

# **Configuration Manual**

MSc Research Project Cyber Security

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### National College of Ireland

#### **MSc Project Submission Sheet**



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Programme: Cyber Security

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Lecturer: Vanessa Ayala-Rivera Submission Due Date: 16/12/2021

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I hereby certify that the information contained in this (my submission) is information pertaining to research I conducted for this project. All information other than my own contribution will be fully referenced and listed in the relevant bibliography section at the rear of the project.

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# **Configuration Manual**

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### 1. Introduction

This configuration document was created to outline the steps for executing the research project and to specify the configuration of the system used to run the models. The stages involve downloading and installing the essential software and packages, as well as the minimal setup required for the project to work well.

### 2. System Configuration

Processor: Intel(R) Core (TM) i7 – 5500U CPU @ 2.40GHz RAM: 16 GB Storage: 500 GB HDD Operating System: Windows 10 64-bit operating system

### 3. Setup

### **Python Libraries version:**

Numpy 1.19.5 Pandas 1.1.15 Seaborn 0.11.2 Mathplotlib 3.2.2 Sklearn 1.0.2

### **Google Colaboratory:**

It is a service that allows users to run Python code in a browser online while requiring little resources and essentially no configuration which needs to be done locally. Only the following requirements for writing code using Google Colab. A browser such as Chrome or Firefox and a google account.

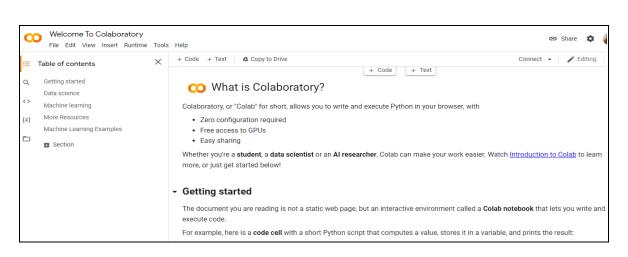


Fig 1. Google Colab

## 4. Implementation

Step 1. Importing necessary python libraries for dataset preprocessing

```
[4] # Importing the required libraries and modules
import numpy as np # linear algebra
import pandas as pd # data processing, CSV file I/O (e.g. pd.read_csv)
import pickle
import seaborn as sns
import matplotlib.pyplot as plt
from sklearn.ensemble import RandomForestClassifier
from sklearn.model_selection import train_test_split
from sklearn.metrics import classification_report, confusion_matrix
```

Fig 2. Libraries required for models

### Step 2. Loading and Preprocessing of the dataset

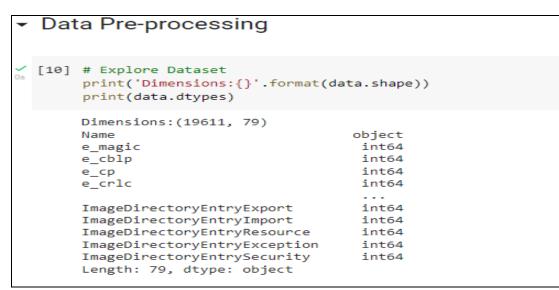


Fig 3. Data Preprocessing

] missing_data = pd.DataFrame(· missing_data	('total_missing	': data.isnull
	total_missing	perc_missing
Name	0	0.0
e_magic	0	0.0
e_cblp	0	0.0
e_cp	0	0.0
e_cric	0	0.0
ImageDirectoryEntryExport	0	0.0
ImageDirectoryEntryImport	0	0.0
ImageDirectoryEntryResource	0	0.0
ImageDirectoryEntryException	0	0.0
ImageDirectoryEntrySecurity	0	0.0
79 rows × 2 columns		

