

# Differential IP Clustering for Vulnerability Detection using applied Machine Learning



UNIVERSITY  

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of  
ABERTAY DUNDEE

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This thesis is dedicated to  
no one  
for no special reason

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# Abstract

plenty of waffle, plenty of waffle, plenty of waffle, plenty of waffle, plenty  
of waffle, plenty of waffle, plenty of waffle, plenty of waffle.

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# Chapter 1

## Introduction

This section will give a brief background and history of Machine Learning before going in to depth with the definitions and types of algorithms used in modern Machine Learning systems. An analyses of the impact Machine Learning has in the field of computer security and how it can be applied directly to penetration testing and red teaming. Research aims and objectives will be provided as well as a statement to the structure of the rest entire thesis.

### 1.1 Background

The term machine learning was first defined by Arthur Samuel in the year 1959 as a "Field of study that gives computers the ability to learn without being explicitly programmed" and later as "a field of study that concentrates on induction algorithms and on other algorithms that can be said to 'learn'", Kohavi & Provost (1998). Machine learning has largely evolved from several subfields of artificial intelligence, specifically, computational learning and pattern recognition but now it stands on its own with its subfield Deep Learning being at the forefront of technology. Machine learning consists of the studying and construction of algorithms that can learn from and make predictions of data, Simon (2013). The early information available for machine learning was almost entirely theoretical due to the lack of processing power at the time. One of the early pioneers was Valiant (1984) whom developed a PAC learning framework and established the theory of a learnable algorithm, Munoz (2014). Modern machine learning algorithms use many calculations from statistics, information theory, theory of algorithms, probability and functional analyses, Munoz (2014).

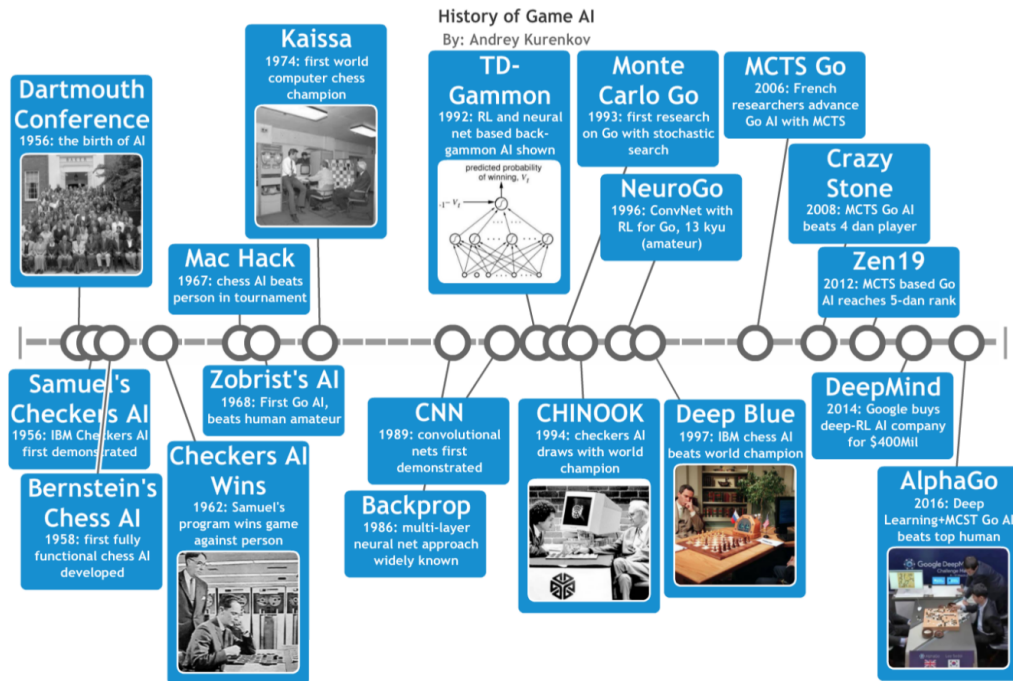


Figure 1.1: Timeline of artificial intelligence for games, Kurenkov (2016)

## 1.2 Current State of Machine Learning

In the recent years, machine learning accuracy and efficiency has increased dramatically which has caused the field to rapidly gain in popularity, especially in deep learning. One of the major reasons for this is due to the improvement in graphics processing units which in some computational tasks can outperform central processing units by an order of magnitude or more, Mattsson (2016). The rapidly growing field of deep learning could be made apparent by Googles recent self-driving car and LeNet image recognition system, however, examples of machine learning can be found in just about every field with a classification or regression problem. Some of which include: natural language processing using morphological analyses for linguistics (Black & Zernik (1994)), document classification and spam detection, better image recognition than humans (Karpathy (2011)) and beating a professional player at the complex Chinese game Go (DeepMind (2016)). Figure 1.1 illustrates the advancement of machine learning using the context of game artificial intelligence players.

The majority of projects over the past few years have focused on Deep Learning due to its performance and versatility with complex problems but also because of the media coverage with large scale projects by international corporations such as Google and Microsoft. The idea of deep learning has been around since the 1980s where a

Japanese scientist, Kunihiro Fukushima, proposed a hierarchical, multi-layered artificial neural network named Neocognitron, designed for handwritten character recognition. Neocognitron was recognised as the inspiration for convolutional neural networks, the most commonly used deep learning algorithm, LeCun et al. (2015). Deep learning attempts to model complex abstractions in data by using a multiple-level architecture most commonly comprising of artificial neural networks and non-linear transformations in its algorithms, Mattsson (2016). As such, the concepts of deep learning are extensions of regular machine learning algorithms.

In the past year a new topic has become of consider interest in the machine learning field, Automated Machine Learning, which can be considered to cover the tasks of algorithm selection, hyperparameter tuning, iterative modelling and model assessment, Mayo (2017). By the end of 2016 the python Auto-sklearn library was created based off of the scikit-learn library which encompasses these tasks, created by a team from the University of Freiburg it won the KDnuggets AutoML challenge, Matthias Feurer (2016). Using these examples, its possible to hypothesize that the future of machine learning will include automatic deep learning, however, these individual fields of machine learning are still in their infancy and far from being used together.

The computer security industry due to its nature, has many examples of machine learning implementations such as intrusion detection systems using machine learning or deep learning for anomaly detection. One of which has been commercialised under the company name Deep Instinct and advertises zero-day detection using deep learning. However, the majority are very similar and are all blue team based security solutions. Researching into Red team or penetration testing tools using machine learning resulted in a disappointing lack of tools or ideas considering the vast amount available for Blue team.

The sole documented research found for red team was for an automatic penetration testing project named Auto Red Team (ART) framework by Lu, Song of Iowa State University in 2008 which used decision trees and hard programmed exploits. This meant the entire hard programmed exploit section had to be reconstructed for each use case, this would not be ideal but also extremely time consuming. Further analyses of the ART framework can be found in the literature review of "Auto Red Team: a network attack automation framework based on decision tree". There have also been tools and libraries created to test the security of software which use machine learning

models such as Deep Pwning. Deep Pwning is an open source metasploit plugin which allows the tricking of machine learning models. This field of research was named Adversarial Machine learning and the first paper of which was respectfully named "Adversarial Machine Learning" and published by ACM in 2011.

There is an extensive amount branches to machine learning and unfortunately, too many to cover in this thesis due to time constraints. Therefore, this introduction will only detail the most popular types of machines learning algorithms.

### **1.3 Significance of study**

The increased demand for penetration testers justifies the need for more effective and advanced tools to conduct their security assessments. The goal of a penetration test being to find vulnerabilities in a system using techniques similar to that of a malicious hacker. This means malicious hackers will continue to use the most advanced methods to gain access to critical systems and thus security teams must also continue to advance their toolset to be as effective and efficient as possible. Penetration tests can last a time scale of anything between one day to several months and more advanced machine learning tools would allow for a more efficient use of this time. The large variety of machine learning models used in the blue team results in a large variety of models required to test them using adversarial machine learning as well as advanced machine learning tools, specifically for the red team in order to bring each team to the same level. Having both red team and blue team on the same level is beneficial for the industry as a whole, providing competition between both sides and to continue to strive for improvement.

This project aims to help correct this unbalance by designing and developing a proof of concept red team tool using machine learning techniques.

The application must be able to be used as an aid to a security professional during a security assessment or capture the flag hacking events in order to be classed as a red teaming tool.

This application must also be scalable and versatile to be used in varying sized network environments.

The tool will be critically analysed and recommendations of related future work will be provided.

## 1.4 Types of Machine learning

As mentioned above there are many different machine learning algorithms available. Each algorithm can be classed based on the problem, required output and several factors of the data set, such as whether it includes labels and the amount of values it includes. The two primary problems machine learning algorithms provide solutions to, can be put in to two categories which are not mutually exclusive and can be combined in certain use cases. These are classification and regression problems. The following describes and states the differences of these types.

### 1.4.1 Classification

With the growth of big data, unstructured data is more prevalent than its structured counterpart. This is because, "while the amount of structured data has grown fast, the amount of unstructured data has grown much faster" Simon (2013). This creates the need for ever more efficient data analytics and is a typical classification problem for machine learning. There are many types of classification, simple types such as a linear classifier and then more complex types such as multiclass and structured classifiers. Classification is largely used in data mining and statistical analyses for these purposes. In short, classification is used when you require an input variable to be identified as part of a group or label, resulting in the full dataset being categorised. Classification can only take a finite set of values such as picking from 1 of N values. In classification, each incorrect answer is equally incorrect, compared to regression where incorrect answer can be varying levels of incorrect.

### 1.4.2 Regression

Regression problems are for when prediction of real continuous values is required, such as predicting stock market values and detecting the age of a person from a picture, Rossant (2014). Regression analyses involves predicting and estimating a response based on previous data and input variables. As mentioned above regression answers can have a varying level of inaccuracy as appose to binary correct or false predictions. This is due to regression using continuous values. Figure 1.2 provides a basic illustration example of these two problems.

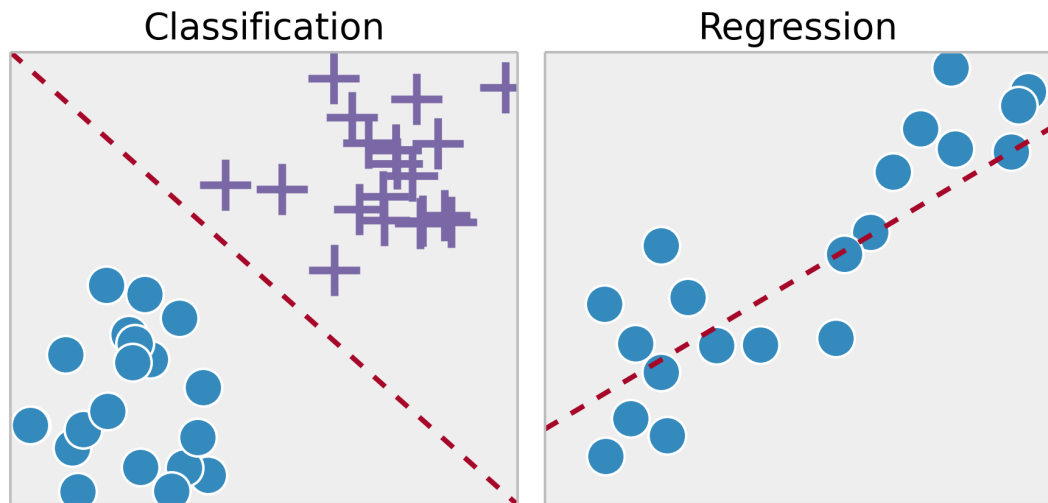


Figure 1.2: Illustration of machine learning types, Rossant (2014)

## 1.5 Types of Learning algorithms

The most commonly found learning methods for algorithms are supervised and unsupervised followed by semi supervised and reinforcement learning which are also fairly common. These are explained briefly as follows.

### 1.5.1 Supervised

In a supervised training model, the entire data-set used for training is pre-labelled so that the algorithm can use pattern analyses to predict given values after training. A use case scenario for supervised learning includes the above stock market example where current values with labels are given and the model is used to predict future labels.

### 1.5.2 Semisupervised

Due to the expensive nature of labelled data in most cases, semisupervised learning uses a split of labelled and unlabelled data to train the model. This is most often split unevenly with the large majority of training data being unlabelled. This model is often used in cases where the cost of labelled data-sets is simply too expensive.

### 1.5.3 Unsupervised

The opposite of supervised learning, where the training data-set has no labels and the model must attempt to determine the correct answer itself. This is often used as a method to determine a structure in the data given. These models can identify segments of similar attributes such as clustering. Unsupervised learning is the primary method used for this thesis methodology.

### 1.5.4 Reinforcement

Reinforcement learning works similarly to heuristic algorithms in the sense that every possibility is attempted and assigned a score in which the iteration with the highest score is used as the model output. Whilst training this model the algorithm uses the highest scores iteration to modify its calculations for greater accuracy and efficiency. This model is often used in robotics as well as game design for path navigation calculation and computer player AI (artificial intelligence). The following is an example use case: A chess player AI calculates each possible move it can make using a reinforcement model for each of its turns. The model assigns a score to each move it could possibly make at that point in time and weights them based on pieces acquired, future strategy prospects and defence risk. Similar models have been used for AIs mentioned in Figure 1.1.

## 1.6 Common Machine Learning Algorithms

As mentioned above, due to the amount of machine learning algorithms available and project time constraints, only a few of the most common algorithms will be described below.

### 1.6.1 Linear Regression

Linear regression predicts real values based off continuous data of two variables. It does this by using a best fit or regression line over the existing data extending in predictions. If the data does not indicate any positive or negative trends then using linear regression will likely not be a very useful model. The trend or direction of data can be calculated using correlation coefficient as follows, where  $(x_2, y_1), (x_n, y_n)$  is the observed data.

$$r = \frac{1}{n-1} \sum \left( \frac{x - \bar{x}}{s_x} \right) \left( \frac{y - \bar{y}}{s_y} \right)$$

A value that is close to 1 would indicate positive correlation where -1 would indicate negative correlation. A normalised covariance calculation may also be used instead.

### 1.6.2 K-Nearest Neighbours

K-nearest neighbours (or KNN) is a widely used computationally expensive supervised learning model for data classification but can also be used for regression problems. The value k refers to the distance is which to weigh the number of class nodes inside. Several distance functions can be used such as Euclidean, Manhattan and Minkowski with the most common being Euclidean. Euclidian distance is the straight-line distance between the two nodes. On a two-dimensional plane it is measured using the following formula where the coordinates are  $\mathbf{p} = (p_1, p_2)$  and  $\mathbf{q} = (q_1, q_2)$ .

$$d(\mathbf{p}, \mathbf{q}) = \sqrt{(q_1 - p_1)^2 + (q_2 - p_2)^2}$$

For e.g. if  $k = 1$  then the current node will be assigned to the same class as its nearest node, where as if  $k = 5$  then the class with the largest number of nodes in a distance of 5 will be selected to represent the current node. KNN is not resistant to bias in data-sets and thus data must be normalised before inputted into the model. KNN models are not to be confused with K-means clustering models as they share a very loose relationship being that KNN can be used to add data into pre-existing K-means clusters known as a nearest centroid classifier.

### 1.6.3 K-means Clustering

K-means Clustering is an unsupervised model in which the number of clusters is specified in advance. K being the number of clusters the model will output for the given data. The value for K can be assigned manually or the optimal value can be calculated by using either the elbow method or the gap statistic, the latter of which is used and explained in the methodology section of this thesis with a practical use of K-means. The K-means algorithm itself is also known as Lloyd's algorithm and uses iterative refinement to determine the best clustering.

### 1.6.4 Dimensionality Reduction

Dimensionality reduction algorithms are exactly what the name suggests. With the dramatic increase in variable types and raw data size being captured leading to the invention of the term Big Data, datasets have become very large. These large datasets have become the main bottleneck for machine learning performance,



especially in computationally expensive models such as KNNs and K-means. There needs to be a way to identify the significance of each variable in the data-set in order to give weights to their values for use in calculations. Principle Component Analyses (PCA) can be used for this purpose as well as for variance maximization. The PCA algorithm attempts to detect correlation between each of these variable types and uses this information to detect the vector directions of maximum variance in high-dimensional data. The algorithm then projects these maximum variance vectors into a smaller dimension whilst attempting to keep the most amount of information and greatest variance as possible. If the measurement scales of the dataset variables are not equal, then the data should be normalised prior to PCA due to the variance maximization functions of the algorithm.

### **1.6.5 Decision Trees**

Decision trees are a classification model that uses supervised learning to calculate either categorical or continuous dependent variables. Decision trees can also be designed manually as was traditionally done. As the name suggests, the model works by splitting the data at each branch of the tree by asking questions and making decisions on each layer. The concept of decision trees is simple however they can be combined together, or include other machine learning models within its layers greatly increasing the complexity of this model.

### **1.6.6 Random Forests**

Random decision forest is one of these complications where the model contains several decision trees built during the training stage and can also be used with unsupervised learning. These random decision forests can be used for classification or regression by simply using the mode of the results from the forest or the mean in the case of a prediction problem. The creation of this model differs from regular decision trees as the training algorithm applies bootstrap aggregating, commonly referred to as bagging, followed by feature bagging resulting in a random decision forest. Bagging increases the stability and accuracy of decision trees by reducing variance and avoiding the overfitting of data. It does this by generating additional sets data based on the given data-set which results in a larger data-set overall reducing the variance but increasing the predictivity of the model. Bagging can also be used on several other machine learning algorithms. Feature bagging also known as the random subspace method, selects random samples from the entire dataset in order to reduce the

correlation between each tree in the forest. This is done to produce varied models for each tree as appose to them being very similar if they were all trained on the entire data-set. Random forests can be complicated once more by adding a neural network within its layers, or even a deep learning model such as recurrent neural networks.

