if registers[i] in li and ('text' in l or 'CODE' in
→l):
#                     registerscount[i]+=1
#         for i in range(len(keywords)):
#             for li in line:
#                 if keywords[i] in li:
#                     keywordcount[i]+=1
#         for prefix in prefixescount:
#             file1.write(str(prefix)+",")
#         for opcode in opcodescount:
#             file1.write(str(opcode)+",")
#         for register in registerscount:
#             file1.write(str(register)+",")
#         for key in keywordcount:
#             file1.write(str(key)+",")
#         file1.write("\n")
#     file1.close()

# def main():
#     #the below code is used for multiprocessing
#     #the number of process depends upon the number of cores present System
#     #process is used to call multiprocessing
#     manager=multiprocessing.Manager()
#     p1=Process(target=firstprocess)
#     p2=Process(target=secondprocess)
#     p3=Process(target=thirdprocess)
#     p4=Process(target=fourthprocess)
#     p5=Process(target=fifthprocess)
#     #p1.start() is used to start the thread execution
#     p1.start()
#     p2.start()
#     p3.start()
#     p4.start()
#     p5.start()
#     #After completion all the threads are joined
#     p1.join()
#     p2.join()
#     p3.join()
#     p4.join()
#     p5.join()

```

```
# if __name__=="__main__":
#     main()
```

[28]: # asmoutputfile.csv(output generated from the above two cells) will contain all
 ↳ the extracted features from .asm files
 # this file will be uploaded in the drive, you can directly use this
 dfasm=pd.read_csv("asmoutputfile.csv")
 Y.columns = ['ID', 'Class']
 result_asm = pd.merge(dfasm, Y,on='ID', how='left')
 result_asm.head()

[28]:

	ID	HEADER:	.text:	.Pav:	.idata:	.data:	.bss:	\
0	01kcPWA9K2B0xQeS5Rju	19	744	0	127	57	0	
1	1E93CpP60RHFNiT5Qfvn	17	838	0	103	49	0	
2	3ekVow2ajZHbTnBcsDfX	17	427	0	50	43	0	
3	3X2nY7iQaPBIWDrAZqJe	17	227	0	43	19	0	
4	460ZzdsSKDCFV8h7XWxf	17	402	0	59	170	0	

	.rdata:	.edata:	.rsrc:	...	edx	esi	eax	ebx	ecx	edi	ebp	esp	eip	\
0	323	0	3	...	18	66	15	43	83	0	17	48	29	
1	0	0	3	...	18	29	48	82	12	0	14	0	20	
2	145	0	3	...	13	42	10	67	14	0	11	0	9	
3	0	0	3	...	6	8	14	7	2	0	8	0	6	
4	0	0	3	...	12	9	18	29	5	0	11	0	11	

	Class
0	1
1	1
2	1
3	1
4	1

[5 rows x 53 columns]

4.2.1.1 Files sizes of each .asm file

[29]: #file sizes of byte files

```
files=os.listdir('asmFiles')
filenames=Y['ID'].tolist()
class_y=Y['Class'].tolist()
class_bytes=[]
sizebytes=[]
fnames=[]
for file in files:
    # print(os.stat('byteFiles/OA32eTdBKayjCWWhZqDOQ.txt'))
    # os.stat_result(st_mode=33206, st_ino=1125899906874507, st_dev=3561571700,
    ↳st_nlink=1, st_uid=0, st_gid=0,
```

```

# st_size=3680109, st_atime=1519638522, st_mtime=1519638522,
→st_ctime=1519638522)
# read more about os.stat: here https://www.tutorialspoint.com/python/→os\_stat.htm
statinfo=os.stat('asmFiles/'+file)
# split the file name at '.' and take the first part of it i.e the file
→name
file=file.split('.')[0]
if any(file == filename for filename in filenames):
    i=filenames.index(file)
    class_bytes.append(class_y[i])
    # converting into Mb's
    sizebytes.append(statinfo.st_size/(1024.0*1024.0))
    fnames.append(file)
asm_size_byte=pd.DataFrame({'ID':fnames,'size':sizebytes,'Class':class_bytes})
print (asm_size_byte.head())

```

	ID	size	Class
0	01azqd4InC7m9JpocGv5	56.229886	9
1	01IsoiSMh5gxyDYTl4CB	13.999378	2
2	01jsnpXSA1gw6aPeDxrU	8.507785	9
3	01kcPWA9K2B0xQeS5Rju	0.078190	1
4	01SuzwMJEIXsK7A8dQbl	0.996723	8

4.2.1.2 Distribution of .asm file sizes

```

[30]: #boxplot of asm files
ax = sns.boxplot(x="Class", y="size", data=asm_size_byte)
plt.title("boxplot of .bytes file sizes")
plt.show()

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```

[31]: # add the file size feature to previous extracted features
print(result_asm.shape)
print(asm_size_byte.shape)
result_asm = pd.merge(result_asm, asm_size_byte.drop(['Class'],
→axis=1),on='ID', how='left')
result_asm.head()

```

(10868, 53)

(10868, 3)

```
[31]:
```

	ID	HEADER:	.text:	.Pav:	.idata:	.data:	.bss:	\
0	01kcPWA9K2B0xQeS5Rju	19	744	0	127	57	0	
1	1E93CpP60RHFNiT5Qfvn	17	838	0	103	49	0	
2	3ekVow2ajZHbTnBcsDfX	17	427	0	50	43	0	
3	3X2nY7iQaPBIWDrAZqJe	17	227	0	43	19	0	
4	460ZzdsSKDCFV8h7XWxf	17	402	0	59	170	0	

	.rdata:	.edata:	.rsrc:	...	esi	eax	ebx	ecx	edi	ebp	esp	eip	\
0	323	0	3	...	66	15	43	83	0	17	48	29	
1	0	0	3	...	29	48	82	12	0	14	0	20	
2	145	0	3	...	42	10	67	14	0	11	0	9	
3	0	0	3	...	8	14	7	2	0	8	0	6	
4	0	0	3	...	9	18	29	5	0	11	0	11	

	Class	size
0	1	0.078190
1	1	0.063400
2	1	0.041695
3	1	0.018757
4	1	0.037567

[5 rows x 54 columns]