Fig 4. Checking for missing data in the dataset

#Statistical description for the data data.describe()												
	e_magic	e_cblp	e_cp	e_crlc	e_cparhdr	e_minalloc	e_maxalloc	e_ss				
count	19611.0	19611.000000	19611.000000	19611.000000	19611.000000	19611.000000	19611.000000	19611.000000				
mean	23117.0	178.615726	71.660752	49.146958	37.370710	37.032635	64178.739687	10.418490				
std	0.0	987.200729	1445.192977	1212.201919	864.515405	915.833139	9110.755873	637.116265				
min	23117.0	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000				
25%	23117.0	144.000000	3.000000	0.000000	4.000000	0.000000	65535.000000	0.000000				
50%	23117.0	144.000000	3.000000	0.000000	4.000000	0.000000	65535.000000	0.000000				
75%	23117.0	144.000000	3.000000	0.000000	4.000000	0.000000	65535.000000	0.000000				
max	23117.0	59448.000000	63200.000000	64613.000000	43690.000000	43690.000000	65535.000000	61436.00000				

Fig 5. Analysis of data in statistical terms

✓ [19] ₀s	fro enc	m sklea	arn.prepro data.appi	ocessing	impor	t LabelE			if x.dtype	== 'ot	ject'	else x)				
	Name e_magic e_cblp e_cp e_crlc e_cparhdr e_minalloc e_maxalloc e_ss e_sp e_csum e_ip e_cs e_lfarlc e_ov														e_ovno	
	0	7037	23117	144	3	0	4	0	65535	0	184	0	0	0	64	0
	1	11957	23117	144	3	0	4	0	65535	0	184	0	0	0	64	0
	2	11448	23117	144	3	0	4	0	65535	0	184	0	0	0	64	0
	3	5367	23117	144	3	0	4	0	65535	0	184	0	0	0	64	0
	4	3489	23117	144	3	0	4	0	65535	0	184	0	0	0	64	0
	(															

Fig 6. Converting nonnumeric to numeric

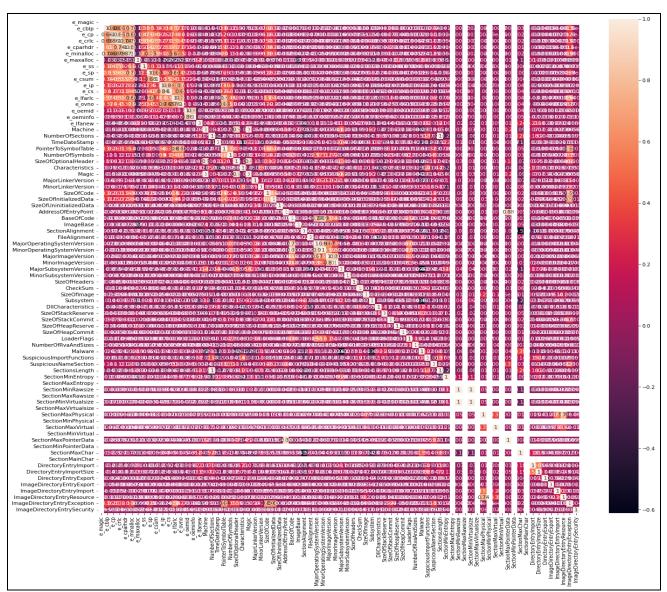


Fig 7. Correlation plot before encoding

fro enc	<pre>converting the non-numeric data into numeric data. com sklearn.preprocessing import LabelEncoder icoded = data.apply(lambda x: LabelEncoder().fit_transform(x) if x.dtype == 'object' else x) icoded.head()</pre>													
	Name	e_magic	e_cblp	e_cp	e_crlc	e_cparhdr	e_minalloc	e_maxalloc	e_ss	e_sp	e_csum	e_ip	e_cs	e_lfar
0	7037	23117	144	3	0	4	0	65535	0	184	0	0	0	(
1	11957	23117	144	3	0	4	0	65535	0	184	0	0	0	(
2	11448	23117	144	3	0	4	0	65535	0	184	0	0	0	(
3	5367	23117	144	3	0	4	0	65535	0	184	0	0	0	(
4	3489	23117	144	3	0	4	0	65535	0	184	0	0	0	(