### **1.6.7 Artificial Neural Networks**

Artificial neural networks (or ANNs) are commonly used as the bases for most deep learning algorithms such as convolutional and recurrent networks and are loosely based on the way a human brain functions. They can be used with several types of learning methods. Neural networks consist of interconnected neurons on a layered scheme in which every neuron has a function and its output can be seen by its connected neurons to then use it in their own functions. Each neural connection has an assigned weight for its output which is calculated during the training stage of the model. The layers consist of three types, an input, an output and hidden layers sandwiched between the two. The number of hidden layers in a neural network is what defines whether it is a deep learning model or not. Models with more than one hidden layer are referred to as deep learning models. How the layers are connected and the way data travels between the layers defines the type of neural network. For e.g. a recurrent neural network allows for sequences to be used for input and output unlike convolutional which can only use fixed values for these. The Google subsidiary DeepMind uses recurrent neural networks in its algorithms to create its AlphaGo AI mentioned previously.

# Chapter 2

## Literature Review

This chapter will include a review of related works in machine learning and network security.

Machine learning has been used in the information security industry ever since Debar et al. (1992) created the first basic network traffic analyser to separate attacks from regular packet traffic using an artificial neural network. Then similarity measurements were used to find anomalies in inputted Unix commands sequences by Lane & Brodley (1997). This lead to the creation of a cluster classifier by Labib & Vemuri (2002) which analysed real-time traffic in a network and plotted it into a GUI graphing interface very similar functionally wise to the application created in this thesis. Like other blue teaming solutions that came before it, Labibs solution allowed for anomalies such as malicious traffic to be detected automatically. Ever since the afore mentioned examples, blue team machine learning solutions were no longer few and far between. The industry has been continually proactive, developing new theories and solutions at an increasing rate. One of such recent developments include the article, Shallow and Deep network intrusion detection systems: A Taxonomy and Survey, Hodo et al. (2017). This article tackles the issue of creating an efficient intrusion detection system to handle large scale data with changing patterns in real time. Another such example is Niyaz et al. (2016), where the authors discuss the optimal approach of using deep learning techniques for intrusion detection systems. All these developments, although great for the industry as a whole have been unbalanced with the majority of research lying strictly within the blue team. This has created a gap in research for the red team. Red teaming in its nature stresses the importance of balance between the two sides. The balance between these teams is important due to the simple fact that malicious individuals will not stop improving and developing new methods. Without the balance of research allowing red team to keep up with

those of malicious intent, red teaming will risk falling too far behind and becoming obsolete. The following paragraphs will now focus on red teaming.

Lu (2008), developed the first machine learning red teaming software using decision tree algorithms. Lu Song named this software Auto Red Team and was successful in creating a semi-automated penetration testing platform. The first step of the framework involved capturing the traffic between the attacker and the victim machines whilst the attacking machine would launch every single exploit supported by the application. For this the framework used 42 exploits from the Metasploit command line edition. All 42 exploits which are hard programmed into the framework must first be executed against the victim manually at this initial learning stage. Each exploit is manually programmed with hard paths due to the fact that, in the year 2008, the Metasploit command line available although having much more than 42 exploits, did not include a query system to find the exploits efficiently. The entire network traffic for this first stage must be captured.

This captured traffic is then audited by the decision tree algorithm which will return a new composed attack strategy and forwarded to a new and improved second attacking unit. Before the strategy is sent to the new unit, it must be parsed and converted to Perl script as the composer outputs the strategies in a human readable format. This new attack unit would then execute this strategy automatically whilst capturing the traffic, however, upon completion it will ask the user to declare whether the exploit was successful or not in order to proceed. At this point the users decision and twelve metrics of data from each captured packet are used as training data from the decision tree, improving its efficiency and accuracy. If the users decision was that the exploit had failed, the framework will repeat the entire process until the user declares that the exploit was successful. The ART framework therefore uses a supervised learning version of a decision tree model. Among Lu Songs future work recommendations is that the Perl converter be replaced with a C++ compiler which could translate the strategy composer output into executable code in real time. This would greatly reduce much of the processing time and convert the application model into a just in time approach. Further, by simply using a recent version of Metasploit the framework would be able to use any exploit from a given library without the need to hard program the paths as variables, greatly increasing the efficiency in the initial stages and overall versatility of framework.

This concludes the literature review chapter and the following chapter will review the thesis methodology.

# Chapter 3

## Methodology

This chapter contains an in-depth analysis of the design and implementation stages carried out during this project. Initially, the design and goals of the application will be enumerated. Subsequently an analysis of the applications infrastructure will be presented followed by an in-depth detail of the modules and submodules within the applications programming. A test case scenario will be defined and executed to provide a proof of concept. The efficiency and accuracy among other factors based on the test scenario will be analysed during the discussion chapter of this thesis.

### 3.1 Design

The application designs primary goal was to be able to detect vulnerable machines on a large-scale network infrastructure regardless of topology or host types by using machine learning techniques and automated tool outputs. However, there are several requirements the application must adhere to for it to be a viable tool during a security assessment. This application was created strictly on a proof of concept bases.

The application was designed with the following requirements in mind:

**Text progress output with multiple verbosity settings** allowing for an experienced tester to understand what the application is doing at any point in time during execution. This is critical as tools used during an assessment on live networks must not hinder or damage the network or its hosts in any way as to disrupt an organisations business.

**Several input type parameters** for which the tester can utilise based on the current information known about the network. Such as, only using one type of scan file and manually selecting the clustering model.

**Several output options including visually in the form of graphs** and to a dot type file to be used with other industry applications and reports.

**Manual overriding of variables via parameters** in order to allow for the application to be scripted and modified by the tester. This will increase the efficiency of using the tool and provide advanced customisation of the algorithms within the application.

**Highly versatile with working conditions and configurability.** The programming of the application to be highly documented allowing a tester to fix and modify the application code to suit the operations needs. By using a primarily interpreted language as oppose to compiled one would allow for this, as well as making the application portable without extra code. For these purposes, the Python programming language was chosen. With the majority of modern tools and scripts used by penetration testers haven been written in Python due to its versatility, reliability and portability, it further enforces this choice.

### 3.1.1 Application Brief

The application requires several parameters to run and has three different global modes; manual, assisted and automatic. These modes can either be run with Nmap, Nessus or both inputs with the majority of the use case scenarios requiring both. The application will then parse these inputs into labels and features, process the data in several ways and cluster the information based on feature similarities (explained further in section 3.2.4). The text output is then displayed which includes the full details of each cluster within the clustering and several statistics such as adjusted silhouette values (more information found in section 3.2.5 and 3.2.6). When using Dual input mode (both Nessus and Nmap files) the application will combine the data from each and subtract the large similarity clusters. By doing this, the application will determine the most unique hosts within the topology and display them in a new clustering. These most unique hosts, based on probability, will be the most vulnerable on the network and should be prioritised by a tester during the manual security assessment, because it is highly unlikely that the large clusters removed

would contain addresses which are solely vulnerable to the same exploit. This is due to the clustering model prioritising the vulnerabilities that each scanner detects, then appending them to the pre-existing host set, thus rendering that host more unique with a greater feature difference than the other hosts without this vulnerability. The application then renders a graphing interface to display the information as such in described in 3.8.

An example output of the application when using **dual input automatic mode** can be found in text form using maximum verbose level at Appendix B.1.1 and a Figure of the graphing interface GUI at Appendix B.1.2. The data-set used for these examples were Nessus and Nmap scan XML outputs generated from a fictional network. Due to the sensitive nature of the data included within these scans such as SSH keys and vulnerability codes, there are no publicly available data-sets.

The difference between modes and parameters will be explained further in the next section, the *infrastructure analysis*.

## 3.2 Infrastructure

Figure 3.1 shows the applications primary class infrastructure when in automatic mode. The infrastructure diagram has been created using standard flow diagram symbols to provide understanding of the process types. The application has been programmed for python version 2.7 interpreters and therefore will not have complete functionality without modification for python 3.0 and above. Due to time constraints placed upon this thesis the library NMAP-Cluster, Blackhat USA (2016), was used to conserve time.

In order to execute this application, it is important to have the correct library dependences. This is done via the python package manager PIP and a requirements file, found in Appendix A.1, by executing the command:

---

```
1 Pip install -r requirements.txt
```

---

The following sections include detailed descriptions of the processes symbolised within the infrastructure Figure 3.1. Beginning from the start circle and ending at the display modules.



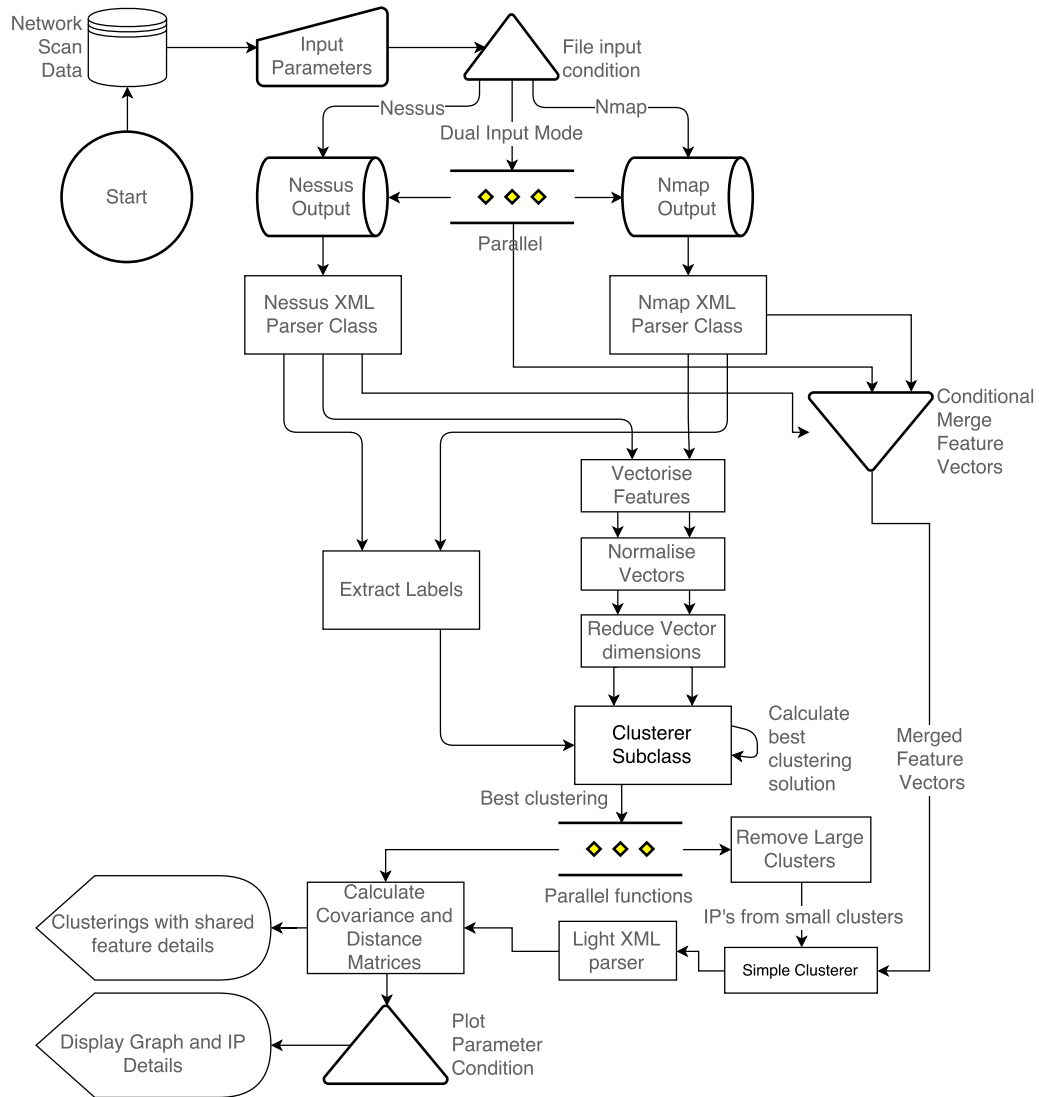


Figure 3.1: Application infrastructure data flow diagram

### 3.2.1 Usage and Parameters

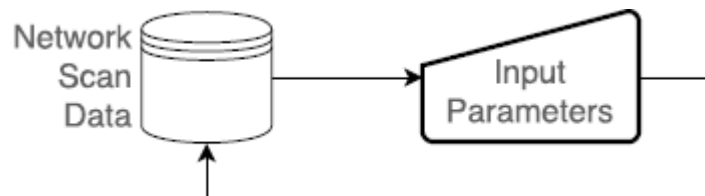


Figure 3.2: Network scan data and input parameter modules from infrastructure diagram at Figure 3.1

The modules shown in Figure 3.2 are those used to store the network scan data and process the applications input parameters. The Network Scan Data module refers

to the XML output of an Nmap scan, Nessus proprietary export file or both. These must be in their respectable formats in order for the parser to recognise them. The files must also be referenced in their correct positional arguments when executing the application such as mentioned in the application usage in Appendix A.2.

The Parameters manual input module on Figure 3.2 refers to the parameters that the application requires to select the correct run configuration. This is required because the application has no execution graphical user interface (GUI), the lack of which was decided for several reasons. Such as allowing for scripting, verbose output and terminal pipe operation commands. This type of interface is generally preferred by professionals due to the speed and reliability it provides over a standard GUI. The graphing stage of the application does however, provide a GUI to allow for manipulation of the graphs in multiple ways. The application usage found in Appendix A.2 includes a full description of the possible parameters. The parameters are passed into the data processing section of the application explained in the next session. The three possible run modes are explained in a later section 3.6.

### **3.2.2 Data Processing**

The data processing modules from the infrastructure diagram have been highlighted in Figure 3.3 and will be described in the following paragraphs.

Depending on the files specified within the input parameters, the application will either feed the Nessus, Nmap or both XML files into the parser classes. These parser classes will take the data from each file and transform it into IP addresses and features which are then passed individually into vectorizers. Vectorization is required for the features to be understood by the clustering algorithm. Vectorization refers to the general process of turning a collection of text (in this case machine attributes) into numerical feature vectors as float values. It is important to note that data from each scanner file is kept separate until the final process. The vectorization class is short and concise due to it having only two purposes, to call the parsers and vectorise the returned results. More information on the vectorization of each file as well as the raw python class code can be found in Appendix A.4.

Once the data has been vectorised it must be normalised in order to avoid large value bias when using dimensionality reduction algorithms such as PCA. Data normalization scales the values to within the same range whilst keeping the data variance

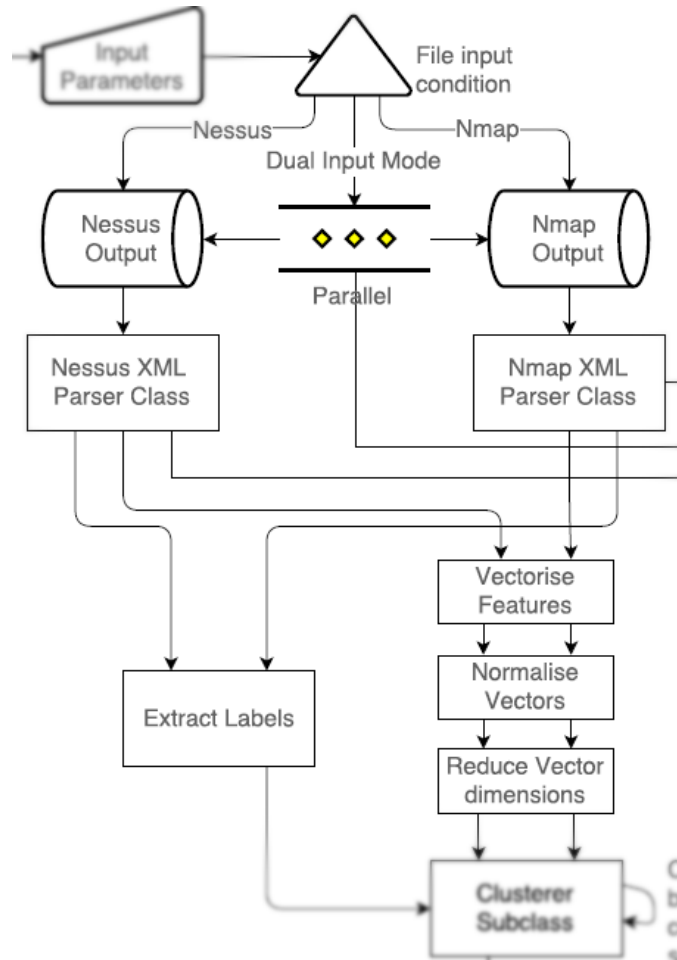


Figure 3.3: The data processing section from the infrastructure diagram at Figure 3.1 with the irrelevant modules blurred out

and eliminating the bias problem. This is programmed immediately after vectorisation in the applications initialisation class found in its raw code form at Appendix A.3.

The major difference between each scanner used is that the Nessus output values include vulnerabilities over Nmap which has superior information on the services and ports of the machine. By using both, the optimal range of information can be achieved, however, this introduces the problem of overfitting which PCA has countered. For more information on PCA refer to dimensionality reduction section 1.6.4 in the introduction. It is possible to greatly modify the output of the two scanners by either using scripts with Nmap or custom plugins for Nessus. Due to the modularity of the application, these modifications to the scanners will not affect the parsers and therefore can be used safely. PCA is also implemented within the initialisation class

which can be found in Appendix A.3. Once PCA is complete, the still separated feature vectors are sent to the clusterer subclass and possibly the conditional merger. The following section will explain the conditional merger of which is represented by symbol in Figure 3.4.