```
[32]: # we normalize the data each column
result_asm = normalize(result_asm)
result_asm.head()
```

```
[32]:
```

	ID	HEADER:	.text:	.Pav:	.idata:	.data:	.bss:	\
0	01kcPWA9K2B0xQeS5Rju	0.107345	0.001092	0.0	0.000761	0.000023	0.0	
1	1E93CpP60RHFNiT5Qfvn	0.096045	0.001230	0.0	0.000617	0.000019	0.0	
2	3ekVow2ajZHbTnBcsDfX	0.096045	0.000627	0.0	0.000300	0.000017	0.0	
3	3X2nY7iQaPBIWDrAZqJe	0.096045	0.000333	0.0	0.000258	0.000008	0.0	
4	460ZzdsSKDCFV8h7XWxf	0.096045	0.000590	0.0	0.000353	0.000068	0.0	

	.rdata:	.edata:	.rsrc:	...	esi	eax	ebx	ecx	\
0	0.000084	0.0	0.000072	...	0.000746	0.000301	0.000360	0.001057	
1	0.000000	0.0	0.000072	...	0.000328	0.000965	0.000686	0.000153	
2	0.000038	0.0	0.000072	...	0.000475	0.000201	0.000560	0.000178	
3	0.000000	0.0	0.000072	...	0.000090	0.000281	0.000059	0.000025	
4	0.000000	0.0	0.000072	...	0.000102	0.000362	0.000243	0.000064	

	edi	ebp	esp	eip	Class	size
0	0.0	0.030797	0.001468	0.003173	1	0.000432
1	0.0	0.025362	0.000000	0.002188	1	0.000327
2	0.0	0.019928	0.000000	0.000985	1	0.000172
3	0.0	0.014493	0.000000	0.000657	1	0.000009
4	0.0	0.019928	0.000000	0.001204	1	0.000143

[5 rows x 54 columns]

4.2.2 Univariate analysis on asm file features

```
[33]: ax = sns.boxplot(x="Class", y=".text:", data=result_asm)
plt.title("boxplot of .asm text segment")
plt.show()
```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```
[34]: ax = sns.boxplot(x="Class", y=".Pav:", data=result_asm)
plt.title("boxplot of .asm pav segment")
plt.show()
```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```
[35]: ax = sns.boxplot(x="Class", y=".data:", data=result_asm)
plt.title("boxplot of .asm data segment")
plt.show()
```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```
[36]: ax = sns.boxplot(x="Class", y=".bss:", data=result_asm)
plt.title("boxplot of .asm bss segment")
plt.show()
```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```
[37]: ax = sns.boxplot(x="Class", y=".rdata:", data=result_asm)
plt.title("boxplot of .asm rdata segment")
plt.show()
```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```
[38]: ax = sns.boxplot(x="Class", y="jmp", data=result_asm)
plt.title("boxplot of .asm jmp opcode")
plt.show()
```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```
[39]: ax = sns.boxplot(x="Class", y="mov", data=result_asm)
plt.title("boxplot of .asm mov opcode")
plt.show()
```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```
[40]: ax = sns.boxplot(x="Class", y="retf", data=result_asm)
plt.title("boxplot of .asm retf opcode")
plt.show()
```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```
[41]: ax = sns.boxplot(x="Class", y="push", data=result_asm)
plt.title("boxplot of .asm push opcode")
plt.show()
```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

4.2.2 Multivariate Analysis on .asm file features

```
[42]: # check out the course content for more explanation on tsne algorithm
# https://www.appliedaicourse.com/course/applied-ai-course-online/lessons/
# → t-distributed-stochastic-neighbourhood-embedding-t-sne-part-1/

# multivariate analysis on byte files
# this is with perplexity 50
```

```

xtsne=TSNE(perplexity=50)
results=xtsne.fit_transform(result_asm.drop(['ID','Class'], axis=1).fillna(0))
vis_x = results[:, 0]
vis_y = results[:, 1]
plt.scatter(vis_x, vis_y, c=data_y, cmap=plt.cm.get_cmap("jet", 9))
plt.colorbar(ticks=range(10))
plt.clim(0.5, 9)
plt.show()

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```

[43]: # by univariate analysis on the .asm file features we are getting very
      →negligible information from
      # 'rtn', '.BSS:', '.CODE' features, so here we are trying multivariate analysis
      →after removing those features
      # the plot looks very messy

xtsne=TSNE(perplexity=30)
results=xtsne.fit_transform(result_asm.drop(['ID','Class', 'rtn', '.BSS:', '.
      →CODE','size'], axis=1))
vis_x = results[:, 0]
vis_y = results[:, 1]
plt.scatter(vis_x, vis_y, c=data_y, cmap=plt.cm.get_cmap("jet", 9))
plt.colorbar(ticks=range(10))
plt.clim(0.5, 9)
plt.show()

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

4.2.3 Conclusion on EDA

We have taken only 52 features from asm files (after reading through many blogs and research papers)

The univariate analysis was done only on few important features.

Take-aways

1. Class 3 can be easily separated because of the frequency of segments, opcodes and keywords being less
2. Each feature has its unique importance in separating the Class labels.


```
[ ]:
```

4.2.4 Adding pixel density features

```
[44]: import numpy, scipy.misc, os, array
```

```
def read_image(filename):  
    f = open(filename, 'rb')  
    ln = os.path.getsize(filename) # length of file in bytes  
    width = 256  
    rem = ln%width  
    a = array.array("B") # uint8 array  
    a.fromfile(f, ln-rem)  
    f.close()  
    g = numpy.reshape(a, (len(a)//width, width))  
    g = numpy.uint8(g)  
    f = g.copy()  
    f.resize((1000,))  
    return list(f)
```

```
[45]: image_1000_features = result_asm.ID.apply(lambda x: (read_image('asmFiles/'+x+'.  
→asm'))).apply(pd.Series)
```

```
[46]: image_1000_features.columns = [x+10000 for x in image_1000_features.columns]
```

```
[47]: image_1000_features.head()
```

```
[47]:
```

	10000	10001	10002	10003	10004	10005	10006	10007	10008	10009	...	\
0	72	69	65	68	69	82	58	49	48	48	...	
1	72	69	65	68	69	82	58	54	67	53	...	
2	72	69	65	68	69	82	58	49	48	48	...	
3	72	69	65	68	69	82	58	49	49	48	...	
4	72	69	65	68	69	82	58	55	54	51	...	