Fig 8. Non numeric to numeric data conversion

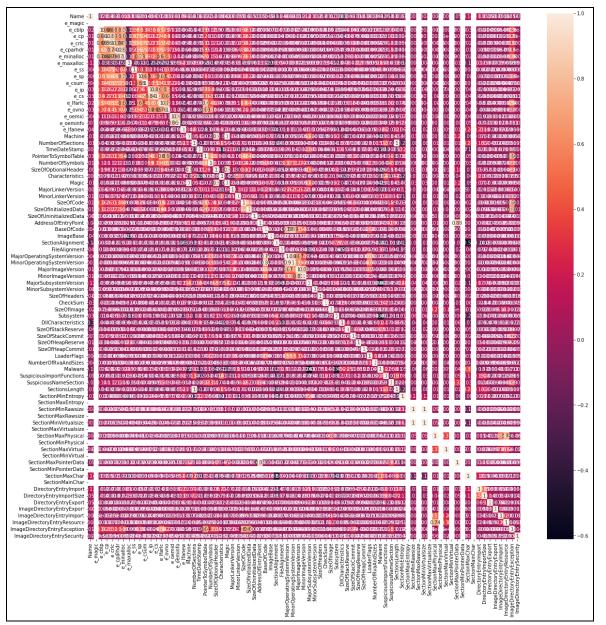


Fig 9. Correlation after label encoding

### **Step 4. Feature Selection**



Fig 9. Feature selection

# **Step 5. Splitting the dataset into Training and Testing in the ratio of 80:20**

```
# Splitting the data into training and testing data subset into 80% and 20% respectively
X_train, X_test, y_train, y_test = train_test_split(data_usage, data['Malware'], test_size=0.2, random_state=0)
print(f'Number of used features is {X_train.shape}')
print(f'Number of used features is {X_test.shape}')
print(f'Number of used features is {y_train.shape}')
print(f'Number of used features is {y_test.shape}')
Number of used features is (15688, 75)
Number of used features is (15688,)
Number of used features is (15688,)
Number of used features is (3923,)
```

Fig 10. Dataset split into training and test

### Step 6. Model building for Binary Classification

Ra	ndom fores	st model \	with bi	nary lab	elled data						
[]	<pre>rfc = RandomForestClassifier(n_estimators=100, random_state=0,</pre>										
[]	<pre># Classificat y_pred = rfc. print(classif</pre>	predict(X_te		:, y_pred,	target_names=['	Benign',	'Malware']))				
		precision	recall	f1-score	support						
	Benign Malware		0.96 1.00	0.97 0.99	1004 2919						
	accuracy macro avg weighted avg	0.99 0.99	0.98 0.99	0.99 0.98 0.99							

Fig 11. Random Forest Binary Model and evaluation metrics

Deo	cision Tree	model w	ith bina	ary labe	led data							
[29]	<pre>from sklearn.tree import DecisionTreeClassifier classifier_dt = DecisionTreeClassifier(criterion = 'entropy', random_state = 42) classifier_dt.fit(X_train, y_train)</pre>											
	DecisionTreeC	lassifier(cr:	iterion='	entropy',	random_state=	42)						
[30]	<pre># Classification report y_pred = classifier_dt.predict(X_test) print(classification_report(y_test, y_pred, target_names=['Benign', 'Malware']))</pre>											
		precision	recall	f1-score	support							
	Benign Malware	0.96 0.99			1004 2919							
	accuracy macro avg weighted avg	0.98 0.98	0.98 0.98	0.98 0.98 0.98	3923 3923 3923							