### 3.2.3 Small Combined Cluster

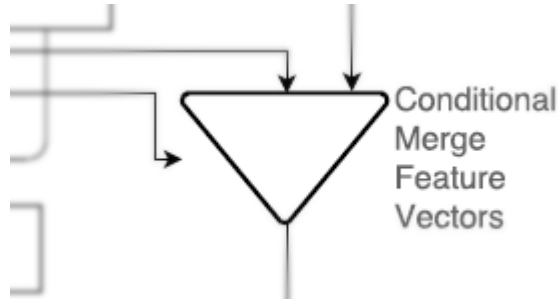


Figure 3.4: Conditional merge vectors process module from the infrastructure diagram at Figure 3.1

The conditional process of merging the feature vectors, referred to by the symbol in Figure 3.4, is executed at this stage, condition dependant on whether dual input mode has been selected when initialising the applications parameters. However, this path must then pause until the main cluster subclass (as represented by the symbol in Figure 3.6.) has completed in order to use its clustering outputs. Once this has occurred, each clustering will be duplicated and the large clusters with greater than three IP addresses will be removed from the clustering. This value can be changed based on the network size however the value of three was found to be optimal for networks of size 10 to 1000 from the algorithm tuning stage (this number can be configured for the users needs by modifying a single variable `maxaddresses` within the display class highlighted in Figure 3.5 below). The result of each is combined then re-clustered and the remaining IPs passed through a simpler file parser (compared to that of the ones previously used) to retrieve the information in an un-vectorised text format. This is done to display individual machine information for the end user within the graphing interface. The clustering of these small cluster IP addresses uses the gap statistic (or Elbow method if preferred by user) algorithm to define the numbers of clustered required as appose to user input as these IPs depend on the cluster subclass algorithm calculated clustering. The main cluster subclass mentioned is explained in

the following section 3.2.4. Gap statistic and Elbow method formulas can be found in the mathematical model at section 3.3.

```

142
143 def remove_large_clusters():
144
145     #originals needed for graph plotting t
146     nessusArray = clusterX
147     nmapArray = NclusterX
148     combined = np.array([])
149     maxaddresses = 3
150     #use biggest array by amount of cluste
151     #loop through largest cluster array, i
152     iterations=0
153     for index in range(len(nessusArray[:,
154         iplistnessus = nessusArray[(index-
155         #delete largest clusters, clus

```

Figure 3.5: Location of the variable which defines the number of maximum IP addresses a cluster is allowed to have for the vulnerability analysis process. The variable is defined within the display.py class remove large clusters function on line 149

### 3.2.4 Clustering Algorithm

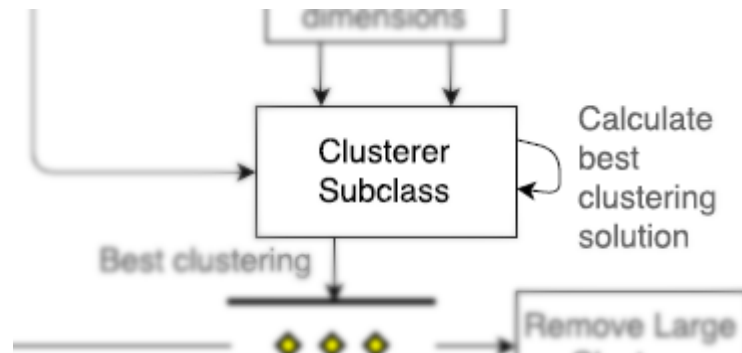


Figure 3.6: Representation of clusterer subclass from infrastructure diagram Figure 3.1 spotlighted

The Clusterer subclass represented in the infrastructure diagram by the module in Figure 3.6, is where the majority of the calculations are done within the application. This includes the three different modes; manual, assisted and automatic. The raw python code for this module can be found in Appendix A.5. Each mode can be used with a variety of clustering algorithms such as K-means, DBscan and agglomerative

clustering. When using dual input mode these functions will execute twice Nessus starting first. A brief summary of each mode is given in the following paragraphs whilst an in-depth algorithm review can be found in section 3.3:

**Manual:** The user supplies all required information to do the clustering. This includes the clustering algorithm and its hyper parameters. If no cluster count  $K$  is provided and K-means was selected, the gap statistic method will be used to calculate the optimal cluster count. An implementation of the elbow method is also included in the application and can be used instead if the gap statistic if the user desires.

**Assisted:** The user assists the algorithm by suggesting that some samples should or should not be clustered together. This is a much slower process than the other modes but provides the user with complete control over the clustering decisions. The user must specify the clustering algorithm to use from the selection mentioned above when using this mode.

**Automatic:** Multiple clustering strategies and parameters are used in an attempt to get the best clustering. Unfortunately, this function can only use the K-means clustering algorithm whilst running in dual input mode, due to limitations of the sklearn librarys centroid functions and the time constraints limiting this development as future work. This function goes through each possible iteration of each clustering algorithm up to a maximum cluster value equalling the number of labels in the given dataset. For e.g. in a dataset with ten IP addresses, the maximum allowed clusters is also then. There are multiple selections the user can decide on in terms of which clustering the application will select. This is calculated via a sorting function in the clustering subclass where the user can select between the average silhouette\* value, the minimum silhouette\* value, the average distance between IP addresses (Euclidian distance by default), the number of clusters in the clustering or the minimum number of common shared features a cluster has in its clustering. These options can also be joined together to make multiple filters. By default, the application uses both the minimum common features and number of overall clusters as filters allowing it to find the clustering with the least number of clusters requiring at least one shared feature. This option is changed near the end of the cluster subclass, changed by uncommenting one line per filter. Each clustering iteration in this loop will calculate the shared positive, negative features, the individual silhouette values and the mean distance to give an overall/per cluster score. Once the iterations are complete for each clustering algorithm the scores and details are used to select one of the clusterings based on the filters and then it is displayed. When using dual input mode, the vulnerability detection on the small clusters explained above in section 3.4 is continued,

due to it requiring these clusters as input. An example of the full text output and graphing interface for the applications automatic mode can be found at Appendix B.1.

\* The silhouette value is used to validate the consistency of clusters in a clustering by indicating whether there are too many or too few clusters. It provides a value of negative one to positive one indicating how well each IP fits in the cluster. A high value close to one indicates a good clustering by showing that the IP address is very similar to the other IP addresses in its cluster, whilst being very different to the IP addresses of the neighbouring clusters. Therefore, having a low or negative silhouette value indicates the wrong number of clusters used. The silhouette value has been calculated using the Euclidian distance (described in section 1.6.2) within the validation class.

### 3.2.5 Covariance and Distance Matrices

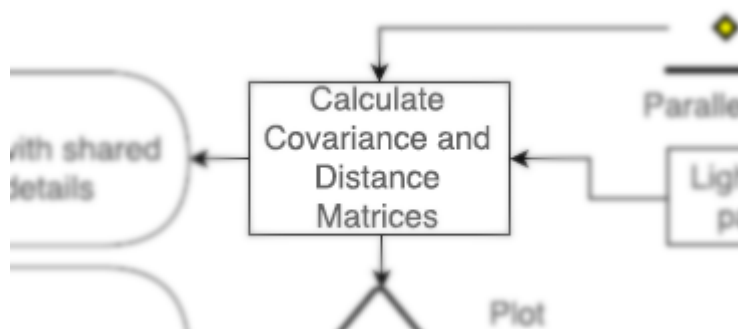


Figure 3.7: Representation of covariance and distance matrices function from infrastructure diagram Figure 3.1 spotlighted

Covariance and distance matrices are calculated just before displaying the graphing interface. The function symbol is shown in Figure 3.7 above. The output from these functions can be found at the end of the text output, an example is shown in Appendix B.1.1. This is shown for the purpose of giving a technical user greater knowledge of the data-set used which can be interpreted and reported to the client in a security assessment. Specifically, the covariance matrix shows how related the centroids for each clustering are to each other, in other words, their similarity. Covariance will highlight any linear relationships that might be apparent within the dataset with the values ranging from negative one to positive one, high values indicating a positive

linear relationship. The Distance matrix shows the literal Euclidian distance between each clustering centroid but by design that also correlates to showing a numericized value defining how different the clusters are from each other. These calculations are done for each clustering result in the current mode (For e.g. three times if using dual mode), however, they are only shown in verbose level one and above modes.

### 3.2.6 Display and Output Modules

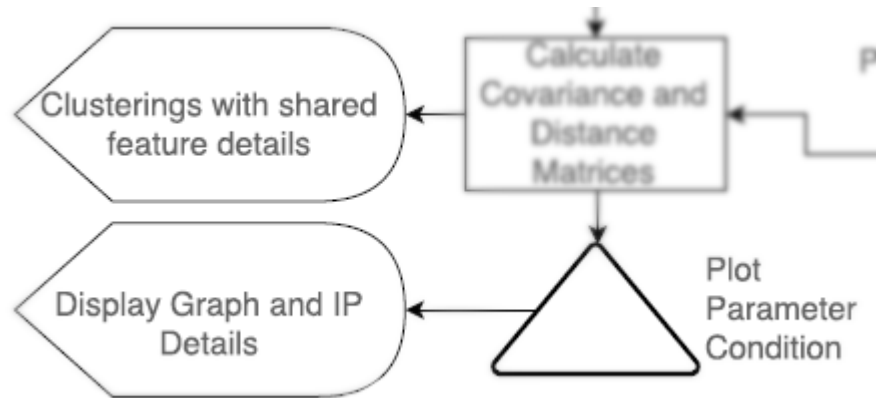


Figure 3.8: Representation of the display modules from infrastructure diagram Figure 3.1 with irrelevant modules blurred out

The application will display the main output at the end of its execution as shown in the infrastructure diagram with the symbol from Figure 3.8. The application will show a running output as the calculations and cluster iterations are performed, providing that the verbosity level is set to one or higher. The verbosity parameter will only effect the text output and not the graphing interface of the application which will remain identical, an example of full text output using dual mode can be found at B.1.1. Without verbose output selected the text output will only show the cluster details (for both formats if using dual mode). In order for the application to display the graphing interface on clustering class completion the application must be run with the parameter -p as mentioned in the usage descriptor found in A.2. The graphs displayed within the graphing GUI are fully manipulatable with zoom, pan and save as functions, however, the graphing interface will differ depending on which mode the application is using.

**dual mode** graphing interface will display five graphs and a small text segment underneath the bottom graph. The five segments are arranged in a grid of 4 with a large graph beneath the grid spanning both columns as the 5th. The first row of two graphs



displays the Nmap output with the right graph solely displaying the centroids and the left displaying the entire Nmap clustering with IP addresses. The second row displays the Nessus graphs in the same configuration as Nmap above. The 3rd row graph displays the IP addresses from the clusters with less than 3 IP addresses (this value is the default and can be changed by following the instruction described in section 3.4 and Figure 3.5). The reason for why this is done has been explained within the application brief at section 3.1.1 and the method used explained within section 3.4. The text segment beneath the 3rd row graph displays the detected operating system information and the probability of that detection being true. This is shown for each IP address in the small cluster IP addresses graph just above the text segment. An example of the dual mode graphing interface can be found in Figure B.1.2 and Figure B.4.

**Standard mode** graphing interfaces are similar to each other as they contain the same two graph split with just different information. The graphs are similar to that of dual mode without the small combined cluster graph of which the left graph includes the full clustering with IP addresses and the right graph solely contains the centroids of those clusters. Two examples of this mode can be found in the Appendix as Figure B.2 for Nessus clustering and Figure B.3 for Nmap clustering, along with the commands used to achieve them.

### 3.3 Mathematical Model and Theory

The following information consists of mathematical representations for the modules in the applications primary class architecture, as shown in figure 3.1. Parse XML files into matrices  $V$  for each input file, consisting of two dimensions  $m$  and  $n$  respectively. The matrices are then vectorised from JSON strings outputted from the parser classes into numerical float representations. These matrices' data is then normalized using the following formula denoted in the journal of Normalization: A Preprocessing Stage, Patro & Sahu (2015).: Where  $V_n$  is the  $n$ th dimension of matrix  $V$  and contains the features for each label  $V_m$ .

$$V_n = \frac{(V_n - \min \{V_n\})}{(\max \{V_n\} - \min \{V_n\})} (new_{\max\{V_n\}} - new_{\min\{V_n\}}) + new_{\min \{V_n\}}$$

This application will normalise the data to the range of 0 to 1 in which the formula is adjusted to the following:

$$V_n = \frac{(V_n - \min \{V_n\})}{(\max \{V_n\} - \min \{V_n\})}$$

The  $V_n$  dimension of matrix  $V$  of each input file is then passed through a principle component analysis formula for dimensionality reduction to two dimensions. This is due to the current number of dimension layers directly correlating to a feature of which there is an average of three hundred per IP address (or  $V_m$ ). This is calculated by using the SKlearn PCA decomposition library which uses the probabilistic model of PCA defined in the journal of Probabilistic Principle Component Analysis, Tipping & Bishop (1999). This algorithm is too complex to be included within the scope of this thesis and therefore is not described here. The result of this PCA algorithm returns a new matrix with two dimensions, which is then used in the K-means clustering algorithm as input with the IP address labels (or  $V_m$ ).

The standard Lloyds algorithm for K-means clustering was then used against the PCA output vectors which can be described as the following five stages. This is carried out for each clustering iteration in the cluster class depending on its current mode.

1. Clusters the Vectors inputs into K number of clusters where K is either assigned manually, iteratively or calculated using the Gap statistic and Elbow values explained below. This depends on the current application mode and user parameters at run time.
2. Select K number of points at random locations to act as cluster centroids.
3. Assign objects to their closest plotted centroid per the selected distance function (Euclidian as default).
4. Calculate the new centroid for each cluster in this new clustering.
5. Repeat steps 2, 3 and 4 until the same points are assigned to each cluster in consecutive rounds and no changes are present in the clustering. In the case of automatic mode, the K value will increment up to a maximum equalling the number of IP addresses or labels in  $V_m$ .

However, the user can also select the use of DBscan or Agglomerative clustering algorithms instead of K-means for any mode apart from dual input. However, due to the limited usage they provide for the applications design goals, will not be described in this thesis.

Each clustering result of this algorithm has several calculations carried out upon it in order to validate and provide scores to the clustering. These calculations are denoted in section 3.6. The primary function used to score these clusterings is a silhouette value which has been defined in the journal, Silhouettes: A graphical aid to the interpretation and validation of cluster analysis, Rousseeuw (1987). A full description of what this value provides for the user is included at the end of section 3.6. The formulas used to calculate the silhouette value are taken from the previously mentioned journal and are described below.

For each vector  $i$ ,  $a(i)$  would be the average dissimilarity of vector  $i$  with all other vectors within the same cluster.  $b(i)$  is the lowest average dissimilarity of vector  $i$  to any other cluster of which vector  $i$  is not a member of. The cluster with the lowest value for  $b(i)$  is the closest neighbouring cluster and therefore the most similar. This defines the following formula for silhouette value  $s(i)$ .

$$s(i) = \frac{b(i) - a(i)}{\max\{a(i), b(i)\}}$$

The resulting clusterings are sorted based on the filters mentioned in 3.6 and the best clustering is chosen. This best clustering is passed through into the small combined cluster algorithm described in section 3.4 as well as the display class in 3.8.

When the application uses the K-means clustering algorithm and a value for K is not defined, the application will attempt to calculate the optimal value for K via either the Gap Statistic or Elbow value methods depending on the users preferences. This is changed within the optimal k means class with the default value being the Gap Statistic method. The python implementation of the Gap Statistic method can be found in the Appendix A.6 with the formulas used were originally developed by Stanford researchers from the journal, Estimating the number of clusters in a data set via the gap statistic, Tibshirani et al. (2001). The implementation of the researchers algorithm within this application can be described by the following points.

1. Cluster the data via K-means from the range of  $k=1$  to  $k=\max$  and calculate the variance quantity variable  $W_k$  (primarily used for elbow method).
2. Generate the reference data sets  $B$  and cluster them with the same  $k$  values as previous step.
3. Calculate the gap statistic value with the following formula:

$$Gap(k) = \left(\frac{1}{B}\right) \sum_{b=1}^B \log W_{kb}^* - \log W_k.$$

4. Calculate the standard deviation using:

$$sd(k) = \left[ \left(\frac{1}{B}\right) \sum_b (\log W_{kb}^* - \left(\frac{1}{B}\right) \sum_b \log W_{kb}^*)^2 \right]^{1/2}$$

5. Then define  $s_k = \sqrt{1 + \frac{1}{B}sd(k)}$  which allows the optimal number of  $K$  to be the smallest  $k$  such that  $Gap(k) \geq Gap(k+1) - s_{k+1}$ .

The Covariance and distance matrices displayed prior to graphing output are calculated using the following methods. More information on these values can be found in 3.2.5. Covariance for a given centroid vector dimensions  $x$  and  $y$  is calculated using the formula:

$$cov(x, y) = \frac{(\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y}))}{n - 1}$$

This is calculated for each centroid in the chosen clustering and displayed in a matrix. The distance matrix displays the Euclidian distance between the  $x$  and  $y$  values for each centroid in the clustering.

### 3.4 Application Testing

Unfortunately due to the sensitive nature of the data required by the application, there are no datasets to be found publicly available online. Therefore, in order to design a proof of concept test procedure a dataset must be created using a significant network size to simulate the proportions of a real organizational network. The size requirements of this dataset limit the use of virtualized machine networks due the

raw processing power required for a network of that size, hence, the permission was granted to allow for the scanning of a private network laboratory in the University of Abertay. The laboratory consisted of approximately 50 online devices at the time of scanning. The raw scan output files for Nmap and Nessus can be found within the thesis artefacts, hacklab analysis folder with the file names hacklab\_new.xml and hacklab\_new.nessus respectfully.

The creation of this dataset required installing the Nmap and Nessus scanner tools on a separate device plugged into the network for this sole purpose. The device used to enter the network and execute the scans was a Dell latitude E6410 laptop running the latest rolling release kernel of Arch Linux. This setup was used specifically to replicate a similar scenario to that of what an industry professional might encounter. The latest stable releases of the required software were used. At the time of writing this thesis (April 2017) those reference numbers were as follows:

- Nmap 7.40
- Nessus 6.10.5
- Python 2.7.13
- Python Libraries
  - backports-abc==0.4
  - bokeh==0.12.1
  - certifi==2016.8.8
  - cyclr==0.10.0
  - futures==3.0.5
  - Jinja2==2.8
  - MarkupSafe==0.23
  - matplotlib==1.5.1
  - numpy==1.11.1
  - pyparsing==2.1.8
  - python-dateutil==2.5.3
  - pytz==2016.6.1

- PyYAML==3.11
- requests==2.11.0
- scikit-learn==0.17.1
- scipy==0.18.0
- singledispatch==3.4.0.3
- six==1.10.0
- tornado==4.4.1
- tabulate==0.7.7

The Nmap scan was created using the following command:

---

```
1 nmap -A -O -oX hacklab.xml 10.0.0.0/24
```

---

The -Ox flag in this command exports the scan as an XML which is the only export format the clustering application currently supports. The Nessus scan was created by using the default scan policy in the advanced scan mode with the same scope as nmap, 10.0.0.0/24. This was then exported as the latest version of the proprietary Nessus XML format from the scan overview page.