	10990	10991	10992	10993	10994	10995	10996	10997	10998	10999
0	71	77	69	78	84	32	72	69	65	68
1	32	32	32	32	58	9	80	111	114	116
2	59	32	70	111	114	109	97	116	9	32
3	32	32	32	32	58	9	80	111	114	116
4	32	32	32	32	58	9	80	111	114	116

[5 rows x 1000 columns]

```
[48]: result_asm = pd.concat([result_asm, image_1000_features], axis=1)
```

```
[49]: result_asm.head()
```

```
[49]:
```

	ID	HEADER:	.text:	.Pav:	.idata:	.data:	.bss:	\
0	01kcPWA9K2B0xQeS5Rju	0.107345	0.001092	0.0	0.000761	0.000023	0.0	
1	1E93CpP60RHFNiT5Qfvn	0.096045	0.001230	0.0	0.000617	0.000019	0.0	
2	3ekVow2ajZHbTnBcsDfX	0.096045	0.000627	0.0	0.000300	0.000017	0.0	
3	3X2nY7iQaPBIWDrAZqJe	0.096045	0.000333	0.0	0.000258	0.000008	0.0	
4	460ZzdsSKDCFV8h7XWxf	0.096045	0.000590	0.0	0.000353	0.000068	0.0	

	.rdata:	.edata:	.rsrc:	...	10990	10991	10992	10993	10994	10995	\
0	0.000084	0.0	0.000072	...	71	77	69	78	84	32	
1	0.000000	0.0	0.000072	...	32	32	32	32	58	9	
2	0.000038	0.0	0.000072	...	59	32	70	111	114	109	
3	0.000000	0.0	0.000072	...	32	32	32	32	58	9	
4	0.000000	0.0	0.000072	...	32	32	32	32	58	9	

	10996	10997	10998	10999
0	72	69	65	68
1	80	111	114	116
2	97	116	9	32
3	80	111	114	116
4	80	111	114	116

[5 rows x 1054 columns]

4.3 Train and test split

```
[50]: asm_y = result_asm['Class']
      asm_x = result_asm.drop(['ID', 'Class', '.BSS:', 'rtn', '.CODE'], axis=1)

[51]: X_train_asm, X_test_asm, y_train_asm, y_test_asm = train_test_split(asm_x, asm_y,
      ↪ stratify=asm_y, test_size=0.20)
      X_train_asm, X_cv_asm, y_train_asm, y_cv_asm = train_test_split(X_train_asm,
      ↪ y_train_asm, stratify=y_train_asm, test_size=0.20)

[52]: #print( X_cv_asm.isnull().all())
```

4.4. Machine Learning models on features of .asm files

4.4.1 K-Nearest Neighbors

```
[53]: # find more about KNeighborsClassifier() here http://scikit-learn.org/stable/
      ↪ modules/generated/sklearn.neighbors.KNeighborsClassifier.html
      # -----
      # default parameter
      # KNeighborsClassifier(n_neighbors=5, weights=uniform, algorithm=auto,
      ↪ leaf_size=30, p=2,
      # metric=minkowski, metric_params=None, n_jobs=1, **kwargs)

      # methods of
      # fit(X, y) : Fit the model using X as training data and y as target values
      # predict(X): Predict the class labels for the provided data
      # predict_proba(X): Return probability estimates for the test data X.
      # -----
      # video link: https://www.appliedaicourse.com/course/applied-ai-course-online/
      ↪ lessons/k-nearest-neighbors-geometric-intuition-with-a-toy-example-1/
      # -----
```

```

# find more about CalibratedClassifierCV here at http://scikit-learn.org/stable/modules/generated/sklearn.calibration.CalibratedClassifierCV.html
# -----
# default paramters
# sklearn.calibration.CalibratedClassifierCV(base_estimator=None,
# →method=sigmoid, cv=3)
#
# some of the methods of CalibratedClassifierCV()
# fit(X, y[, sample_weight])          Fit the calibrated model
# get_params([deep])                 Get parameters for this estimator.
# predict(X)                         Predict the target of new samples.
# predict_proba(X)                   Posterior probabilities of classification
#-----
# video link:
#-----

alpha = [x for x in range(1, 21,2)]
cv_log_error_array=[]
for i in alpha:
    print('For k = ',i)
    k_cfl=KNeighborsClassifier(n_neighbors=i)
    k_cfl.fit(X_train_asm,y_train_asm)
    sig_clf = CalibratedClassifierCV(k_cfl, method="sigmoid")
    sig_clf.fit(X_train_asm, y_train_asm)
    predict_y = sig_clf.predict_proba(X_cv_asm)
    cv_log_error_array.append(log_loss(y_cv_asm, predict_y, labels=k_cfl.
→classes_, eps=1e-15))

for i in range(len(cv_log_error_array)):
    print ('log_loss for k = ',alpha[i],'is',cv_log_error_array[i])

best_alpha = np.argmin(cv_log_error_array)

fig, ax = plt.subplots()
ax.plot(alpha, cv_log_error_array,c='g')
for i, txt in enumerate(np.round(cv_log_error_array,3)):
    ax.annotate((alpha[i],np.round(txt,3)), (alpha[i],cv_log_error_array[i]))
plt.grid()
plt.title("Cross Validation Error for each alpha")
plt.xlabel("Alpha i's")
plt.ylabel("Error measure")
plt.show()

k_cfl=KNeighborsClassifier(n_neighbors=alpha[best_alpha])
k_cfl.fit(X_train_asm,y_train_asm)
sig_clf = CalibratedClassifierCV(k_cfl, method="sigmoid")
sig_clf.fit(X_train_asm, y_train_asm)