Fig 12. Decision Tree Binary Model and evaluation metrics

Support Vector Machine (SVM) model with binary labelled data

```
#Import svm model
     from sklearn import svm
    svm = svm.SVC(kernel='rbf',random_state=1,C=1,gamma='auto')
    svm.fit(X_train, y_train)
    SVC(C=1, gamma='auto', random_state=1)
[33] # Classification report
    y pred = svm.predict(X test)
    print(classification_report(y_test, y_pred, target_names=['Benign', 'Malware']))
                  precision recall f1-score support
                    0.92 0.01 0.02
0.75 1.00 0.85
                                                     1004
          Benign
                                          0.85
         Malware
                                                     2919
       macro avg 0.83 0.51 0.44 3923
ighted avg 0.79 0.75 0.64
    weighted avg
```

Fig 13. SVM Binary Model and evaluation metrics

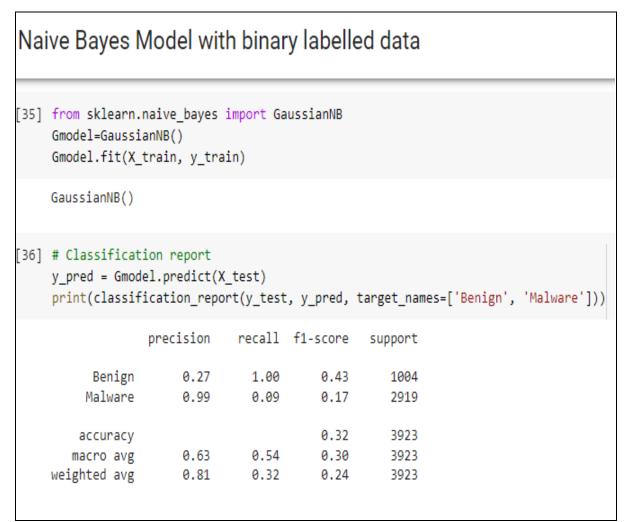


Fig 14. Naïve Bayes Binary Model and evaluation metrics

## K-Nearest Neighbour model with binary labelled data

```
[38] from sklearn.neighbors import KNeighborsClassifier
knn = KNeighborsClassifier(n_neighbors=1)
knn.fit(X_train,y_train)
KNeighborsClassifier(n_neighbors=1)
```

```
[39] # Classification report
y_pred = knn.predict(X_test)
print(classification_report(y_test, y_pred, target_names=['Benign', 'Malware']))
precision recall f1-score support
Benign 0.93 0.94 0.94 1004
Malware 0.98 0.98 0.98 2919
```

0.97

0.96

0.97

3923

3923

3923

Fig 15. KNN Binary Model and evaluation metrics

0.96

0.97

### Step 7: Model building for Multi classification

0.95

0.97

accuracy

macro avg

weighted avg

```
Model Building for Multi Classification
[41] # class distribution for original data
     print(data.groupby('Subsystem').size())
     Subsystem
    1
            438
          15451
     2
     3
          3711
     9
              4
    10
              4
    16
              3
    dtype: int64
[42] # sample distribution print in markdown table format
     #label = [Adware, Virus, Spyware, Trojan, Worm, Ransomware]
     label = 'Subsystem'
     lblTypes = set(data[label])
     for lbl in lblTypes:
         print('| {} | {} | '.format(lbl, len(data[data[label] == lbl].index)))
```

Fig 16. Labels for malware classification

[42]	1   438     2   15451     3   3711     9   4     10   4     16   3
[43]	<pre># Splitting the data into training and testing data subset into 80% and 20% respectively X_train, X_test, y_train, y_test = train_test_split(data_usage, data['Subsystem'], test_size=0.2, random_state=0)</pre>
[44]	<pre>print(f'Number of used features is {X_train.shape}') print(f'Number of used features is {X_test.shape}') print(f'Number of used features is {y_train.shape}') print(f'Number of used features is {y_test.shape}')</pre>
	Number of used features is (15688, 75) Number of used features is (3923, 75) Number of used features is (15688,) Number of used features is (3923,)