Once both scans have completed the next step is to remove the testing device used from the XML files manually due to each scanner retrieving greatly differing results from the testing machine compared to other machines on the network. This step is crucial and ignoring it will manipulate the final result of the clustering algorithm rendering it inaccurate. The artefact files have had this step done prior to submission.

The scan files have then been passed through the clustering application using the following commands for each input type in automatic mode:

### Nmap

---

```
1 cluster.py -s automatic -vv -p "../hacklab analyses/hacklab_new.xml"
```

---

### Nessus

---

```
1 cluster.py -s automatic -vv -p -N "../hacklab analyses/hacklab_new.nessus"
```

---

### Dual Input

---

```
1 cluster.py -s automatic -vv -p -t -N -tp "../hacklab
analyses/hacklab_new.xml" "../hacklab analyses/hacklab_new.nessus"
```

---

More information and the results for this proof of concept test can be found in Appendix section B.2. These results will be validated within the thesis conclusions.

## Chapter 4

# Discussion and Results

In this chapter, the application developed and the thesis as a whole will be reviewed.

The aim of this dissertation was to design and develop a proof of concept tool to be utilised by the red teaming individuals such as penetration testers during a security assessment. The application was required to be versatile, portable and reliable allowing it to be used in any size of network topology. From the Application Brief 3.1.1, the application serves as an additional tool to be used during the initial enumeration stage of a penetration test. It will provide the user with several graphs and statistics on the network which are not provided by currently available commercial tools, such as a visual network overview of machine differences. The application primarily targets are security professionals and network administrators whom want a visual representation of their overall network security due to how well the application scales to network sizes.

The main findings of this thesis show that red teaming can benefit greatly from machine learning techniques and more research needs to be carried out to move the industry forward towards more accurate, faster and efficient software tools. The results from the proof of concept testing of the application against a network with approximately fifty hosts can be found at Appendix B.2 and information on how the test was carried out can be found in section 3.4. The results show that the application was successful in classifying the difference and vulnerability level of the hosts on the network. The following paragraphs will analyse the results in order to show how this conclusion was achieved.

The application relies on the two scan files used, Nmap and Nessus, for its accuracy as they provide the bases of all the information used for every calculation involved in



10.0.0.1
10.0.0.185
10.0.0.3
10.0.0.5
10.0.0.7

Figure 4.1: The five detected vulnerable IP addresses from the Hacklab Analysis testing section in 3.4

the software. This means that these two scan files can be used to validate the application results by manually checking each scanner output within their respective applications.

First, the results of the application must be outlined and analysed to then be compared against the files, this is carried out in the following subsection.

#### 4.0.1 Test Results Analysis and Validation

This subsection will summarise the results of the application testing followed by the validation and in depth explanation of those results.

##### 4.0.1.1 Analysis of Results

The application detected the five IP addresses in Figure 4.0.1.1 as the most vulnerable based on their vulnerabilities and properties. This is shown in both the applications GUI output in Figure 4.2 and its text output in B.2.1:

Figure 4.2 shows the lower segment of the full results in Figure B.4, specifically, the vulnerability clustering section of the application explained in 3.4. The top graph represents the IP addresses from the small clusters, in this case clusters with less than three IP addresses, clustered via how similar they are to each other and their vulnerability properties, if any were detected by the Nessus scanner. This means that these hosts have the greatest difference from the majority of others in the network and most likely have known vulnerabilities. However, if the majority of hosts on the network have the same vulnerability, then the application will only display the *different* ones, resulting in those hosts not being shown. This scenario is discussed further on in the discussion.

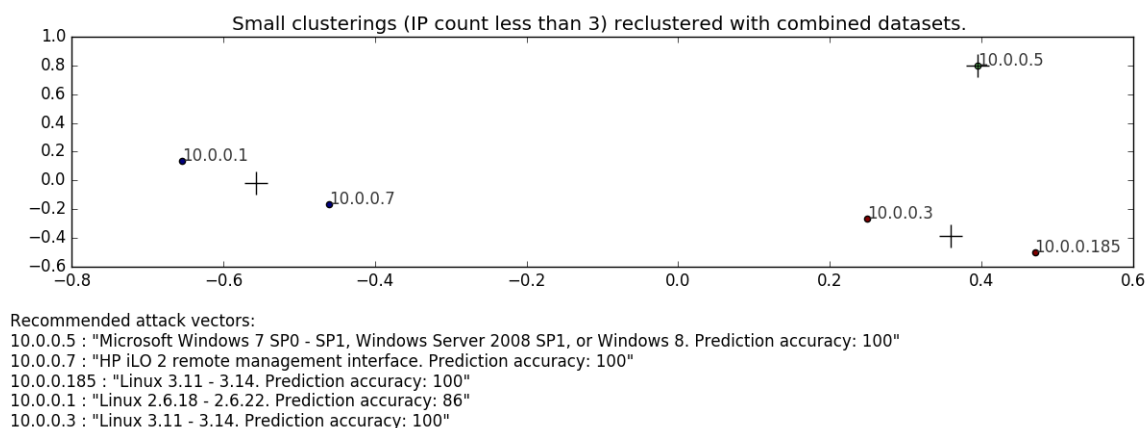


Figure 4.2: Short excerpt from the applications graphing output in automatic dual mode from Abertay Universitys Hacklab network analysis 3.4

From the applications maximum verbosity text output in B.2.1, line 237 shows that the five selected hosts were selected based on 491 different features per host, which was achieved by joining both feature sets together for this part of the application. The features are passed through PCA, similarly to that of the regular clustering section of the application to reduce these 495 features into a two-dimensional dataset. The split for this feature set in this particular test was 288 for Nessus (line 7) and 203 for Nmap (line 11), this relates to a 59% bias in the clustering towards the Nessus parsed features. Due to Nessus providing all the information about the vulnerabilities and Nmap providing a greater overall amount of system information, the two sets of data work together to give the application a more complete overview of the network. The Nessus bias allows the application to prioritise the vulnerabilities in the clustering algorithm over general system information. However, this ratio is fluid and changes depending on the Nessus configuration used and the networked hosts themselves. This unequal ratio is due to the fact that, while the Nmap scanner returns the same amount of data per machine without variance, the Nessus scanner outputs vary greatly depending on the number and detail of the vulnerabilities detected. This means that a network with a greater number of detailed vulnerabilities on its hosts will provide a much greater clustering bias towards the Nessus data, rather than the Nmap data. This bias only occurs when both data-sets are combined during the final vulnerability detection stage of the application explained in 3.4 and not during individual tool clusterings. Based off this information, the application recommends these afore mentioned five hosts to be targeted first during a manual security assessment as they will provide the greatest chance of success.

The text information beneath the graph in Figure 4.2 shows the operating systems detected for each host in 4.0.1.1 as well as the probability percentage of this prediction being accurate. The four graphs in figure 4.3 represent the other half of the applications output in automatic dual mode. Each graph is titled to show the user what they each represent. As mentioned previously these clusterings use their respectable tool files and thus, do not contain the bias that Figure 4.2 has. Two of these four graphs are the exact same as those found if the application were to execute using the single dataset mode for each tool as such demonstrated in Figures B.2 and B.3. This section of the output can be found explained in section 3.8. The IP addresses of hosts in Figure 4.0.1.1 can be seen within the two clusterings, showing how those hosts relate to the other hosts of larger clusters, in the same terms as Figure 4.2 but keeping the tool output features to their individual graphs.

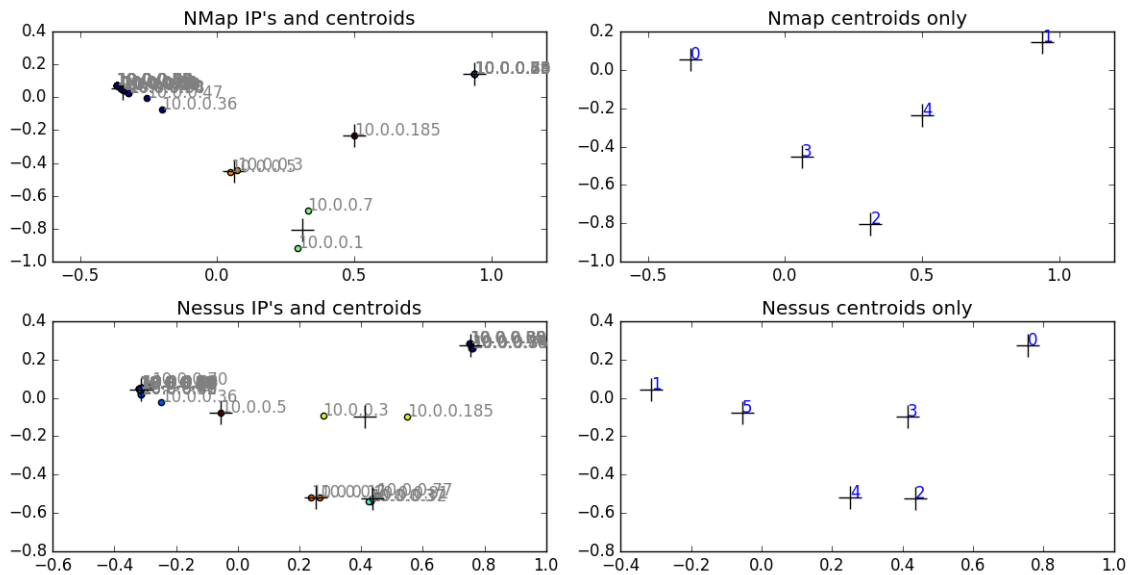


Figure 4.3: standard tool output graphs from applications graphing output in automatic dual mode from Abertay University Hacklab network analysis 3.4

#### 4.0.1.2 validation

In order to determine whether the detected IP addresses in Figure 4.0.1.1 are that of the most differential hosts, each tool output must be investigated individually. By uploading the Nessus file onto a Nessus web interface of version 6 or above, it is possible to see an overview of the file with colour coded bars for each host. This is represented in Appendix Figures B.5 and B.6. The original image was split into two

Figures simply due to the size being too large for this paper. The key for the colour coded bars in the Nessus overview is as follows:

Red	Critical rank vulnerability
Yellow	Medium rank vulnerability
Green	Low rank vulnerability
Blue	Host information

Through analysis of the Nessus overview using the key above, the differences in vulnerabilities per host are made apparent as visually intended by the developer, Tenable<sup>TM</sup>. By specifically viewing the selected hosts in this overview, it can be observed that **they are indeed, the most unique compared to the other hosts** in terms of Critical, Medium and Low ranked vulnerabilities. When using this method to compare hosts, it is important to not provide as much weight to the key colour blue representing the host information as to the vulnerability codes. This is because the blue host information is given much lower importance than the vulnerabilities in the Nessus parser class. If the keys shown above were weighted equally, the clustering would not have the bias towards the vulnerabilities and result in a less accurate prediction by the algorithm. Unfortunately, there is no similar visualisation method to validate the Nmap graph and requires manually reading the output XML file and comparing It to the application graphs.

Another method to validate the network is a manual approach, which requires in-depth prior knowledge of the network used. This approach is possible for individuals such as network administrators who might use this tool for the purpose of confirming the results of a security assessment, and not to discover new information. This approach is not possible for scenarios such as capture the flag events where there is no prior knowledge of the network and its hosts. The manual approach involves comparing the applications verbose text output, such as each clusters shared features, to the individuals prior knowledge of the network. The following section will give a critical evaluation of the application.

#### 4.0.2 Critical Evaluation

This section will provide a short critical evaluation of the application and its performance in its intended role.

As previously mentioned in this discussion, if the used network contains a large number of hosts with the same vulnerability, then those hosts become the majority and since the application, in its essence, finds the differences in hosts, the clustering might not reflect the hosts with that vulnerability accurately. This problem is difficult to solve completely as it would involve changing the applications core design architecture. A possible untested solution however, is as follows. By adding a new layer on top of the parser classes to make it aware of common vulnerabilities in the dataset, then displaying them to the user directly would mitigate this problem. Another possible workaround is simply for the user to manually check for any common, major vulnerabilities within the Nessus overview as mentioned in the previous section and take them into account when considering the applications results.

Unfortunately, due to the nature of the datasets used including sensitive information, there are no online datasets that the application could use. This makes the creation of a proper testing procedure difficult as all data used for this application within this project was created specifically for this project. This means that the application is to be taken simply as a proof of concept and not a finalised design.

Due to project time constraints, the primary automatic dual mode of the application was only able to be implemented with the K-means algorithm however, both DBscan and agglomerative implementations are already fully programmed and integrated into the model. The only issue stopping these algorithms to be integrated into this mode is the fact the sklearn python library (used to calculate several of the clustering functions) does not include centroid support for them and one would have to be manually programmed, which would take more time than was available for the project. Once that feature has been implemented, the two algorithms can be re-enabled with minimal difficulty.

The application has been optimised for model prediction accuracy over performance times, however, each class module within the application has been programmed modularly allowing for the easy modification by others without prior knowledge of how it was created. Variables may be changed easily, such as the IP cut off limit for the vulnerability detection stage explained in 3.4 without fear of it causing problems elsewhere in the application. Another such example of modularity is allowing the option of modifying the parsing classes to change what it taken as features and labels from each tool output. When using a network with less than a hundred hosts such as that

of in the testing chapter, the application will take between one and three minutes to complete on the dual automatic mode. Other modes such as manual and assisted can be completed in less than thirty seconds. This performance could be improved by converting parts of the application models to a language such as C++ although this would greatly reduce the customization level of the application, which is one of the primary design features. Upon successful completion of the clusterer, the application will write to two files. One of which is a dot file including the primary clustering and the other is a target file including the selected hosts from the results. The latter file can be imported into many tools such as Metasploit and Armitage to manually assess those hosts. The former can be imported into graphing tools such as Gephi to interface with them in a different manner than the integrated interface. Both these output files can be easily disabled they are not needed by modifying one line at the end of the initialization class.

### **4.0.3 Discussion Summary**

The findings in this thesis contribute towards developing a machine learning arsenal of tools that can be used by security professionals, as well as help balance the research in the security industry by providing a unique red teaming proof of concept tool. The results of the testing are of direct practical relevance to the red team, and such align with the design goals highlighted in 3.1.1. Only a couple of other studies, to our knowledge, have examined the use of machine learning in red teaming. It is clear that further research in this area is necessary before machine learning can become common place in this industry sector, however, this paper has helped cleared ground for this research to take place in. This paragraph marks the end of the discussion and the following section will provide conclusions to the paper.

# Chapter 5

## Conclusions

# Appendix A

## Code

### A.1 Requirements python file

---

```
1 backports-abc==0.4
2 bokeh==0.12.1
3 certifi==2016.8.8
4 cyclers==0.10.0
5 futures==3.0.5
6 Jinja2==2.8
7 MarkupSafe==0.23
8 matplotlib==1.5.1
9 numpy==1.11.1
10 pyparsing==2.1.8
11 python-dateutil==2.5.3
12 pytz==2016.6.1
13 PyYAML==3.11
14 requests==2.11.0
15 scikit-learn==0.17.1
16 scipy==0.18.0
17 singledispatch==3.4.0.3
18 six==1.10.0
19 tornado==4.4.1
20 tabulate==0.7.7
```

---

### A.2 Application Usage Parameters

The following includes the usage and parameter definitions for the application.

---

```
1 usage: cluster.py [-h] [-s {manual,automatic,assisted}]
2                  [-c {kmeans,dbscan,agglomerative}]
3                  [--metric {euclidean,cosine,jaccard}] [-N] [-n N_CLUSTERS]
4                  [-e EPSILON] [-m MIN_SAMPLES] [-cent] [-t] [-tp twinpath]
```



```

5         [-p] [-v]
6         path [path ...]
7 Cluster NMap/Nessus Output
8 positional arguments:
9   path                Paths to files or directories to scan
10 optional arguments:
11   -h, --help          show this help message and exit
12   -s {manual,automatic,assisted}, --strategy {manual,automatic,assisted}
13   -c {kmeans,dbscan,agglomerative}, --method {kmeans,dbscan,agglomerative}
14   --metric {euclidean,cosine,jaccard}
15   -N, --nessus        use .nessus file input
16   -n N_CLUSTERS, --n_clusters N_CLUSTERS
17                       Number of kmeans clusters to aim for
18   -e EPSILON, --epsilon EPSILON
19                       DBSCAN Epsilon
20   -m MIN_SAMPLES, --min_samples MIN_SAMPLES
21                       DBSCAN Minimum Samples
22   -cent, --centroids  plot only centroids graph, requires the use of "-p"
23   -t, --twin          use both input formats to calculate vulnerable single
24                       clusters, use with -tp and -N
25   -tp twinpath, --twinpath twinpath
26                       path to nmap xml if using twin clustering
27   -p, --plot          Plot clusters on 2D plane
28   -v, --verbosity    increase output verbosity

```

---

## A.3 Initialization Class - cluster.py

The following contains the raw code for the primary initializer class of file 'cluster.py' in the application parent folder. It is used to start the application using parameters from the usage, in Appendix A.2. The code has been thoroughly commented with the intention of being modified by a potential tester for individual operational requirements. This is further enforced by the application being entirely modular with each module able to be modified without harming the others. The cluster subclass has been removed from this code and placed in A.5 in order to promote the legibility of this appendix thus must not be executed direction without concatenation with the afore mentioned subclass first.