```

```

pred_y=sig_clf.predict(X_test_asm)

predict_y = sig_clf.predict_proba(X_train_asm)
print ('log loss for train data',log_loss(y_train_asm, predict_y))
predict_y = sig_clf.predict_proba(X_cv_asm)
print ('log loss for cv data',log_loss(y_cv_asm, predict_y))
predict_y = sig_clf.predict_proba(X_test_asm)
print ('log loss for test data',log_loss(y_test_asm, predict_y))
plot_confusion_matrix(y_test_asm,sig_clf.predict(X_test_asm))

```

```

For k = 1
For k = 3
For k = 5
For k = 7
For k = 9
For k = 11
For k = 13
For k = 15
For k = 17
For k = 19
log_loss for k = 1 is 0.03957063826268944
log_loss for k = 3 is 0.051404140483720065
log_loss for k = 5 is 0.060458399090755545
log_loss for k = 7 is 0.06926503500170093
log_loss for k = 9 is 0.07662482937608156
log_loss for k = 11 is 0.08533517515537771
log_loss for k = 13 is 0.09201851136153923
log_loss for k = 15 is 0.09951338808060389
log_loss for k = 17 is 0.10600066326872025
log_loss for k = 19 is 0.11164566751230305

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```

log loss for train data 0.014493418765645175
log loss for cv data 0.03957063826268944
log loss for test data 0.060795224711157655
Number of misclassified points 0.8739650413983441
----- Confusion matrix
-----

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

----- Precision matrix

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Sum of columns in precision matrix [1. 1. 1. 1. 1. 1. 1. 1. 1.]
----- Recall matrix

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Sum of rows in precision matrix [1. 1. 1. 1. 1. 1. 1. 1. 1.]

4.4.2 Logistic Regression

```
[54]: # read more about SGDClassifier() at http://scikit-learn.org/stable/modules/generated/sklearn.linear\_model.SGDClassifier.html
# -----
# default parameters
# SGDClassifier(loss=hinge, penalty=l2, alpha=0.0001, l1_ratio=0.15,
#   →fit_intercept=True, max_iter=None, tol=None,
# shuffle=True, verbose=0, epsilon=0.1, n_jobs=1, random_state=None,
#   →learning_rate=optimal, eta0=0.0, power_t=0.5,
# class_weight=None, warm_start=False, average=False, n_iter=None)

# some of methods
# fit(X, y[, coef_init, intercept_init, ])          Fit linear model with
#   →Stochastic Gradient Descent.
# predict(X)          Predict class labels for samples in X.

#-----
# video link: https://www.appliedaicourse.com/course/applied-ai-course-online/lessons/geometric-intuition-1/
#-----

alpha = [10 ** x for x in range(-5, 4)]
cv_log_error_array=[]
```

```

for i in alpha:
    print('For C = ',i)
    logisticR=LogisticRegression(penalty='l2',C=i,class_weight='balanced')
    logisticR.fit(X_train_asm,y_train_asm)
    sig_clf = CalibratedClassifierCV(logisticR, method="sigmoid")
    sig_clf.fit(X_train_asm, y_train_asm)
    predict_y = sig_clf.predict_proba(X_cv_asm)
    cv_log_error_array.append(log_loss(y_cv_asm, predict_y, labels=logisticR.
    →classes_, eps=1e-15))

for i in range(len(cv_log_error_array)):
    print ('log_loss for c = ',alpha[i],'is',cv_log_error_array[i])

best_alpha = np.argmin(cv_log_error_array)

fig, ax = plt.subplots()
ax.plot(alpha, cv_log_error_array,c='g')
for i, txt in enumerate(np.round(cv_log_error_array,3)):
    ax.annotate((alpha[i],np.round(txt,3)), (alpha[i],cv_log_error_array[i]))
plt.grid()
plt.title("Cross Validation Error for each alpha")
plt.xlabel("Alpha i's")
plt.ylabel("Error measure")
plt.show()

logisticR=LogisticRegression(penalty='l2',C=alpha[best_alpha],class_weight='balanced')
logisticR.fit(X_train_asm,y_train_asm)
sig_clf = CalibratedClassifierCV(logisticR, method="sigmoid")
sig_clf.fit(X_train_asm, y_train_asm)

predict_y = sig_clf.predict_proba(X_train_asm)
print ('log loss for train data',(log_loss(y_train_asm, predict_y,
    →labels=logisticR.classes_, eps=1e-15)))
predict_y = sig_clf.predict_proba(X_cv_asm)
print ('log loss for cv data',(log_loss(y_cv_asm, predict_y, labels=logisticR.
    →classes_, eps=1e-15)))
predict_y = sig_clf.predict_proba(X_test_asm)
print ('log loss for test data',(log_loss(y_test_asm, predict_y,
    →labels=logisticR.classes_, eps=1e-15)))
plot_confusion_matrix(y_test_asm,sig_clf.predict(X_test_asm))

```

```

For C = 1e-05
For C = 0.0001
For C = 0.001
For C = 0.01
For C = 0.1
For C = 1

```

```

For C = 10
For C = 100
For C = 1000
log_loss for c = 1e-05 is 0.6124787071799669
log_loss for c = 0.0001 is 0.4751208970123429
log_loss for c = 0.001 is 0.3977660473853282
log_loss for c = 0.01 is 0.3760540596894935
log_loss for c = 0.1 is 0.3611079524704754
log_loss for c = 1 is 0.34044976162745205
log_loss for c = 10 is 0.34172723104482733
log_loss for c = 100 is 0.33636416411211234
log_loss for c = 1000 is 0.35344162276452423

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

log loss for train data 0.3212305712115807
log loss for cv data 0.33636416411211234
log loss for test data 0.3252260819309329
Number of misclassified points 10.027598896044159
----- Confusion matrix
-----

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

----- Precision matrix
-----

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Sum of columns in precision matrix [ 1.  1.  1.  1. nan  1.  1.  1.  1.]
----- Recall matrix
-----

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Sum of rows in precision matrix [1.  1.  1.  1.  1.  1.  1.  1.  1.]

```

4.4.3 Random Forest Classifier

```

[55]: # -----
# default parameters
# sklearn.ensemble.RandomForestClassifier(n_estimators=10, criterion=gini,
#     ↳max_depth=None, min_samples_split=2,
# min_samples_leaf=1, min_weight_fraction_leaf=0.0, max_features=auto,
#     ↳max_leaf_nodes=None, min_impurity_decrease=0.0,
# min_impurity_split=None, bootstrap=True, oob_score=False, n_jobs=1,
#     ↳random_state=None, verbose=0, warm_start=False,
# class_weight=None)

# Some of methods of RandomForestClassifier()
# fit(X, y, [sample_weight])      Fit the SVM model according to the given
#     ↳training data.
# predict(X)      Perform classification on samples in X.
# predict_proba(X)      Perform classification on samples in X.