Fig 17. Dataset split for Multi classifier models

Rar	ndom forest	model	with mu	ti class	s data						
[45]	<pre>rfc = RandomFore rfc.fit(X_train,</pre>		oob_score = max_depth =	True,	0, random_sta†	e=0,					
	/usr/local/lib/p "X does not ha RandomForestClas	ave valid	feature nam	nes, but"		: UserWarning: X does not have valid feature names, but RandomForestClassion					
[46]	<pre># Classification report #"1": "Adware", "2": "Virus", "3": "Spyware", "9": "Trojan", "10": "Worm", "16" : "Ransomware"}, inplace=True)</pre>										
	<pre>y_pred = rfc.pre print(classifica</pre>	· -		y_pred, t	target_names=	['Adware', 'Virus', 'Spyware', 'Trojan', 'Worm', 'Ransomware']))					
	pr	recision	recall <del>(</del>	1-score	support						
	Adware Virus Spyware Thoian	0.99 1.00 1.00 1.00	0.98 1.00 1.00 1.00	0.98 1.00 1.00 1.00	90 3044 786						
	Trojan Worm Ransomware	1.00 1.00 1.00	1.00 1.00 1.00	1.00 1.00 1.00	1 1 1						

Fig 18. Multi classifier model using Random Forest and evlauation metrics

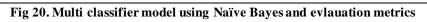
c	rom sklearn. lassifier_dt lassifier_dt	= DecisionT	reeClassifi			opy', randon	1_state = 4	42)			
D	ecisionTreeC	lassifier(cr	iterion='er	ntropy', r	andom_state	=42)					
y.	Classificat _pred = clas rint(classif	sifier_dt.pr		y_pred, t	arget_names	=['Adware',	'Virus',	'Spyware',	'Trojan',	'Worm',	'Ransomware']
	Adware Virus	1.00	1.00 1.00	1.00	90 3044						
	Spyware Trojan Worm				786 1 1						
	Ransomware	1.00	1.00	1.00	1						
	accuracy	1.00	1.00	1.00	3923						
	macro avg eighted avg	1.00 1.00	1.00 1.00	1.00 1.00	3923 3923						



SV	M model f	or multi c	lass da	ta							
[51]	#Import svm from sklearn svm = svm.SV svm.fit(X_tr	import svm C(kernel='rb	-	state=1,C	=1,gamma='au	ito')					
	SVC(C=1, gam	ma='auto', r	andom_stat	e=1)							
[52]	<pre># Classifica y_pred = svm print(classi</pre>	.predict(X_t		, y_pred,	target_name	es=['Adware',	'Virus', '	Spyware',	'Trojan',	'Worm',	'Ransomware']))
		precision	recall	f1-score	support						
	Adware	1.00	0.09	0.16	90						
	Virus	0.78	1.00	0.88	3044						
	Spyware	1.00	0.03	0.05	786						
	Trojan	0.00	0.00	0.00	1						
	Worm	0.00	0.00	0.00	1						
	Ransomware	0.00	0.00	0.00	1						
	accuracy			0.78	3923						
	macro avg	0.46	0.19	0.18	3923						
	weighted avg	0.83	0.78	0.70	3923						