---

```

1 import logging
2
3 from sklearn.preprocessing import normalize
4
5 from clusterer_parts.optimal_k_k_means import optimalK

```

```

6 from clusterer_parts.analysis import get_common_features_from_cluster,
   get_common_feature_stats
7 from clusterer_parts.clustering import cluster_with_dbscan,
   cluster_with_kmeans, precompute_distances, \
8   cluster_with_agglomerative, cluster_interactive, get_centroids,
   cluster_single_kmeans, get_k
9 from clusterer_parts.display import print_cluster_details,
   generate_dot_graph_for_gephi, create_plot, \
10  create_plot_centroids, create_plot_only_centroids, twin,
   remove_large_clusters
11 from clusterer_parts.optimizing import sort_items_by_multiple_keys
12 from clusterer_parts.reduction import pca
13 from clusterer_parts.validation import validate_clusters,
   get_average_distance_per_cluster
14 from clusterer_parts.vectorize import vectorize
15 import numpy as np
16 from tabulate import tabulate
17 firstpass = True
18
19 if __name__ == "__main__":
20     import argparse
21
22     parser = argparse.ArgumentParser(description=u'Cluster NMap/Nessus
   Output')
23
24     parser.add_argument('path', metavar='path', type=str, nargs='+',
   default=None,
25                         help="Paths to files or directories to scan")
26
27     parser.add_argument('-s', '--strategy', default="automatic",
   choices=["manual", "automatic", "assisted"])
28     parser.add_argument('-c', '--method', default="kmeans",
   choices=["kmeans", "dbscan", "agglomerative"])
29     parser.add_argument('--metric', default="euclidean",
   choices=["euclidean", "cosine", "jaccard"])
30     parser.add_argument('-N', '--nessus', default="false", required=False,
   action='store_true',
31                         help='use .nessus file input')
32
33     parser.add_argument('-n', '--n_clusters', type=int, default=2,
   help='Number of kmeans clusters to aim for')
34     parser.add_argument('-e', '--epsilon', type=float, default=0.5,
   help='DBSCAN Epsilon')
35     parser.add_argument('-m', '--min_samples', type=int, default=5,
   help='DBSCAN Minimum Samples')
36     parser.add_argument('-cent', '--centroids', default=False,
   required=False, action='store_true',

```

```

37         help='plot only centroids graph, requires the use of
38         "-p"')
39     parser.add_argument('-t', '--twin', default=False, required=False,
40         action='store_true',
41         help='use both input formats to calculate vulnerable
42         single clusters, use with -tp and -N')
43     parser.add_argument('-tp', '--twinpath', metavar='twinpath', type=str,
44         required=False,
45         help='path to nmap xml if using twin clustering')
46     parser.add_argument('-p', '--plot', default=False, required=False,
47         action='store_true',
48         help='Plot clusters on 2D plane')
49     parser.add_argument("-v", "--verbosity", action="count",
50         help="increase output verbosity")
51     args = parser.parse_args()
52     logging.basicConfig(format='%(asctime)s %(process)s %(module)s
53         %(funcName)s %(levelname)-8s :%(message)s',
54         datefmt='%m-%d %H:%M')
55
56     if args.verbosity == 1:
57         logging.getLogger().setLevel(logging.INFO)
58     elif args.verbosity > 1:
59         logging.getLogger().setLevel(logging.DEBUG)
60
61     if (args.twin == False):
62
63         # Vectorize our input
64         logging.info("Vectorizing Stage")
65         vector_names, vectors, vectorizer = vectorize(args.path,
66             args.nessus)
67         logging.debug("Loaded {0} vectors with {1}
68             features".format(len(vector_names), vectors.shape[1]))
69         logging.info("Vectorizing complete")
70
71         # normalise vectors first before passing them through PCA. PCA uses
72         2 dimensions
73         logging.info("Normalising the vectors")
74         normalized_vectors = normalize(vectors)
75         logging.info("Reducing vectors to two dimensions with PCA")
76         reduced_vectors = pca(normalized_vectors)
77         logging.debug(
78             "reduced to {0} vectors with {1}
79             dimensions".format((reduced_vectors.shape[0]),
80                 reduced_vectors.shape[1]))

```

```

72
73 # Cluster the vectors
74 logging.info("Clustering")
75 labels = cluster(vector_names, vectors, reduced_vectors,
76                 normalized_vectors, vectorizer, args.strategy,
77                 args.method, args.n_clusters, args.epsilon,
78                 args.min_samples, args.metric)
79 logging.info("Clustering Complete")
80 # Test cluster validity
81 overall_score, per_cluster_score = validate_clusters(vectors,
82 labels)
83
84 # Analysis relevant to the person reading results
85 universal_positive_features, universal_negative_features,
86 shared_features = get_common_features_from_cluster(
87 vectors, labels, vectorizer)
88
89 # logging.debug("Shared features: {0}".format(shared_features))
90
91 # Reduce results and relevant information to per cluster data
92 cluster_details = {}
93 for cluster_id in per_cluster_score.keys():
94     cluster_details[cluster_id] = {
95         "silhouette": per_cluster_score[cluster_id],
96         "shared_positive_features":
97             shared_features[cluster_id]['positive'],
98         # "shared_negative_features":
99             shared_features[cluster_id]['negative'],
100         "ips": [vector_names[x] for x in xrange(len(vector_names))
101                 if labels[x] == cluster_id]
102     }
103 print "Note: shared features does not retain keys from XML and
104       therefore wont always be human readable."
105 print_cluster_details(cluster_details, shared_features)
106
107 if args.plot:
108     # only kmeans centroids for now
109     if no_clusters.startswith("kmeans") :
110         logging.debug("Getting centroids using reduced vectors:")
111         # global centroidskmeans
112         # take just cluster number from result string
113         n_clusters = no_clusters.split("=", 1)[1]
114         n_clusters = int(n_clusters.rstrip(','), 1)[0])
115         logging.debug("nclusters: " + str(n_clusters))
116
117         centroidskmeans = get_centroids(reduced_vectors, n_clusters)
118         logging.debug("attempting to plot the following centroids:\n

```

```

111         " + str(centroidskmeans))
112
113     # covariance
114     x = centroidskmeans[:, 0]
115     y = centroidskmeans[:, 1]
116     X = np.vstack((x, y))
117     cov = np.cov(X)
118     logging.info("Centroids Covariance Matrix:\n
119                  {0}".format(cov))
120
121     # print similarity distance between centroids
122     matrix = precompute_distances(centroidskmeans,
123                                  metric=args.metric)
124     matrixTable = tabulate(matrix)
125     logging.info(
126         "distance matrix between centroids using metric: {0}
127         :\n{1}".format(args.metric, matrixTable))
128
129     if args.centroids:
130         create_plot_only_centroids(reduced_vectors, labels,
131                                   vector_names, centroidskmeans, n_clusters)
132     else:
133         create_plot_centroids(reduced_vectors, labels,
134                               vector_names, centroidskmeans, n_clusters,
135                               cluster_details)
136
137     # manually selected kmeans though arguments
138     elif args.method == "kmeans" and args.strategy != "automatic":
139         if get_k() > 0:
140             centroidskmeans = get_centroids(reduced_vectors, get_k())
141         else:
142             centroidskmeans = get_centroids(reduced_vectors,
143                                             args.n_clusters)
144
145     logging.debug("attempting to plot the following centroids:
146                  \n" + str(centroidskmeans))
147
148     # covariance
149     x = centroidskmeans[:, 0]
150     y = centroidskmeans[:, 1]
151     X = np.vstack((x, y))
152     cov = np.cov(X)
153     logging.info("Centroids Covariance Matrix:\n
154                  {0}".format(cov))
155
156     # print similarity distance between centroids
157     matrix = precompute_distances(centroidskmeans,

```

```

        metric=args.metric)
149     matrixTable = tabulate(matrix)
150     logging.info(
151         "distance matrix between centroids using metric: {0}
           :\n{1}".format(args.metric, matrixTable))
152
153     if args.centroids:
154         if get_k() > 0:
155             create_plot_only_centroids(reduced_vectors, labels,
                vector_names, centroidskmeans, get_k())
156         else:
157             create_plot_only_centroids(reduced_vectors, labels,
                vector_names, centroidskmeans, args.n_clusters)
158     else:
159         if get_k() > 0:
160             create_plot_centroids(reduced_vectors, labels,
                vector_names, centroidskmeans, get_k(),
161                 cluster_details)
162         else:
163             create_plot_centroids(reduced_vectors, labels,
                vector_names, centroidskmeans, args.n_clusters,
164                 cluster_details)
165     else:
166         logging.debug("plotting standard graph")
167         create_plot(reduced_vectors, labels, vector_names)
168     # Write DOT diagram out to cluster.dot, designed for input into
        Gephi (https://gephi.org/)
169     with open("cluster.dot", "w") as f:
170         f.write(
171             generate_dot_graph_for_gephi(precompute_distances(vectors,
                metric=args.metric), vector_names, labels))
172
173     elif args.twin == True and args.strategy == "automatic":
174
175         logging.debug("twin flag enabled")
176         logging.debug("tp: {0} , path: {1}".format(args.twinpath,
            args.path))
177
178     # Vectorize our input for nessus
179     logging.info("Vectorizing Stage for Nessus")
180     Nvector_names, Nvectors, Nvectorizer = vectorize(args.path,
        args.nessus)
181     logging.debug("Loaded {0} vectors with {1}
        features".format(len(Nvector_names), Nvectors.shape[1]))
182     logging.info("Vectorizing complete\n")
183
184     # Vectorize our input for nmap

```

```

185 logging.info("Vectorizing Stage for nmap")
186 twinpath = list()
187 twinpath.append(args.twinpath)
188 vector_names, vectors, vectorizer = vectorize(twinpath, False)
189 logging.debug("Loaded {0} vectors with {1}
      features".format(len(vector_names), vectors.shape[1]))
190 logging.info("Vectorizing complete\n")
191
192 # normalise vectors first before passing them through PCA. PCA uses
      2 dimensions
193 # nessus
194 logging.info("Normalising the nessus vectors")
195 Nnormalized_vectors = normalize(Nvectors)
196 logging.info("Reducing vectors to two dimensions with PCA")
197 Nreduced_vectors = pca(Nnormalized_vectors)
198 logging.debug(
199     "reduced to {0} vectors with {1}
      dimensions".format((Nreduced_vectors.shape[0]),
      Nreduced_vectors.shape[1]))
200 logging.info("Normalising complete\n")
201
202 # normalise vectors first before passing them through PCA. PCA uses
      2 dimensions
203 # nmap
204 logging.info("Normalising the nmap vectors")
205 normalized_vectors = normalize(vectors)
206 logging.info("Reducing vectors to two dimensions with PCA")
207 reduced_vectors = pca(normalized_vectors)
208 logging.debug(
209     "reduced to {0} vectors with {1}
      dimensions".format((reduced_vectors.shape[0]),
      reduced_vectors.shape[1]))
210 logging.info("Normalising complete\n")
211
212 # Cluster the vectors
213 logging.info("Clustering Nessus")
214 Nlabels = cluster(Nvector_names, Nvectors, Nreduced_vectors,
      Nnormalized_vectors, Nvectorizer, args.strategy,
215     args.method, args.n_clusters, args.epsilon,
      args.min_samples, args.metric)
216 logging.info("Clustering Complete\n\n")
217 # Test cluster validity
218 Noverall_score, Nper_cluster_score = validate_clusters(Nvectors,
      Nlabels)
219
220 # Cluster the vectors
221 logging.info("Clustering Nmap")

```

```

222     labels = cluster(vector_names, vectors, reduced_vectors,
223                     normalized_vectors, vectorizer, args.strategy,
224                     args.method, args.n_clusters, args.epsilon,
225                     args.min_samples, args.metric)
226 logging.info("Clustering Complete\n\n")
227 # Test cluster validity
228 overall_score, per_cluster_score = validate_clusters(vectors,
229 labels)
230
231 # Analysis relevant to the person reading results
232 # nessus
233 Nuniversal_positive_features, Nuniversal_negative_features,
234 Nshared_features = get_common_features_from_cluster(
235     Nvectors, Nlabels, Nvectorizer)
236
237 # Analysis relevant to the person reading results
238 # nmap
239 universal_positive_features, universal_negative_features,
240 shared_features = get_common_features_from_cluster(
241     vectors, labels, vectorizer)
242
243 # Reduce results and relevant information to per cluster data
244 # nessus
245 Ncluster_details = {}
246 for cluster_id in Nper_cluster_score.keys():
247     Ncluster_details[cluster_id] = {
248         "silhouette": Nper_cluster_score[cluster_id],
249         "shared_positive_features":
250             Nshared_features[cluster_id]['positive'],
251         "ips": [Nvector_names[x] for x in xrange(len(Nvector_names))
252                if Nlabels[x] == cluster_id]
253     }
254 print "Note: shared features does not retain keys from XML and
255     therefore wont always be human readable."
256 print "Printing Nessus cluster details\n"
257 print_cluster_details(Ncluster_details, Nshared_features)
258
259 print "\n\n"
260
261 # Reduce results and relevant information to per cluster data
262 cluster_details = {}
263 for cluster_id in per_cluster_score.keys():
264     cluster_details[cluster_id] = {
265         "silhouette": per_cluster_score[cluster_id],
266         "shared_positive_features":
267             shared_features[cluster_id]['positive'],
268         # "shared_negative_features":

```



```

260         shared_features[cluster_id]['negative'],
        "ips": [vector_names[x] for x in xrange(len(vector_names))
                if labels[x] == cluster_id]
261     }
262     print "Printing Nmap cluster details\n"
263     print_cluster_details(cluster_details, shared_features)
264
265     if args.plot:
266         # Nmap
267         logging.debug("Getting centroids using reduced vectors for
                        Nmap:")
268         # take just cluster number from result string
269         n_clusters = Nno_clusters.split("=", 1)[1]
270         n_clusters = int(n_clusters.rsplit(')', 1)[0])
271         logging.debug("nclusters: " + str(n_clusters))
272
273         centroidskmeans = get_centroids(reduced_vectors, n_clusters)
274         k = get_k()
275         logging.debug("attempting to plot the following centroids:\n
                        " + str(centroidskmeans) + "\n\n")
276
277         # Nessus
278         logging.debug("Getting centroids using reduced vectors for
                        Nessus:")
279         # take just cluster number from result string
280         logging.debug("nclusters: " + str(Nno_clusters))
281
282         Nn_clusters = no_clusters.split("=", 1)[1]
283         Nn_clusters = int(Nn_clusters.rsplit(')', 1)[0])
284         logging.debug("nclusters: " + str(Nn_clusters))
285
286         Ncentroidskmeans = get_centroids(Nreduced_vectors,
                                           Nn_clusters)
287         logging.debug("attempting to plot the following centroids:\n
                        " + str(Ncentroidskmeans) + "\n\n")
288         Nk = get_k()
289
290         # covariance for Nmap
291         x = centroidskmeans[:, 0]
292         y = centroidskmeans[:, 1]
293         X = np.vstack((x, y))
294         cov = np.cov(X)
295         logging.info("Nmap Centroids Covariance Matrix:\n
                        {0}".format(cov))
296
297         # covariance for Nessus
298         Nx = Ncentroidskmeans[:, 0]

```

```

299     Ny = Ncentroidsmeans[:, 1]
300     NX = np.vstack((Nx, Ny))
301     Ncov = np.cov(NX)
302     logging.info("Nessus Centroids Covariance Matrix:\n
        {0}".format(Ncov))
303
304     # print similarity distance between centroids
305     # Nessus
306     matrix = precompute_distances(centroidsmeans,
        metric=args.metric)
307     matrixTable = tabulate(matrix)
308     logging.info(
309         "distance matrix between centroids using metric for
        Nmap: {0} :\n{1}".format(args.metric, matrixTable))
310
311     # print similarity distance between centroids
312     # Nmap
313     Nmatrix = precompute_distances(Ncentroidsmeans,
        metric=args.metric)
314     NmatrixTable = tabulate(Nmatrix)
315     logging.info(
316         "distance matrix between centroids using metric for
        Nessus: {0} :\n{1}".format(args.metric,
        NmatrixTable))
317
318     small_ips = remove_large_clusters()
319
320     logging.info("IP's from clusters with less than 3 IP's:\n
        {0}".format((small_ips)))
321
322
323     #creates large array with 2nd dimension as large enough to
        hold both feature vectors
324     nesmap = np.zeros((len(small_ips),
        (Nvectors.shape[1]+vectors.shape[1])))
325
326
327     #for each single ip
328     for index in range(len(small_ips)):
329         # for each ip in vectors
330         features = 0
331         for index2 in range(vectors.shape[0]):
332             #if ip is equal to vector ip
333             #logging.debug("if {0} = {1}
        ".format(small_ips[index], vector_names[index2]))
334             if small_ips[index] == vector_names[index2]:
335                 #logging.debug("ip is equal to vector ip")

```

```

336         #for every one of this vectors features
337         for index3 in range(vectors.shape[1]):
338             #assign its features to single ip vector
339             nesmap[index,features] = vectors[index2,
340                 index3]
341             features +=1
342         break
343
344     for index2 in range(Nvectors.shape[0]):
345         #logging.debug("if {0} = {1}"
346             ".format(small_ips[index], Nvector_names[index2]))
347         if small_ips[index] == Nvector_names[index2]:
348             #logging.debug("ip is equal to vector ip")
349             for index3 in range(Nvectors.shape[1]):
350                 #append nessus features onto nmap features
351                 nesmap[index,features] = Nvectors[index2,
352                     index3]
353                 features += 1
354             break
355
356 logging.debug("Loaded {0} vectors with {1}
357     features".format(nesmap.shape[0], nesmap.shape[1]))
358 small_normalized_vectors = normalize(nesmap)
359 logging.info("Normalizing input and reducing vectors to two
360     dimensions with PCA")
361 final = pca(small_normalized_vectors)
362
363 logging.info("Resulting single IP vectors:\n
364     {0}".format(final))
365
366 Smatrix = precompute_distances(final, metric=args.metric)
367 SmatrixTable = tabulate(Smatrix)
368 logging.info(
369     "distance matrix between centroids of small combined
370     clusters: {0} :\n{1}".format(args.metric,
371         SmatrixTable))
372 clusterz = cluster_single_kmeans(final, 2)
373
374 logging.info("Writing recommended attack IP's to targets.txt
375     for exploitation\n {0}")
376 f = open('targets.txt', 'w')
377 for index in range(len(small_ips)):
378     f.write('{0}\n'.format(small_ips[index])) # python will
379         convert \n to os.linesep
380 f.close() # you can omit in most cases as the destructor
381     will call it

```

```

372         twin(reduced_vectors, labels, vector_names, centroidskmeans,
              n_clusters, cluster_details, Nreduced_vectors, Nlabels,
              Nvector_names, Ncentroidskmeans, Nn_clusters,
              Ncluster_details, small_ips, final, clusterz, twinpath)
373
374     else: print "not yet implemented #todo"

```

---

## A.4 Vectorization Class - vectorize.py

The following contains the raw code for the vectorize.py class. This class is called by the main cluster.py class at appendix A.3 in order to extract the information from the scan files as well as vectorize that information to be returned to the primary class.