# some of attributes of RandomForestClassifier()
# feature_importances_ : array of shape = [n_features]
# The feature importances (the higher, the more important the feature).

# -----
# video link: https://www.appliedaicourse.com/course/applied-ai-course-online/
#     ↳lessons/random-forest-and-their-construction-2/
# -----

alpha=[10,50,100,500,1000,2000,3000]
cv_log_error_array=[]
for i in alpha:
    print('For estimator = ',i)
    r_cfl=RandomForestClassifier(n_estimators=i,random_state=42,n_jobs=-1)
    r_cfl.fit(X_train_asm,y_train_asm)
    sig_clf = CalibratedClassifierCV(r_cfl, method="sigmoid")
    sig_clf.fit(X_train_asm, y_train_asm)
    predict_y = sig_clf.predict_proba(X_cv_asm)
    cv_log_error_array.append(log_loss(y_cv_asm, predict_y, labels=r_cfl.
    ↳classes_, eps=1e-15))

for i in range(len(cv_log_error_array)):
    print ('log_loss for c = ',alpha[i], 'is',cv_log_error_array[i])

best_alpha = np.argmin(cv_log_error_array)

fig, ax = plt.subplots()
ax.plot(alpha, cv_log_error_array,c='g')
for i, txt in enumerate(np.round(cv_log_error_array,3)):
    ax.annotate((alpha[i],np.round(txt,3)), (alpha[i],cv_log_error_array[i]))

```



```

plt.grid()
plt.title("Cross Validation Error for each alpha")
plt.xlabel("Alpha i's")
plt.ylabel("Error measure")
plt.show()

r_cfl=RandomForestClassifier(n_estimators=alpha[best_alpha],random_state=42,n_jobs=-1)
r_cfl.fit(X_train_asm,y_train_asm)
sig_clf = CalibratedClassifierCV(r_cfl, method="sigmoid")
sig_clf.fit(X_train_asm, y_train_asm)
predict_y = sig_clf.predict_proba(X_train_asm)
print ('log loss for train data',(log_loss(y_train_asm, predict_y,
→labels=sig_clf.classes_, eps=1e-15)))
predict_y = sig_clf.predict_proba(X_cv_asm)
print ('log loss for cv data',(log_loss(y_cv_asm, predict_y, labels=sig_clf.
→classes_, eps=1e-15)))
predict_y = sig_clf.predict_proba(X_test_asm)
print ('log loss for test data',(log_loss(y_test_asm, predict_y, labels=sig_clf.
→classes_, eps=1e-15)))
plot_confusion_matrix(y_test_asm,sig_clf.predict(X_test_asm))

```

```

For estimator = 10
For estimator = 50
For estimator = 100
For estimator = 500
For estimator = 1000
For estimator = 2000
For estimator = 3000
log_loss for c = 10 is 0.019683804059731778
log_loss for c = 50 is 0.017718381172568694
log_loss for c = 100 is 0.01644343749170818
log_loss for c = 500 is 0.016054768918798396
log_loss for c = 1000 is 0.01579015124916087
log_loss for c = 2000 is 0.015750874531983823
log_loss for c = 3000 is 0.015864170478007575

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```

log loss for train data 0.01033319416987003
log loss for cv data 0.015750874531983823
log loss for test data 0.021136720289532153
Number of misclassified points 0.36798528058877644
----- Confusion matrix
-----

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

----- Precision matrix

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Sum of columns in precision matrix [1. 1. 1. 1. 1. 1. 1. 1. 1.]
----- Recall matrix

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Sum of rows in precision matrix [1. 1. 1. 1. 1. 1. 1. 1. 1.]

4.4.4 XgBoost Classifier

```
[56]: # Training a hyper-parameter tuned Xg-Boost regressor on our train data

# find more about XGBClassifier function here http://xgboost.readthedocs.io/en/latest/python/python\_api.html?#xgboost.XGBClassifier
# -----
# default paramters
# class xgboost.XGBClassifier(max_depth=3, learning_rate=0.1, n_estimators=100,
#   silent=True,
#   objective='binary:logistic', booster='gbtree', n_jobs=1, nthread=None,
#   gamma=0, min_child_weight=1,
#   max_delta_step=0, subsample=1, colsample_bytree=1, colsample_bylevel=1,
#   reg_alpha=0, reg_lambda=1,
#   scale_pos_weight=1, base_score=0.5, random_state=0, seed=None, missing=None,
#   **kwargs)

# some of methods of RandomForestRegressor()
# fit(X, y, sample_weight=None, eval_set=None, eval_metric=None,
#   early_stopping_rounds=None, verbose=True, xgb_model=None)
# get_params([deep])          Get parameters for this estimator.
```

```

# predict(data, output_margin=False, ntree_limit=0) : Predict with data. NOTE:
→ This function is not thread safe.
# get_score(importance_type='weight') -> get the feature importance
# -----
# video link2: https://www.appliedaicourse.com/course/applied-ai-course-online/lessons/what-are-ensembles/
# -----

alpha=[10,50,100,500,1000,2000,3000]
cv_log_error_array=[]
for i in alpha:
    print('For estimator = ',i)
    x_cfl=XGBClassifier(n_estimators=i,nthread=-1)
    x_cfl.fit(X_train_asm,y_train_asm)
    sig_clf = CalibratedClassifierCV(x_cfl, method="sigmoid")
    sig_clf.fit(X_train_asm, y_train_asm)
    predict_y = sig_clf.predict_proba(X_cv_asm)
    cv_log_error_array.append(log_loss(y_cv_asm, predict_y, labels=x_cfl.
→ classes_, eps=1e-15))

for i in range(len(cv_log_error_array)):
    print ('log_loss for c = ',alpha[i],'is',cv_log_error_array[i])

best_alpha = np.argmin(cv_log_error_array)

fig, ax = plt.subplots()
ax.plot(alpha, cv_log_error_array,c='g')
for i, txt in enumerate(np.round(cv_log_error_array,3)):
    ax.annotate((alpha[i],np.round(txt,3)), (alpha[i],cv_log_error_array[i]))
plt.grid()
plt.title("Cross Validation Error for each alpha")
plt.xlabel("Alpha i's")
plt.ylabel("Error measure")
plt.show()

x_cfl=XGBClassifier(n_estimators=alpha[best_alpha],nthread=-1)
x_cfl.fit(X_train_asm,y_train_asm)
sig_clf = CalibratedClassifierCV(x_cfl, method="sigmoid")
sig_clf.fit(X_train_asm, y_train_asm)

predict_y = sig_clf.predict_proba(X_train_asm)

print ('For values of best alpha = ', alpha[best_alpha], "The train log loss is:
→ ",log_loss(y_train_asm, predict_y))
predict_y = sig_clf.predict_proba(X_cv_asm)