Fig 19. Multi classifier model using SVM and evlauation metrics

N 1 1											
Naive	Bayes r	model wit	th multi	class d	ata						
	,										
[54] fro	om sklearn.	.naive_bayes	import Ga	ussianNB							
Gmo	del=Gaussi	lanNB()									
Gmo	del.fit(X_	_train, y_tra	ain)								
6											
Gau	ussianNB()										
		ion report									
	pred = Gmod	del.predict(X	(test)								
pri	nt(classif	fication_repo		, y_pred,	target_names	=['Adware',	'Virus',	'Spyware',	'Trojan', '	'Worm',	'Ransomware'])
pri	nt(classif		ort(y_test			=['Adware',	'Virus',	'Spyware',	'Trojan', '	'Worm',	'Ransomware'])
pri	nt(classif	fication_repo	ort(y_test	, y_pred, f1-score	target_names support	=['Adware',	'Virus',	'Spyware',	'Trojan', '	'Worm',	'Ransomware'])
pri	nt(classif Adware		ort(y_test			=['Adware',	'Virus',	'Spyware',	'Trojan', '	'Worm',	'Ransomware'])
pri		precision	recall 0.01	f1-score	support	=['Adware',	'Virus',	'Spyware',	'Trojan', '	'Worm',	'Ransomware'])
pri	Adware	precision 0.17 1.00	recall 0.01	f1-score 0.02	support 90	=['Adware',	'Virus',	'Spyware',	'Trojan', '	'Worm',	'Ransomware'])
pri	Adware Virus	precision 0.17 1.00 0.49	recall 0.01 0.08	f1-score 0.02 0.15	support 90 3044	=['Adware',	'Virus',	'Spyware',	'Trojan', '	'Worm',	'Ransomware'])
pri	Adware Virus Spyware	precision 0.17 1.00 0.49	recall 0.01 0.08 0.69	f1-score 0.02 0.15 0.58	support 90 3044 786	=['Adware',	'Virus',	'Spyware',	'Trojan', '	'Worm',	'Ransomware'])
	Adware Virus Spyware Trojan	precision 0.17 1.00 0.49 0.00	0.01 0.01 0.08 0.69 0.00	f1-score 0.02 0.15 0.58 0.00	support 90 3044 786 1	=['Adware',	'Virus',	'Spyware',	'Trojan', '	'Worm',	'Ransomware'])
	Adware Virus Spyware Trojan Worm ansomware	precision 0.17 1.00 0.49 0.00 0.00	recall 0.01 0.08 0.69 0.00 1.00	f1-score 0.02 0.15 0.58 0.00 0.00 0.00	support 90 3044 786 1 1 1	=['Adware',	'Virus',	'Spyware',	'Trojan', '	'Worm',	'Ransomware'])
R	Adware Virus Spyware Trojan Worm ansomware accuracy	precision 0.17 1.00 0.49 0.00 0.00 0.00	recall 0.01 0.08 0.69 0.00 1.00 1.00	f1-score 0.02 0.15 0.58 0.00 0.00 0.00 0.20	support 90 3044 786 1 1 1 3923	=['Adware',	'Virus',	'Spyware',	'Trojan', '	'Worm',	'Ransomware'])
R	Adware Virus Spyware Trojan Worm ansomware	precision 0.17 1.00 0.49 0.00 0.00	recall 0.01 0.08 0.69 0.00 1.00 1.00	f1-score 0.02 0.15 0.58 0.00 0.00 0.00	support 90 3044 786 1 1 1	=['Adware',	'Virus',	'Spyware',	'Trojan', '	'Worm',	'Ransomware'])



K-nearest Neighbour model with multi class data											
[57]	from sklearn.n knn = KNeighbo knn.fit(X_trai	rsClassifie			sifier						
	KNeighborsClas	sifier(n_ne	ighbors=1)								
	<pre># Classificati y_pred = knn.p print(classifi</pre>	redict(X_te		y_pred, t	target_names:	['Adware',	'Virus',	'Spyware',	'Trojan',	'Worm',	'Ransomware']))
		precision	recall f	f1-score	support						
	Adware Virus Spyware Trojan Worm	0.78	0.92 0.95 0.75 0.00 1.00	0.93 0.94 0.77 0.00 1.00	90 3044 786 1 1						
	Ransomware	1.00	1.00	1.00	1						
	accuracy macro avg weighted avg	0.77 0.90	0.77 0.91	0.91 0.77 0.91	3923 3923 3923						

Fig 21. Multi classifier model using KNN and evlauation metrics