---

```

1  import logging
2  from parsing import parsers
3  from parse_nessus import Nparsers
4  from single_ip_parsing_nmap import single_ip_parsers
5  import numpy as np
6
7
8  class Vectorizer:
9      """
10     This class handles the vectorizing of input
11     It additionally stores all pseudo vectors until we are ready for the
12     finished vectors
13     """
14
15     def __init__(self):
16         self.tokenized_strings = []
17         self.pseudo_vectors = {}
18
19     def add_string_to_ip(self, ip, string):
20         if ip not in self.pseudo_vectors:
21             self.pseudo_vectors[ip] = []
22
23         if string not in self.tokenized_strings:
24             self.tokenized_strings.append(string)
25
26         s_id = self.tokenized_strings.index(string)
27         self.pseudo_vectors[ip].append(s_id)
28
29     def parse_input(self, input_string, n):
30         ##n boolean value refers to nessus input only
31         if (n==True):
32             for parser in Nparsers:

```

```

32         if parser.can_parse_input(input_string):
33             results = parser.parse_input(input_string)
34             for key in results.keys():
35                 for s in results[key]:
36                     self.add_string_to_ip(key, s)
37     else:
38         for parser in parsers:
39             if parser.can_parse_input(input_string):
40                 results = parser.parse_input(input_string)
41                 for key in results.keys():
42                     for s in results[key]:
43                         self.add_string_to_ip(key, s)
44
45     def output_vectors(self):
46         vector_names = []
47         vectors = np.zeros((len(self.pseudo_vectors.keys()),
48                             len(self.tokenized_strings)), dtype=np.float)
49
50         for ip_index, ip in enumerate(self.pseudo_vectors.keys()):
51             vector_names.append(ip)
52             for s_index in self.pseudo_vectors[ip]:
53                 # Just set it to one, we want to ignore any case we see a
54                 # value more than once
55                 vectors[ip_index, s_index] = 1
56
57         return vector_names, vectors
58
59     def vectorize(files_to_vectorize, n):
60         vectorizer = Vectorizer()
61         for file_path in files_to_vectorize:
62             with open(file_path, "r") as f:
63                 vectorizer.parse_input(f.read(), n)
64
65         vector_names, vectors = vectorizer.output_vectors()
66
67         return vector_names, vectors, vectorizer
68
69     def parse_single_ips(files_to_vectorize, ips):
70         for file_path in files_to_vectorize:
71             with open(file_path, "r") as f:
72                 for parser in single_ip_parsers:
73                     #logging.debug("vectorisor selecting single ips")
74                     results = parser.parse_input(f.read(), ips)
75                     return results

```

---

## A.5 Clustering Algorithm subclass

The following code consists of the clustering algorithm used for all the modes within the application. It uses the provided parameters to decide between these modes as well as calls several other subclasses for code modularity.

---

```
1 def cluster(  
2     vector_names,  
3     vectors,  
4     reduced_vectors,  
5     normalized_vectors,  
6     vectorizer,  
7     strategy="automatic",  
8     cluster_method="kmeans",  
9     n_clusters=2,  
10    epsilon=0.5,  
11    min_samples=5,  
12    metric="euclidean",  
13 ):  
14     """  
15     Clustering options:  
16  
17     Manual:  
18     The user supplies all required information to do the clustering. This  
19     includes the clustering algorithm and  
20     hyper parameters,  
21     if no cluster count is provided the gap_statistic method will be used  
22     to calculate the optimal cluster count  
23  
24     Assisted:  
25     The user assists the algorithm by suggesting that some samples should  
26     or should not be clustered together  
27  
28     Automatic:  
29     The multiple clustering strategies and parameters are used in an  
30     attempt to get the best clusters  
31  
32     finds the least amount of clusters with atleast one shared feature  
33  
34     only uses gap statistic for small IP clusters  
35     """  
36  
37     global centroidskmeans, centroidagglo, centroiddb, no_clusters,  
38         Nno_clusters, Ncentroidskmeans
```

```

37     if strategy == "manual":
38         no_clusters = ""
39         if cluster_method == "kmeans":
40             #centroids_kmeans = get_centroids(reduced_vectors,
41                 n_clusters=n_clusters)
42             #logging.debug("centroids for kmeans:
43                 {0}".format(centroids_kmeans))
44             k, gapdf = optimalK(vectors, nrefs=3,
45                 maxClusters=reduced_vectors.shape[0])
46             return cluster_with_kmeans(reduced_vectors, n_clusters=k)
47
48         elif cluster_method == "dbscan":
49             return cluster_with_dbscan(reduced_vectors, epsilon=epsilon,
50                 min_samples=min_samples, metric=metric)
51
52         elif cluster_method == "agglomerative":
53             return cluster_with_agglomerative(reduced_vectors,
54                 n_clusters=n_clusters, metric=metric)
55
56         else:
57             # Unknown clustering method
58             raise NotImplementedError()
59
60     elif strategy == "assisted":
61         """
62         To display a information about a vector to a user, you can use the
63         following:
64         display_vector_index_details(vector_index, vectors, vector_names,
65             vectorizer)
66         """
67
68         return cluster_interactive(reduced_vectors, vectorizer, vectors,
69             vector_names)
70
71     elif strategy == "automatic":
72         results = []
73         smallest_cluster_count = vectors.shape[0]
74         # centroids works for only kmeans atm
75         for cluster_method in [
76             #todo add agglo and dbscan back in after they can return
77             centroids.
78             "kmeans" # ,
79             # "agglomerative",
80             # "dbscan",
81         ]:
82             if cluster_method == "kmeans":
83                 #this method is called X-means clustering
84                 logging.debug("Starting prospective KMeans clusterings")

```

```

75     move_to_next_method = False
76     # start at 2 clusters and end at smallest_cluster_count
77     for n_clusters in xrange(2, smallest_cluster_count):
78         logging.debug("Trying
79             {0}".format("kmeans(n_clusters={0})".format(n_clusters)))
80         labels = cluster_with_kmeans(reduced_vectors,
81             n_clusters=n_clusters)
82         overall_score, per_cluster_score =
83             validate_clusters(vectors, labels)
84         mean_distance =
85             get_average_distance_per_cluster(vectors, labels)[0]
86
87         tsp, msp, msn = get_common_feature_stats(vectors,
88             labels, vectorizer)
89
90         # If any cluster has 0 shared features, we just ignore
91         # the result
92         if msp <= tsp:
93             logging.debug("Not all clusters are informative (a
94                 cluster has 0 shared features) ")
95             continue
96         if len(set(labels)) > smallest_cluster_count:
97             move_to_next_method = True
98             # logging.debug("len(set(labels)): {0} >
99                 smallest_cluster_count:
100                 {1}".format(len(set(labels)),
101                     smallest_cluster_count))
102             break
103         if len(set(labels)) < smallest_cluster_count:
104             smallest_cluster_count = len(set(labels))
105         #too verbose
106         # logging.debug(repr((
107             #     overall_score,
108             #     min(per_cluster_score.values()),
109             #     mean_distance,
110             #     labels,
111             #     len(set(labels)),
112             #     tsp,
113             #     msp,
114             #     msn,
115             #     "kmeans(n_clusters={0})".format(n_clusters)
116             # )))
117         results.append(
118             (
119                 overall_score,
120                 min(per_cluster_score.values()),
121                 mean_distance,

```



```

112         labels,
113         len(set(labels)),
114         tsp,
115         msp,
116         msn,
117         "kmeans(n_clusters={0})".format(n_clusters)
118     )
119 )
120 if move_to_next_method:
121     continue
122
123 if cluster_method == "agglomerative":
124     logging.debug("Starting prospective Agglomerative
125                    clusterings")
126     move_to_next_method = False
127     for n_clusters in xrange(2, smallest_cluster_count):
128         logging.debug("Trying
129                        {0}".format("agglomerative(n_clusters={0})".format(n_clusters)))
130         labels = cluster_with_agglomerative(reduced_vectors,
131                                             n_clusters=n_clusters, metric=metric)
132         overall_score, per_cluster_score =
133             validate_clusters(vectors, labels)
134         mean_distance =
135             get_average_distance_per_cluster(vectors, labels)[0]
136
137         tsp, msp, msn = get_common_feature_stats(vectors,
138                                                  labels, vectorizer)
139
140         # If any cluster has 0 shared features, we just ignore
141         # the result
142         if msp <= tsp:
143             logging.debug("Not all clusters are informative (a
144                            cluster has 0 shared features) ")
145             continue
146         if len(set(labels)) > smallest_cluster_count:
147             move_to_next_method = True
148             break
149         if len(set(labels)) < smallest_cluster_count:
150             smallest_cluster_count = len(set(labels))
151
152     logging.debug(repr((
153         overall_score,
154         min(per_cluster_score.values()),
155         mean_distance,
156         labels,
157         len(set(labels)),
158         tsp,

```

```

151         msp,
152         msn,
153         "agglomerative(n_clusters={0})".format(n_clusters)
154     )))
155     results.append(
156         (
157             overall_score,
158             min(per_cluster_score.values()),
159             mean_distance,
160             labels,
161             len(set(labels)),
162             tsp,
163             msp,
164             msn,
165             "agglomerative(n_clusters={0})".format(n_clusters)
166         )
167     )
168     if move_to_next_method:
169         continue
170
171 if cluster_method == "dbscan":
172     logging.debug("Starting prospective DBSCAN clusterings")
173     distance_matrix = precompute_distances(vectors,
174                                           metric=metric)
175     min_distance =
176         sorted(set(list(distance_matrix.flatten())))[1]
177     max_distance =
178         sorted(set(list(distance_matrix.flatten())))[-1]
179     num_steps = 25.0
180     step_size = float(max_distance - min_distance) /
181         float(num_steps)
182     epsilon = min_distance
183     while True:
184         logging.debug("Trying
185             {0}".format("dbscan(epsilon={0})".format(epsilon)))
186         labels = cluster_with_dbscan(reduced_vectors,
187                                     epsilon=epsilon, min_samples=1,
188                                     distances=distance_matrix)
189         if len(set(labels)) == 1 and list(set(labels))[0] == 0:
190             break
191         overall_score, per_cluster_score =
192             validate_clusters(vectors, labels)
193         mean_distance =
194             get_average_distance_per_cluster(vectors, labels)[0]
195
196     tsp, msp, msn = get_common_feature_stats(vectors,
197                                             labels, vectorizer)

```

```

189
190         # If any cluster has 0 shared features, we just ignore
           the result
191     if msp <= tsp:
192         logging.debug("Not all clusters are informative (a
           cluster has 0 shared features) ")
193         epsilon += step_size
194         continue
195
196     logging.debug(repr((
197         overall_score,
198         min(per_cluster_score.values()),
199         mean_distance,
200         labels,
201         len(set(labels)),
202         tsp,
203         msp,
204         msn,
205         "dbscan(epsilon={0})".format(epsilon)
206     )))
207     results.append(
208         (
209             overall_score,
210             min(per_cluster_score.values()),
211             mean_distance,
212             labels,
213             len(set(labels)),
214             tsp,
215             msp,
216             msn,
217             "dbscan(epsilon={0})".format(epsilon)
218         )
219     )
220     epsilon += step_size
221
222     # Choose best clustering result based on the following attributes
223     sorted_results = sort_items_by_multiple_keys(
224         results,
225         {
226             # 0: True, # AVG Silhouette
227             # 1: True, # Min Silhouette
228             # 2: False, # Average distance
229             4: False, # Number of clusters
230             # 6: True, # Min common features per cluster
231         },
232         {
233             # 0: 1,

```

```

234         # 1: 1,
235         # 2: 1,
236         4: 1,
237         # 6: 1
238     }
239 )
240 # logging.debug(sorted_results)
241 best_result = results[sorted_results[0][0]]
242 # logging.debug(best_result)
243
244 best_method = best_result[-1]
245 best_silhouette = best_result[0]
246 best_labels = best_result[3]
247 global firstpass
248 if firstpass:
249     no_clusters = best_result[-1]
250     firstpass = False
251 else:
252     Nno_clusters = best_result[-1]
253
254     # no_clusters = best_result[-1]
255
256     logging.info("Best clustering method: {0} (adjusted silhouette ==
257                  {1})".format(best_method, best_silhouette))
258     return best_labels
259
260 else:
261     # Unknown strategy
262     raise NotImplementedError()

```

---

## A.6 Gap Statistic Implementation

The following segment of code is the defined function 'optimalK' which calculates the optimal value for K in K-means via the Gap Statistic method. This implementation is within the optimal\_k\_k\_means.py class and is thoroughly commented. The parameters and returned value are explained at the top of the function.

---

```

1
2 def optimalK(data, nrefs, maxClusters):
3     """
4     Calculates KMeans optimal K using Gap Statistic from
5     http://web.stanford.edu/~hastie/Papers/gap.pdf
6     Params:
7     data: ndarray of shape (n_samples, n_features)
8     nrefs: number of sample reference datasets to create

```

```

8         maxClusters: Maximum number of clusters to test for
9 Returns: (gaps, optimalK)
10 """
11 gaps = np.zeros((len(range(1, maxClusters)),))
12 resultsdf = pd.DataFrame({'clusterCount': [], 'gap': []})
13 for gap_index, k in enumerate(range(1, maxClusters)):
14
15     # Holder for reference dispersion results
16     refDisps = np.zeros(nrefs)
17
18     # For n references, generate random sample and perform kmeans
19     # getting resulting dispersion of each loop
20     for i in range(nrefs):
21         # Create new random reference set
22         randomReference = np.random.random_sample(size=data.shape)
23
24         # Fit to it
25         km = KMeans(k)
26         km.fit(randomReference)
27
28         refDisp = km.inertia_
29         refDisps[i] = refDisp
30
31     # Fit cluster to original data and create dispersion
32     km = KMeans(k)
33     km.fit(data)
34
35     origDisp = km.inertia_
36
37     # Calculate gap statistic
38     gap = np.log(np.mean(refDisps)) - np.log(origDisp)
39
40     # Assign this loop's gap statistic to gaps
41     gaps[gap_index] = gap
42
43     resultsdf = resultsdf.append({'clusterCount': k, 'gap': gap},
44                                  ignore_index=True)
45
46 return (gaps.argmax() + 1, resultsdf) # Plus 1 because index of 0
47 means 1 cluster is optimal, index 2 = 3 clusters are optimal

```

---

# Appendix B

## Example Application Output

### B.1 Fictional Small Network Output

This output was generated by running the application in automatic, maximum verbosity, dual input with graph plotting mode. The command used to execute the application with this configuration is as follows:

---

```
1 cluster.py -s automatic -vv -p -t -N -tp ../cw.xml ../cw.nessus
```

---

The network in which the scans were taken place is an entirely fictional and uses four machines in the scope. This scope is described below.