```

```

print('For values of best alpha = ', alpha[best_alpha], "The cross validation_
→log loss is:",log_loss(y_cv_asm, predict_y))
predict_y = sig_clf.predict_proba(X_test_asm)
print('For values of best alpha = ', alpha[best_alpha], "The test log loss is:
→",log_loss(y_test_asm, predict_y))
plot_confusion_matrix(y_test_asm,sig_clf.predict(X_test_asm))

```

```

For estimator = 10
For estimator = 50
For estimator = 100
For estimator = 500
For estimator = 1000
For estimator = 2000
For estimator = 3000
log_loss for c = 10 is 0.04992282929194357
log_loss for c = 50 is 0.02598101172114864
log_loss for c = 100 is 0.01779345310522223
log_loss for c = 500 is 0.014743518272656544
log_loss for c = 1000 is 0.014555198646745888
log_loss for c = 2000 is 0.014459489185114926
log_loss for c = 3000 is 0.01445476662318132

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```

For values of best alpha = 3000 The train log loss is: 0.008733764783017739
For values of best alpha = 3000 The cross validation log loss is:
0.01445476662318132
For values of best alpha = 3000 The test log loss is: 0.03316661864010893
Number of misclassified points 0.45998160073597055

```

```

----- Confusion matrix
-----

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

```

----- Precision matrix
-----

```

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Sum of columns in precision matrix [1. 1. 1. 1. 1. 1. 1. 1. 1.]
----- Recall matrix

<IPython.core.display.Javascript object>

<IPython.core.display.HTML object>

Sum of rows in precision matrix [1. 1. 1. 1. 1. 1. 1. 1. 1.]

4.4.5 Xgboost Classifier with best hyperparameters

```
[57]: x_cfl=XGBClassifier()

prams={
    'learning_rate': [0.01,0.03,0.05,0.1,0.15,0.2],
    'n_estimators': [100,200,500,1000,2000],
    'max_depth': [3,5,10],
    'colsample_bytree': [0.1,0.3,0.5,1],
    'subsample': [0.1,0.3,0.5,1]
}
random_cfl=RandomizedSearchCV(x_cfl,param_distributions=prams,verbose=1,n_jobs=-1,)
random_cfl.fit(X_train_asm,y_train_asm)
```

Fitting 3 folds for each of 10 candidates, totalling 30 fits

[Parallel(n_jobs=-1)]: Using backend LokyBackend with 6 concurrent workers.

[Parallel(n_jobs=-1)]: Done 30 out of 30 | elapsed: 13.5min finished

```
[57]: RandomizedSearchCV(cv='warn', error_score='raise-deprecating',
                        estimator=XGBClassifier(base_score=0.5, booster='gbtree',
                                                colsample_bylevel=1,
                                                colsample_bynode=1,
                                                colsample_bytree=1, gamma=0,
                                                learning_rate=0.1, max_delta_step=0,
                                                max_depth=3, min_child_weight=1,
                                                missing=None, n_estimators=100,
                                                n_jobs=1, nthread=None,
                                                objective='binary:logistic',
                                                random_state=0, reg_al...
                                                seed=None, silent=None, subsample=1,
                                                verbosity=1),
                        iid='warn', n_iter=10, n_jobs=-1,
                        param_distributions={'colsample_bytree': [0.1, 0.3, 0.5, 1],
```

```

        'learning_rate': [0.01, 0.03, 0.05, 0.1,
                           0.15, 0.2],
        'max_depth': [3, 5, 10],
        'n_estimators': [100, 200, 500, 1000,
                          2000],
        'subsample': [0.1, 0.3, 0.5, 1]},
    pre_dispatch='2*n_jobs', random_state=None, refit=True,
    return_train_score=False, scoring=None, verbose=1)

```

[58]: `print (random_cfl.best_params_)`

```

{'subsample': 1, 'n_estimators': 1000, 'max_depth': 5, 'learning_rate': 0.15,
 'colsample_bytree': 0.1}

```

[59]: *# Training a hyper-parameter tuned Xg-Boost regressor on our train data*

find more about XGBClassifier function here http://xgboost.readthedocs.io/en/latest/python/python_api.html?#xgboost.XGBClassifier

-----

default paramters

*# class xgboost.XGBClassifier(max_depth=3, learning_rate=0.1, n_estimators=100,
→silent=True,*

*# objective='binary:logistic', booster='gbtree', n_jobs=1, nthread=None,
→gamma=0, min_child_weight=1,*

*# max_delta_step=0, subsample=1, colsample_bytree=1, colsample_bylevel=1,
→reg_alpha=0, reg_lambda=1,*

*# scale_pos_weight=1, base_score=0.5, random_state=0, seed=None, missing=None,
→**kwargs)*

some of methods of RandomForestRegressor()

*# fit(X, y, sample_weight=None, eval_set=None, eval_metric=None,
→early_stopping_rounds=None, verbose=True, xgb_model=None)*

get_params([deep]) Get parameters for this estimator.

*# predict(data, output_margin=False, ntree_limit=0) : Predict with data. NOTE:
→This function is not thread safe.*

get_score(importance_type='weight') -> get the feature importance

-----

video link2: <https://www.appliedaicourse.com/course/applied-ai-course-online/lessons/what-are-ensembles/>

-----

```

x_cfl=XGBClassifier(subsample= 0.3, n_estimators= 1000, max_depth= 10,   
→learning_rate= 0.03, colsample_bytree= 0.5)
x_cfl.fit(X_train_asm,y_train_asm)
c_cfl=CalibratedClassifierCV(x_cfl,method='sigmoid')
c_cfl.fit(X_train_asm,y_train_asm)

```

```

predict_y = c_cfl.predict_proba(X_train_asm)
print ('train loss',log_loss(y_train_asm, predict_y))
predict_y = c_cfl.predict_proba(X_cv_asm)
print ('cv loss',log_loss(y_cv_asm, predict_y))
predict_y = c_cfl.predict_proba(X_test_asm)
print ('test loss',log_loss(y_test_asm, predict_y))

```

```

train loss 0.009759075491612286
cv loss 0.015404153604484504
test loss 0.027258792256670696

```

4.5. Machine Learning models on features of both .asm and .bytes files

4.5.1. Merging both asm and byte file features

[60]: result.head()

```

[60]: Unnamed: 0      00      0000      0001      0002      0003      0004  \
0      0.000000  0.262806  0.127389  0.079943  0.054323  0.088980  0.064972
1      0.001001  0.017358  0.009262  0.057364  0.004340  0.003270  0.010955
2      0.002002  0.040827  0.007479  0.047232  0.010648  0.010951  0.035070
3      0.003003  0.009209  0.004620  0.016276  0.004001  0.005248  0.007097
4      0.004004  0.008629  0.007132  0.004627  0.001356  0.008366  0.000551

      0005      0006      0007  ...      FFF9      FFFA      FFFB      FFFC  \
0  0.090303  0.109255  0.121901  ...  0.001933  0.003526  0.001031  0.001188
1  0.001075  0.000583  0.001215  ...  0.019884  0.031735  0.009574  0.001619
2  0.063427  0.053511  0.017744  ...  0.022370  0.076164  0.017381  0.007125
3  0.003655  0.001845  0.002552  ...  0.002486  0.009168  0.002504  0.009284
4  0.001183  0.000291  0.000608  ...  0.006628  0.001410  0.000147  0.000000

      FFFD      FFFE      FFFF      ID      size  Class
0  0.001294  0.000759  0.001227  01azqd4InC7m9JpocGv5  0.092219      9
1  0.014518  0.015811  0.006936  01IsoiSMh5gxyDYTl4CB  0.121236      2
2  0.013943  0.010625  0.003393  01jsnpXSAlgW6aPeDxrU  0.084499      9
3  0.003450  0.007969  0.001954  01kcPWA9K2B0xQeS5Rju  0.010759      1
4  0.000575  0.000379  0.000108  01SuzwMJEIXsK7A8dQbl  0.006233      8

```

[5 rows x 66036 columns]

[61]: result_asm.head()

```

[61]: ID  HEADER:  .text:  .Pav:  .idata:  .data:  .bss:  \
0  01kcPWA9K2B0xQeS5Rju  0.107345  0.001092  0.0  0.000761  0.000023  0.0
1  1E93CpP60RHFNiT5Qfvn  0.096045  0.001230  0.0  0.000617  0.000019  0.0
2  3ekVow2ajZHbTnBcsDfX  0.096045  0.000627  0.0  0.000300  0.000017  0.0
3  3X2nY7iQaPBIWDrAZqJe  0.096045  0.000333  0.0  0.000258  0.000008  0.0
4  460ZzdsSKDCFV8h7XWxf  0.096045  0.000590  0.0  0.000353  0.000068  0.0

      .rdata:  .edata:  .rsrc:  ...  10990  10991  10992  10993  10994  10995  \
0  0.000084  0.0  0.000072  ...  71  77  69  78  84  32

```

1	0.000000	0.0	0.000072	...	32	32	32	32	58	9
2	0.000038	0.0	0.000072	...	59	32	70	111	114	109
3	0.000000	0.0	0.000072	...	32	32	32	32	58	9
4	0.000000	0.0	0.000072	...	32	32	32	32	58	9

	10996	10997	10998	10999
0	72	69	65	68
1	80	111	114	116
2	97	116	9	32
3	80	111	114	116
4	80	111	114	116

[5 rows x 1054 columns]

```
[62]: print(result.shape)
      print(result_asm.shape)
```

(10868, 66036)

(10868, 1054)

```
[63]: result_x = pd.merge(result,result_asm.drop(['Class'], axis=1),on='ID',
      ↪how='left')
      result_y = result_x['Class']
      result_x = result_x.drop(['ID','rtn','.BSS:','.CODE','Class'], axis=1)
      result_x.head()
```

[63]:	Unnamed: 0	00	0000	0001	0002	0003	0004	\
0	0.000000	0.262806	0.127389	0.079943	0.054323	0.088980	0.064972	
1	0.001001	0.017358	0.009262	0.057364	0.004340	0.003270	0.010955	
2	0.002002	0.040827	0.007479	0.047232	0.010648	0.010951	0.035070	
3	0.003003	0.009209	0.004620	0.016276	0.004001	0.005248	0.007097	
4	0.004004	0.008629	0.007132	0.004627	0.001356	0.008366	0.000551	

	0005	0006	0007	...	10990	10991	10992	10993	10994	\
0	0.090303	0.109255	0.121901	...	116	101	120	116	58	
1	0.001075	0.000583	0.001215	...	10	46	116	101	120	
2	0.063427	0.053511	0.017744	...	116	101	120	116	58	
3	0.003655	0.001845	0.002552	...	71	77	69	78	84	
4	0.001183	0.000291	0.000608	...	116	101	120	116	58	

	10995	10996	10997	10998	10999
0	48	48	52	48	49
1	116	58	48	48	52
2	48	48	52	48	49
3	32	72	69	65	68
4	48	48	52	48	49

[5 rows x 67083 columns]

4.5.2. Train and Test split