Table B.1: Fictional Network Scope and Machine Descriptions

	Windows 2008 server running Apache 2 web server with MySQL database. Hosting a DNS server.
192.168.0.1	Domain Controller for UADTARGETNET domain running Lightweight Directory Access Protocol NETBIOS name: SERVER1
	Windows 2008 server running empty web server and hosting a DNS server running Lightweight Directory Access Protocol NETBIOS name: SERVER2
192.168.0.10	Windows 7 Professional 7600 (Windows 7 Professional 6.1) NETBIOS name: CLIENT1
192.168.0.11	Windows 7 Professional 7600 (Windows 7 Professional 6.1) NETBIOS name: CLIENT2

#### B.1.1 Text Output

---

```

1 04-19 01:32 19624 cluster <module> DEBUG :twin flag enabled
2 04-19 01:32 19624 cluster <module> DEBUG :tp: ../cw.xml , path: ['../cw.nessus']
3 04-19 01:32 19624 cluster <module> INFO :Vectorizing Stage for Nessus
4 04-19 01:32 19624 parse_nessus parse_input INFO :Parsing Nessus XML * BETA *
5 no of IP's taken from nessus: 4
6 04-19 01:32 19624 parse_nessus parse_input INFO :Done Nessus parsing
7 04-19 01:32 19624 cluster <module> DEBUG :Loaded 4 vectors with 45 features
8 04-19 01:32 19624 cluster <module> INFO :Vectorizing complete
9
10 04-19 01:32 19624 cluster <module> INFO :Vectorizing Stage for nmap
11 no of IP's taken from NMAP: 4
12 04-19 01:32 19624 cluster <module> DEBUG :Loaded 4 vectors with 81 features
13 04-19 01:32 19624 cluster <module> INFO :Vectorizing complete
14
15 04-19 01:32 19624 cluster <module> INFO :Normalising the nessus vectors
16 04-19 01:32 19624 cluster <module> INFO :Reducing vectors to two dimensions with PCA
17 04-19 01:32 19624 cluster <module> DEBUG :reduced to 4 vectors with 2 dimensions
18 04-19 01:32 19624 cluster <module> INFO :Normalising complete
19
20 04-19 01:32 19624 cluster <module> INFO :Normalising the nmap vectors
21 04-19 01:32 19624 cluster <module> INFO :Reducing vectors to two dimensions with PCA
22 04-19 01:32 19624 cluster <module> DEBUG :reduced to 4 vectors with 2 dimensions
23 04-19 01:32 19624 cluster <module> INFO :Normalising complete
24
25 04-19 01:32 19624 cluster <module> INFO :Clustering Nessus
26 04-19 01:32 19624 cluster cluster DEBUG :Starting prospective KMeans clusterings
27 04-19 01:32 19624 cluster cluster DEBUG :Trying kmeans(n_clusters=2)
28 04-19 01:32 19624 cluster cluster DEBUG :Trying kmeans(n_clusters=3)
29 04-19 01:32 19624 cluster cluster INFO :Best clustering method: kmeans(n_clusters=2) (adjusted
    silhouette == 0.176059056707)
30 04-19 01:32 19624 cluster <module> INFO :Clustering Complete
31
32
33 04-19 01:32 19624 cluster <module> INFO :Clustering Nmap
34 04-19 01:32 19624 cluster cluster DEBUG :Starting prospective KMeans clusterings
35 04-19 01:32 19624 cluster cluster DEBUG :Trying kmeans(n_clusters=2)
36 04-19 01:32 19624 cluster cluster DEBUG :Trying kmeans(n_clusters=3)
37 04-19 01:33 19624 cluster cluster INFO :Best clustering method: kmeans(n_clusters=2) (adjusted
    silhouette == 0.441640409673)
38 04-19 01:33 19624 cluster <module> INFO :Clustering Complete
39
40
41 Note: shared features does not retain keys from XML and therefore wont always be human readable.
42 Printing Nessus cluster details
43
44 Cluster ID: 0
45 Silhouette Score: 0.299967117317
46 IPs: 192.168.0.10, 192.168.0.11
47
48 Shared Features:
49 general-purpose
50 windows
51 445
52 cpe:/o:microsoft:windows_7::professional
53 Mon Apr 03 11:25:23 2017
54 Microsoft Windows 7 Professional
55 9
56 5355
57
58
59
60 Cluster ID: 1
61 Silhouette Score: 0.0521509960965
62 IPs: 192.168.0.1, 192.168.0.2
63
64 Shared Features:
65 Mon Apr 03 11:25:06 2017

```

```

66     cpe:/o:microsoft:windows
67     general-purpose
68     windows
69     53
70     23
71     445
72
73
74
75     cluster ID : amount of IP's
76     0 : 2
77     1 : 2
78
79
80
81     Printing Nmap cluster details
82
83     Cluster ID: 0
84     Silhouette Score: 0.213624573736
85     IPs: 192.168.0.1, 192.168.0.2
86
87     Shared Features:
88     tcp23open telnet
89     tcp42open tcpwrapped
90     tcp53open domain
91     tcp80open http
92     tcp88open kerberos-sec
93     tcp135open
94     tcp139open
95     tcp389open
96     tcp445open
97     tcp464open
98     tcp593open
99     tcp636open tcpwrapped
100    tcp3268open
101    tcp3269open tcpwrapped
102    tcp49152open
103    tcp49153open
104    tcp49154open
105    tcp49155open
106    tcp49157open
107    tcp49158open
108
109
110
111    Cluster ID: 1
112    Silhouette Score: 0.66965624561
113    IPs: 192.168.0.10, 192.168.0.11
114
115    Shared Features:
116    tcp135open msrpcMicrosoft Windows RPC
117    tcp139open netbios-ssnMicrosoft Windows 98 netbios-ssn
118    tcp445open microsoft-dsMicrosoft Windows 10 microsoft-ds
119    tcp49152open msrpcMicrosoft Windows RPC
120    tcp49153open msrpcMicrosoft Windows RPC
121    tcp49154open msrpcMicrosoft Windows RPC
122    tcp49175open msrpcMicrosoft Windows RPC
123    tcp49176open msrpcMicrosoft Windows RPC
124
125
126
127    cluster ID : amount of IP's
128    0 : 2
129    1 : 2
130    04-19 01:33 19624 cluster <module> DEBUG :Getting centroids using reduced vectors for Nmap:
131    04-19 01:33 19624 cluster <module> DEBUG :nclusters: 2
132    04-19 01:33 19624 clustering get_centroids DEBUG :K = 0
133

```



```

134 04-19 01:33 19624 cluster <module> DEBUG :attempting to plot the following centroids:
135 [[-0.56736301 -0.01841586]
136 [ 0.56736301 0.01841586]]
137
138
139 04-19 01:33 19624 cluster <module> DEBUG :Getting centroids using reduced vectors for Nessus:
140 04-19 01:33 19624 cluster <module> DEBUG :nclusters: kmeans(n_clusters=2)
141 04-19 01:33 19624 cluster <module> DEBUG :nclusters: 2
142 04-19 01:33 19624 clustering get_centroids DEBUG :K = 0
143
144 04-19 01:33 19624 cluster <module> DEBUG :attempting to plot the following centroids:
145 [[-0.49416507 -0.0089938 ]
146 [ 0.49416507 0.0089938 ]]
147
148
149 04-19 01:33 19624 cluster <module> INFO :Nmap Centroids Covariance Matrix:
150 [[ 0.64380156 0.02089695]
151 [ 0.02089695 0.00067829]]
152 04-19 01:33 19624 cluster <module> INFO :Nessus Centroids Covariance Matrix:
153 [[ 4.88398234e-01 8.88884549e-03]
154 [ 8.88884549e-03 1.61776945e-04]]
155 04-19 01:33 19624 cluster <module> INFO :distance matrix between centroids using metric for Nmap:
156 euclidean :
157 -----
158 0          1.13532
159 1.13532 0
160 -----
161 04-19 01:33 19624 cluster <module> INFO :distance matrix between centroids using metric for Nessus:
162 euclidean :
163 -----
164 0          0.988494
165 0.988494 0
166 -----
167 04-19 01:33 19624 cluster <module> INFO :IP's from clusters with less than 3 IP's:
168 ['192.168.0.1' '192.168.0.10' '192.168.0.11' '192.168.0.2']
169 04-19 01:33 19624 cluster <module> DEBUG :Loaded 4 vectors with 126 features
170 04-19 01:33 19624 cluster <module> INFO :Normalizing input and reducing vectors to two dimensions
171 with PCA
172 04-19 01:33 19624 cluster <module> INFO :Resulting single IP vectors:
173 [[-0.50300301 -0.49063882]
174 [ 0.52888708 -0.00269094]
175 [ 0.56066029 0.03834554]
176 [-0.58654436 0.45498422]]
177 04-19 01:33 19624 cluster <module> INFO :distance matrix between centroids of small combined
178 clusters: euclidean :
179 -----
180 0          1.14144    1.18794    0.949306
181 1.14144 0          0.0518992    1.20568
182 1.18794 0.0518992 0          1.22052
183 0.949306 1.20568    1.22052    0
184 -----
185 04-19 01:33 19624 clustering cluster_single_kmeans INFO :Calculating gap statistic value, this can
186 take a while...
187
188 04-19 01:33 19624 clustering cluster_single_kmeans INFO :gap statistics recommends number of
189 clusters: 3
190
191 04-19 01:33 19624 clustering cluster_single_kmeans DEBUG :No K value specified, using Gap Statistic
192
193 04-19 01:33 19624 cluster <module> INFO :Writing recommended attack IP's to targets.txt for
194 exploitation
195 {0}
196 04-19 01:33 19624 display twin DEBUG :twin cluster plot.
197 04-19 01:33 19624 single_ip_parsing_nmap parse_input DEBUG :found ip 192.168.0.1
198 04-19 01:33 19624 single_ip_parsing_nmap parse_input DEBUG :found ip 192.168.0.2
199 04-19 01:33 19624 single_ip_parsing_nmap parse_input DEBUG :found ip 192.168.0.10
200 04-19 01:33 19624 single_ip_parsing_nmap parse_input DEBUG :found ip 192.168.0.11

```

```

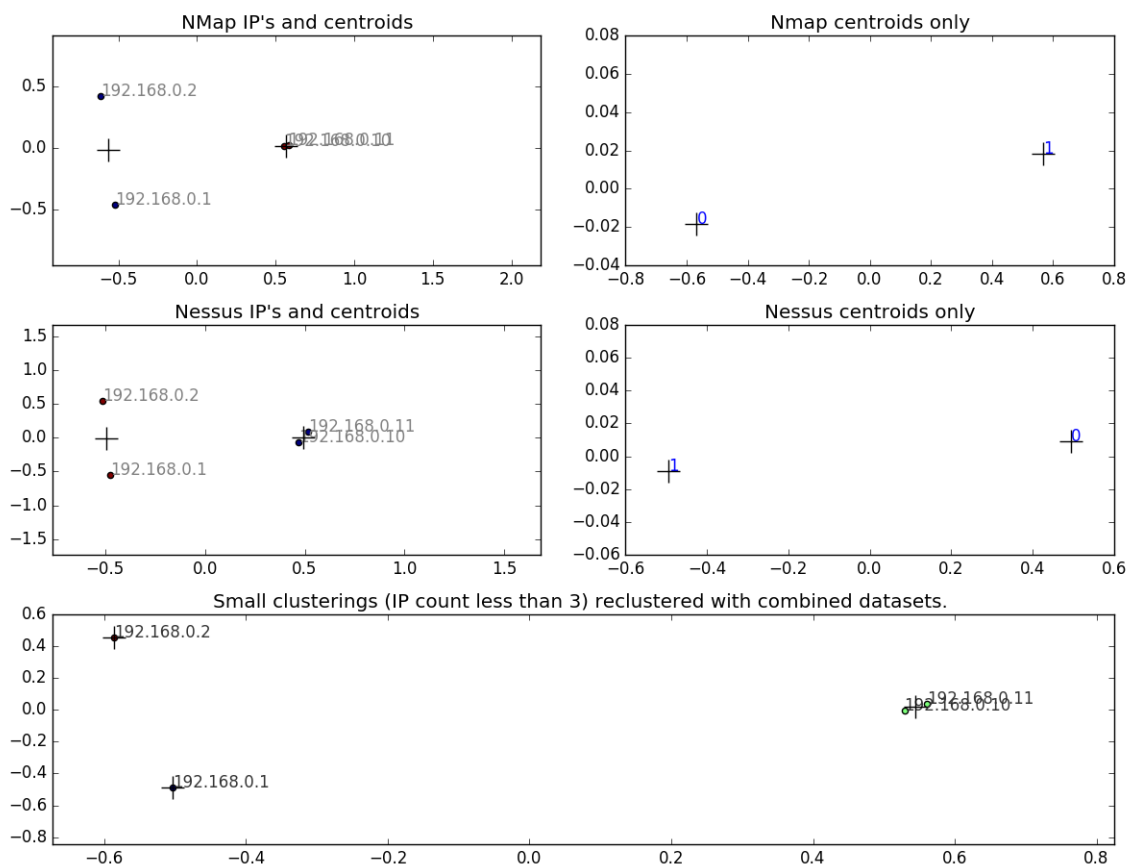
194 04-19 01:33 19624 display twin DEBUG :{'192.168.0.2': ['"Microsoft Windows 7 SP0 - SP1, Windows
Server 2008 SP1, Windows 8, or Windows 8.1 Update 1. Prediction accuracy: 100"'], '192.168.0.1':
[], '192.168.0.10': ['"Microsoft Windows 7 SP0 - SP1, Windows Server 2008 SP1, Windows 8, or
Windows 8.1 Update 1. Prediction accuracy: 100"'], '192.168.0.11': ['"Microsoft Windows 7 SP0 -
SP1, Windows Server 2008 SP1, Windows 8, or Windows 8.1 Update 1. Prediction accuracy: 100"']}

```

---

## B.1.2 Figures

The Figure B.1 below shows the example graphing interface output using the configuration mentioned previously in Appendix B.1.



Recommended attack vectors:  
192.168.0.2 : "Microsoft Windows 7 SP0 - SP1, Windows Server 2008 SP1, Windows 8, or Windows 8.1 Update 1. Prediction accuracy: 100"  
192.168.0.1 : OS detection failed, Didn't receive UDP response  
192.168.0.10 : "Microsoft Windows 7 SP0 - SP1, Windows Server 2008 SP1, Windows 8, or Windows 8.1 Update 1. Prediction accuracy: 100"  
192.168.0.11 : "Microsoft Windows 7 SP0 - SP1, Windows Server 2008 SP1, Windows 8, or Windows 8.1 Update 1. Prediction accuracy: 100"

Figure B.1: Example of graphing interface GUI from a fictional network to be used in conjunction with text output at B.1.1

## B.2 Hacklab Analysis Results

The following results contain outputs of the application created in this thesis when using the University Hacklab network as a dataset for scanning. The scans were obtained after permission was given on the 21st of March 2017. The IP addresses included within the scope is the entire /24 subnet with a mask of 255.255.255.0.

The nmap command used to obtain the scan was as follows:

---

```
1 nmap -A -O -oX hacklab.xml 10.0.0.0/24
```

---

The nessus configuration used was the default policy on an advanced scan. The target IP address configuration of 10.0.0.0/24 was used.

### B.2.1 Text Output

The following command was issued to the application in order to achieve the output shown below. It contains the application output in very verbose dual input mode using the hacklab dataset.

---

```
1 cluster.py -s automatic -vv -p -t -N -tp "../hacklab
  analyses/hacklab_new.xml" "../hacklab analyses/hacklab_new.nessus"
```

---

```
1 04-25 02:51 3220 cluster <module> DEBUG :twin flag enabled
2 04-25 02:51 3220 cluster <module> DEBUG :tp: ../hacklab analyses/hacklab_new.xml , path: ['../hacklab
  analyses/hacklab_new.nessus']
3 04-25 02:51 3220 cluster <module> INFO :Vectorizing Stage for Nessus
4 04-25 02:51 3220 parse_nessus parse_input INFO :Parsing Nessus XML * BETA *
5 04-25 02:51 3220 parse_nessus parse_input INFO :Done Nessus parsing
6 no of IP's taken from nessus: 46
7 04-25 02:51 3220 cluster <module> DEBUG :Loaded 46 vectors with 288 features
8 04-25 02:51 3220 cluster <module> INFO :Vectorizing complete
9 04-25 02:51 3220 cluster <module> INFO :Vectorizing Stage for nmap
10 no of IP's taken from NMAP: 47
11 04-25 02:51 3220 cluster <module> DEBUG :Loaded 42 vectors with 203 features
12 04-25 02:51 3220 cluster <module> INFO :Vectorizing complete
13 04-25 02:51 3220 cluster <module> INFO :Normalising the nessus vectors
14 04-25 02:51 3220 cluster <module> INFO :Reducing vectors to two dimensions with PCA
15 04-25 02:51 3220 cluster <module> DEBUG :reduced to 46 vectors with 2 dimensions
16 04-25 02:51 3220 cluster <module> INFO :Normalising complete
17 04-25 02:51 3220 cluster <module> INFO :Normalising the nmap vectors
18 04-25 02:51 3220 cluster <module> INFO :Reducing vectors to two dimensions with PCA
19 04-25 02:51 3220 cluster <module> DEBUG :reduced to 42 vectors with 2 dimensions
20 04-25 02:51 3220 cluster <module> INFO :Normalising complete
21 04-25 02:51 3220 cluster <module> INFO :Clustering Nessus
22 04-25 02:51 3220 cluster cluster DEBUG :Starting prospective KMeans clusterings
23 04-25 02:51 3220 cluster cluster DEBUG :Trying kmeans(n_clusters=2)
24 04-25 02:52 3220 cluster cluster DEBUG :Not all clusters are informative (a cluster has 0 shared
  features)
25 04-25 02:52 3220 cluster cluster DEBUG :Trying kmeans(n_clusters=3)
26 04-25 02:52 3220 cluster cluster DEBUG :Not all clusters are informative (a cluster has 0 shared
  features)
27 04-25 02:52 3220 cluster cluster DEBUG :Trying kmeans(n_clusters=4)
28 04-25 02:52 3220 cluster cluster DEBUG :Not all clusters are informative (a cluster has 0 shared
  features)
```

---

```

29 04-25 02:52 3220 cluster cluster DEBUG :Trying kmeans(n_clusters=5)
30 04-25 02:52 3220 cluster cluster DEBUG :Not all clusters are informative (a cluster has 0 shared
    features)
31 04-25 02:52 3220 cluster cluster DEBUG :Trying kmeans(n_clusters=6)
32 04-25 02:52 3220 cluster cluster DEBUG :Trying kmeans(n_clusters=7)
33 04-25 02:53 3220 cluster cluster INFO :Best clustering method: kmeans(n_clusters=6) (adjusted
    silhouette == 0.281637496409)
34 04-25 02:53 3220 cluster <module> INFO :Clustering Complete
35 04-25 02:53 3220 cluster <module> INFO :Clustering Nmap
36 04-25 02:53 3220 cluster cluster DEBUG :Starting prospective KMeans clusterings
37 04-25 02:53 3220 cluster cluster DEBUG :Trying kmeans(n_clusters=2)
38 04-25 02:53 3220 cluster cluster DEBUG :Not all clusters are informative (a cluster has 0 shared
    features)
39 04-25 02:53 3220 cluster cluster DEBUG :Trying kmeans(n_clusters=3)
40 04-25 02:53 3220 cluster cluster DEBUG :Not all clusters are informative (a cluster has 0 shared
    features)
41 04-25 02:53 3220 cluster cluster DEBUG :Trying kmeans(n_clusters=4)
42 04-25 02:53 3220 cluster cluster DEBUG :Not all clusters are informative (a cluster has 0 shared
    features)
43 04-25 02:53 3220 cluster cluster DEBUG :Trying kmeans(n_clusters=5)
44 04-25 02:53 3220 cluster cluster DEBUG :Trying kmeans(n_clusters=6)
45 04-25 02:53 3220 cluster cluster INFO :Best clustering method: kmeans(n_clusters=5) (adjusted
    silhouette == 0.411583679893)
46 04-25 02:53 3220 cluster <module> INFO :Clustering Complete
47 Note: shared features does not retain keys from XML and therefore wont always be human readable.
48 Printing Nessus cluster details
49 Cluster ID: 0
50 Silhouette Score: 0.25819092251
51 IPs: 10.0.0.28, 10.0.0.34, 10.0.0.35, 10.0.0.69, 10.0.0.74, 10.0.0.78, 10.0.0.80, 10.0.0.85
52 Shared Features:
53 general-purpose
54 cpe:/o:linux:linux_kernel:3.10
55 cpe:/o:linux:linux_kernel:3.13
56 cpe:/o:linux:linux_kernel:4.2
57 cpe:/o:linux:linux_kernel:4.8
58 cpe:/a:openbsd:openssh:7.2
59 SSH-2.0-OpenSSH_7.2
60 Linux Kernel 3.10\\nLinux Kernel 3.13\\nLinux Kernel 4.2\\nLinux Kernel 4.8
61 1
62 linux
63
64
65 Cluster ID: 1
66 Silhouette Score: 0.220046730181
67 IPs: 10.0.0.24, 10.0.0.26, 10.0.0.27, 10.0.0.30, 10.0.0.33, 10.0.0.36, 10.0.0.39, 10.0.0.40,
    10.0.0.41, 10.0.0.42, 10.0.0.43, 10.0.0.46, 10.0.0.47, 10.0.0.48, 10.0.0.50, 10.0.0.51,
    10.0.0.53, 10.0.0.54, 10.0.0.55, 10.0.0.56, 10.0.0.58, 10.0.0.64, 10.0.0.65, 10.0.0.66,
    10.0.0.70, 10.0.0.76, 10.0.0.79, 10.0.0.83, 10.0.0.84
68 Shared Features:
69 cpe:/o:microsoft:windows_7::professional
70 cpe:/a:openbsd:openssh:7.1 -> OpenBSD OpenSSH 7.1
71 SSH-2.0-OpenSSH_7.1p1 Microsoft_Win32_port_with_VS
72 general-purpose
73 Microsoft Windows 7 Professional
74 windows
75 445
76
77
78 Cluster ID: 2
79 Silhouette Score: 0.18063333287
80 IPs: 10.0.0.32, 10.0.0.37, 10.0.0.77, 10.0.0.81
81 Shared Features:
82 1
83
84
85 Cluster ID: 3
86 Silhouette Score: 0.0374285781879
87 IPs: 10.0.0.185, 10.0.0.3