```
[64]: X_train, X_test_merge, y_train, y_test_merge = train_test_split(result_x,
    ↳ result_y, stratify=result_y, test_size=0.20)
X_train_merge, X_cv_merge, y_train_merge, y_cv_merge =
    ↳ train_test_split(X_train, y_train, stratify=y_train, test_size=0.20)
```

4.5.3. Xgboost Classifier with best hyperparameters

```
[65]: x_cfl=XGBClassifier()

prams={
    'learning_rate': [0.01, 0.03, 0.05, 0.1, 0.15, 0.2],
    'n_estimators': [100, 200, 500, 1000, 2000],
    'max_depth': [3, 5, 10],
    'colsample_bytree': [0.1, 0.3, 0.5, 1],
    'subsample': [0.1, 0.3, 0.5, 1]
}
random_cfl=RandomizedSearchCV(x_cfl, param_distributions=prams, verbose=1, n_jobs=4,)
random_cfl.fit(X_train_asm, y_train_asm)
```

Fitting 3 folds for each of 10 candidates, totalling 30 fits

[Parallel(n_jobs=4)]: Using backend LokyBackend with 4 concurrent workers.

[Parallel(n_jobs=4)]: Done 30 out of 30 | elapsed: 20.5min finished

```
[65]: RandomizedSearchCV(cv='warn', error_score='raise-deprecating',
    estimator=XGBClassifier(base_score=0.5, booster='gbtree',
        colsample_bylevel=1,
        colsample_bynode=1,
        colsample_bytree=1, gamma=0,
        learning_rate=0.1, max_delta_step=0,
        max_depth=3, min_child_weight=1,
        missing=None, n_estimators=100,
        n_jobs=1, nthread=None,
        objective='binary:logistic',
        random_state=0, reg_al...
        seed=None, silent=None, subsample=1,
        verbosity=1),
    iid='warn', n_iter=10, n_jobs=4,
    param_distributions={'colsample_bytree': [0.1, 0.3, 0.5, 1],
        'learning_rate': [0.01, 0.03, 0.05, 0.1,
            0.15, 0.2],
        'max_depth': [3, 5, 10],
        'n_estimators': [100, 200, 500, 1000,
            2000],
        'subsample': [0.1, 0.3, 0.5, 1]},
    pre_dispatch='2*n_jobs', random_state=None, refit=True,
    return_train_score=False, scoring=None, verbose=1)
```

```
[66]: print (random_cfl.best_params_)
```

```
{'subsample': 1, 'n_estimators': 1000, 'max_depth': 10, 'learning_rate': 0.15, 'colsample_bytree': 0.1}
```

```
[67]: # Training a hyper-parameter tuned Xg-Boost regressor on our train data

# find more about XGBClassifier function here http://xgboost.readthedocs.io/en/latest/python/python\_api.html?#xgboost.XGBClassifier
# -----
# default parameters
# class xgboost.XGBClassifier(max_depth=3, learning_rate=0.1, n_estimators=100,
#     silent=True,
# objective='binary:logistic', booster='gbtree', n_jobs=1, nthread=None,
#     gamma=0, min_child_weight=1,
# max_delta_step=0, subsample=1, colsample_bytree=1, colsample_bylevel=1,
#     reg_alpha=0, reg_lambda=1,
# scale_pos_weight=1, base_score=0.5, random_state=0, seed=None, missing=None,
#     **kwargs)

# some of methods of RandomForestRegressor()
# fit(X, y, sample_weight=None, eval_set=None, eval_metric=None,
#     early_stopping_rounds=None, verbose=True, xgb_model=None)
# get_params([deep])          Get parameters for this estimator.
# predict(data, output_margin=False, ntree_limit=0) : Predict with data. NOTE:
#     This function is not thread safe.
# get_score(importance_type='weight') -> get the feature importance
# -----
# video link2: https://www.appliedaicourse.com/course/applied-ai-course-online/lessons/what-are-ensembles/
# -----

x_cfl=XGBClassifier(subsample= 0.3, n_estimators= 1000, max_depth= 10,
    learning_rate= 0.03, colsample_bytree= 0.5)
x_cfl.fit(X_train_asm,y_train_asm)
c_cfl=CalibratedClassifierCV(x_cfl,method='sigmoid')
c_cfl.fit(X_train_asm,y_train_asm)

predict_y = c_cfl.predict_proba(X_train_asm)
print ('train loss',log_loss(y_train_asm, predict_y))
predict_y = c_cfl.predict_proba(X_cv_asm)
print ('cv loss',log_loss(y_cv_asm, predict_y))
predict_y = c_cfl.predict_proba(X_test_asm)
print ('test loss',log_loss(y_test_asm, predict_y))
```

```
train loss 0.009759075491612286
```

```
cv loss 0.015404153604484504
test loss 0.027258792256670696
```

```
[68]: from prettytable import PrettyTable
x = PrettyTable()
x.field_names = ["Data", "Model", "Train logloss", "Test logloss"]
x.add_row(["Only byte files", "Logistic Regression", round(0.
→06645698722998557,3), round(0.18994698178499667,3)])
x.add_row(["asm files with image features", "XGBoost", round(0.
→009477617552245923,3), round(0.021151928630058183,3)])
x.add_row(["Both byte and asm files with image features", "XGboost", round(0.
→009759075491612286,3), round(0.027258792256670696,3)])
x.border=True
print(x)
```

```
+-----+-----+-----+
---+-----+
|          Data          |          Model          | Train
logloss | Test logloss |
+-----+-----+-----+
---+-----+
|          Only byte files          | Logistic Regression |      0.066
|      0.19      |
|          asm files with image features          |          XGBoost          |      0.009
|      0.021      |
| Both byte and asm files with image features | Logistic Regression |      0.11
|      0.122      |
+-----+-----+-----+
---+-----+
```

Clearly asm files with image fetures is the best Model. Using both byte files and asm files, did not impacted much on model.