```

```

88     Shared Features:
89     general-purpose
90     cpe:/a:openbsd:openssh:7.2
91     cpe:/o:linux:linux_kernel
92     linux
93     cpe:/o:canonical:ubuntu_linux:16.04
94     SSH-2.0-OpenSSH_7.2p2 Ubuntu-4ubuntu2.1
95     Linux Kernel 4.4 on Ubuntu 16.04 (xenial)
96
97
98     Cluster ID: 4
99     Silhouette Score: -0.00647458529864
100    IPs: 10.0.0.1, 10.0.0.7
101    Shared Features:
102    443
103    3
104
105
106    Cluster ID: 5
107    Silhouette Score: 1.0
108    IPs: 10.0.0.5
109    Shared Features:
110    cpe:/o:microsoft:windows_7::professional
111    general-purpose
112    Microsoft Windows 7 Professional
113    windows
114    445
115    1490105876
116    Hacklab2
117    Tue Mar 21 14:17:56 2017
118    Tue Mar 21 14:15:50 2017
119    HACKLAB2
120    10.0.0.5
121    9
122    00:25:b3:28:d9:14
123    49158
124    5355
125
126
127    cluster ID : amount of IP's
128    0 : 8
129    1 : 29
130    2 : 4
131    3 : 2
132    4 : 2
133    5 : 1
134    Printing Nmap cluster details
135    Cluster ID: 0
136    Silhouette Score: 0.528486221388
137    IPs: 10.0.0.24, 10.0.0.26, 10.0.0.27, 10.0.0.30, 10.0.0.33, 10.0.0.36, 10.0.0.39, 10.0.0.41,
        10.0.0.42, 10.0.0.43, 10.0.0.46, 10.0.0.47, 10.0.0.48, 10.0.0.50, 10.0.0.51, 10.0.0.53,
        10.0.0.54, 10.0.0.55, 10.0.0.56, 10.0.0.58, 10.0.0.64, 10.0.0.65, 10.0.0.66, 10.0.0.70,
        10.0.0.76, 10.0.0.79, 10.0.0.83, 10.0.0.84
138    Shared Features:
139    tcp22opensshprotocol 2.0
140    tcp135openmsrpcMicrosoft Windows RPC
141    tcp139opennetbios-ssn
142    tcp443openhttpVMware VirtualCenter Web service
143    tcp443openhttpssl-certlocalityNamePalo Alto
144    tcp443openhttpssl-certorganizationalUnitNameVMware
145    tcp443openhttpssl-certcommonNameVMware
146    tcp443openhttpssl-certemailAddressnone@vmware.com
147    tcp443openhttpssl-certcountryNameUS
148    tcp443openhttpssl-certtypersa
149    tcp443openhttpssl-certbits2048
150    tcp443openhttpssl-certnotAfter2017-08-03T11:59:23+00:00
151    tcp443openhttpssl-certnotBefore2016-08-03T11:59:23+00:00
152    tcp443openhttpssl-certmd5ad6e5f01314d1518b4051099cd81a1f0

```

```

153     tcp443openhttpssl-certsha183a0d223f33891ba5192d94c26c44e05cfe72a51
154     tcp443openhttpssl-certpem-----BEGIN
155         CERTIFICATE-----\nMIIIEjCCAvqgAwIBAgIJAP9VV+H40SLcMAOGCSqGSIb3DQEBCwUAMGMxCzAJBgNV\\nBAYTA1VTMRIwEAYDVQQHDA1Q
156         CERTIFICATE-----\n
157     tcp445opennetbios-ssn
158     Cluster ID: 1
159     Silhouette Score: 1.0
160     IPs: 10.0.0.28, 10.0.0.34, 10.0.0.35, 10.0.0.40, 10.0.0.69, 10.0.0.74, 10.0.0.78, 10.0.0.80, 10.0.0.85
161     Shared Features:
162     tcp22opensshOpenSSH7.2protocol 2.0
163
164     Cluster ID: 2
165     Silhouette Score: -0.222222631321
166     IPs: 10.0.0.1, 10.0.0.7
167     Shared Features:
168     tcp443open
169     tcp80openhttp
170
171     Cluster ID: 3
172     Silhouette Score: -0.248345190601
173     IPs: 10.0.0.3, 10.0.0.5
174     Shared Features:
175     tcp139opennetbios-ssn
176     tcp445opennetbios-ssn
177
178     Cluster ID: 4
179     Silhouette Score: 1.0
180     IPs: 10.0.0.185
181     Shared Features:
182     tcp22opensshprotocol
183         2.0SF-Port22-TCP:V=6.47%I=7%D=3/21%Time=58D136BB%P=i686-pc-windows-windows%r(NULL,29,"SSH-2\\\0-OpenSSH_7\\\
184
185     cluster ID : amount of IP's
186     0 : 28
187     1 : 9
188     2 : 2
189     3 : 2
190     4 : 1
191     04-25 02:53 3220 cluster <module> DEBUG :Getting centroids using reduced vectors for Nmap:
192     04-25 02:53 3220 cluster <module> DEBUG :nclusters: 5
193     04-25 02:53 3220 cluster <module> DEBUG :attempting to plot the following centroids:
194     [[-0.34575755 0.05203286]
195     [ 0.93722406 0.14279432]
196     [ 0.31235087 -0.80310291]
197     [ 0.49977173 -0.23454181]
198     [ 0.06086076 -0.45066068]]
199     04-25 02:53 3220 cluster <module> DEBUG :Getting centroids using reduced vectors for Nessus:
200     04-25 02:53 3220 cluster <module> DEBUG :nclusters: kmeans(n_clusters=5)
201     04-25 02:53 3220 cluster <module> DEBUG :nclusters: 6
202     04-25 02:53 3220 cluster <module> DEBUG :attempting to plot the following centroids:
203     [[ 0.75587225 0.27174194]
204     [-0.31269697 0.04260005]
205     [ 0.43641262 -0.52606411]
206     [ 0.41356214 -0.09467375]
207     [ 0.2518518 -0.51866881]
208     [-0.05524435 -0.07839544]]
209     04-25 02:53 3220 cluster <module> INFO :Nmap Centroids Covariance Matrix:
210     [[ 0.23001336 0.02479788]
211     [ 0.02479788 0.14788991]]
212     04-25 02:53 3220 cluster <module> INFO :Nessus Centroids Covariance Matrix:
213     [[ 0.14543938 0.00427455]
214     [ 0.00427455 0.10009763]]

```

```

218 04-25 02:53 3220 cluster <module> INFO :distance matrix between centroids using metric for Nmap:
      euclidean :
219 -----
220 0          1.28619  1.07906  0.892774 0.64656
221 1.28619  0          1.13366  0.577708 1.0584
222 1.07906  1.13366  0          0.598655 0.43297
223 0.892774 0.577708 0.598655 0          0.489234
224 0.64656  1.0584   0.43297  0.489234 0
225 -----
226 04-25 02:53 3220 cluster <module> INFO :distance matrix between centroids using metric for Nessus:
      euclidean :
227 -----
228 0          1.09286  0.859389 0.501435 0.937436 0.883463
229 1.09286  0          0.940502 0.739119 0.796077 0.284468
230 0.859389 0.940502 0          0.431995 0.184709 0.664931
231 0.501435 0.739119 0.431995 0          0.453786 0.469089
232 0.937436 0.796077 0.184709 0.453786 0          0.536795
233 0.883463 0.284468 0.664931 0.469089 0.536795 0
234 -----
235 04-25 02:53 3220 cluster <module> INFO :IP's from clusters with less than 3 IP's:
236 ['10.0.0.1' '10.0.0.185' '10.0.0.3' '10.0.0.5' '10.0.0.7']
237 04-25 02:53 3220 cluster <module> DEBUG :Loaded 5 vectors with 491 features
238 04-25 02:53 3220 cluster <module> INFO :Normalizing input and reducing vectors to two dimensions with
      PCA
239 04-25 02:53 3220 cluster <module> INFO :Resulting single IP vectors:
240 [[-0.65409661 0.13743411]
241 [ 0.47045184 -0.50265125]
242 [ 0.24966134 -0.26531969]
243 [ 0.39487565 0.79836834]
244 [-0.46089222 -0.16783152]]
245 04-25 02:53 3220 cluster <module> INFO :distance matrix between centroids of small combined clusters:
      euclidean :
246 -----
247 0          1.29395  0.989439 1.23983 0.361269
248 1.29395  0          0.324152 1.30321 0.9897
249 0.989439 0.324152 0          1.07355 0.71721
250 1.23983  1.30321  1.07355  0          1.29069
251 0.361269 0.9897   0.71721  1.29069 0
252 -----
253 04-25 02:53 3220 clustering cluster_single_kmeans INFO :Calculating gap statistic value, this can
      take a while...
254 04-25 02:53 3220 clustering cluster_single_kmeans INFO :gap statistics recommends number of clusters:
      3
255 04-25 02:53 3220 clustering cluster_single_kmeans DEBUG :No K value specified, using Gap Statistic
256 04-25 02:53 3220 cluster <module> INFO :Writing recommended attack IP's to targets.txt for
      exploitation
257 04-25 02:53 3220 display twin DEBUG :twin cluster plot.

```

---

## B.2.2 Figures

Figure B.2 shows the application output in standard mode using the Nessus dataset from the hacklab. The command used is as follows:

---

```
1 cluster.py -s automatic -vv -p -N "../hacklab analyses/hacklab_new.nessus"
```

---

The Figure B.3 shows output to that similar of the Figure B.2 above however this used the Nmap dataset as appose to the Nessus one used in that figure. The command used to acheive this result is as follows:

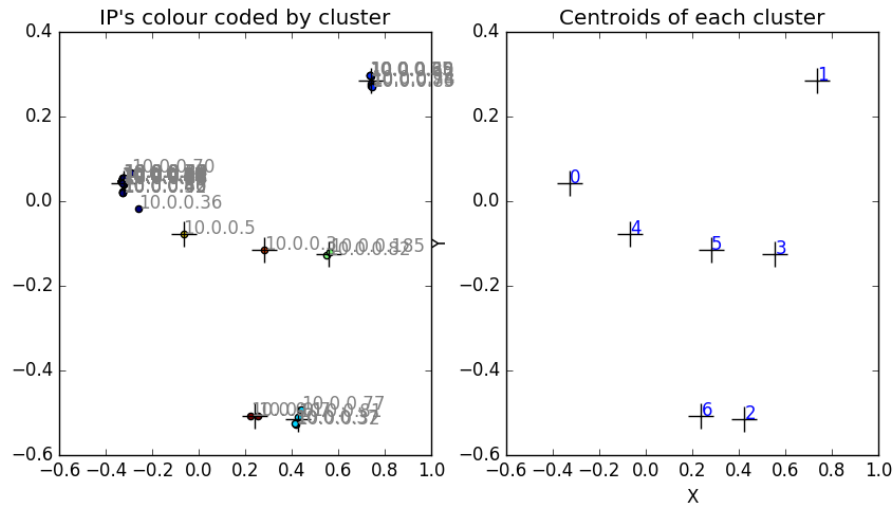


Figure B.2: Example graphing interface output using the the Abertay University Hacklab Nessus scan as a dataset.

---

```
1 cluster.py -s automatic -vv -p "../hacklab analyses/hacklab_new.xml"
```

---

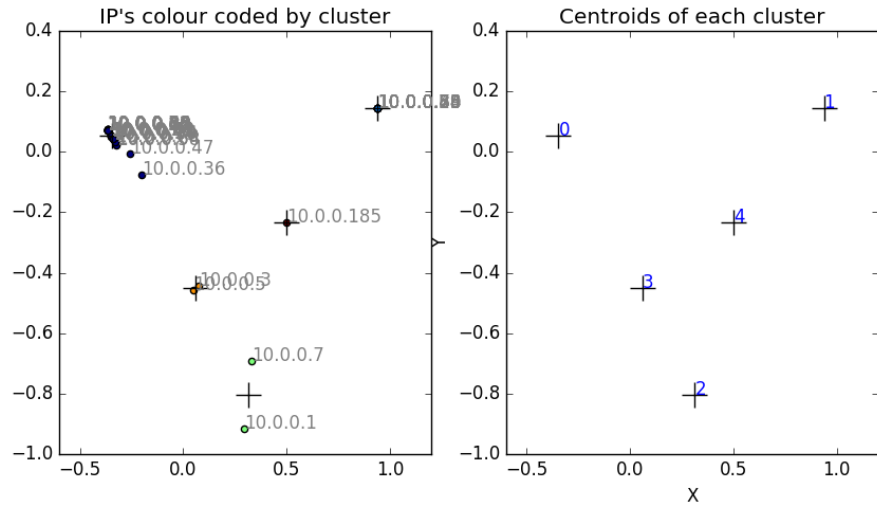
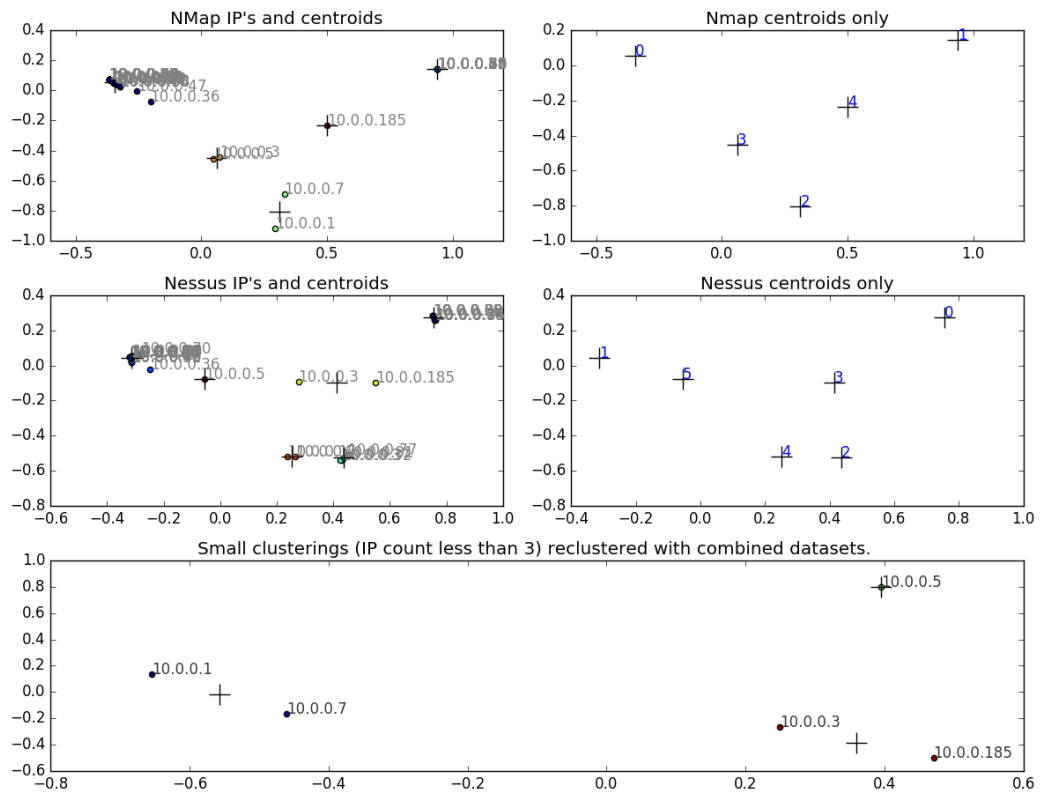


Figure B.3: Example graphing interface output using the the Abertay University Hacklab Nmap scan as a dataset.

Figure B.4 shows the graphing interface opened after the completion of the clustering algorithm. This was achieved using dual mode and the command use can be found above in B.2.1.





Recommended attack vectors:  
 10.0.0.5 : "Microsoft Windows 7 SP0 - SP1, Windows Server 2008 SP1, or Windows 8. Prediction accuracy: 100"  
 10.0.0.7 : "HP iLO 2 remote management interface. Prediction accuracy: 100"  
 10.0.0.185 : "Linux 3.11 - 3.14. Prediction accuracy: 100"  
 10.0.0.1 : "Linux 2.6.18 - 2.6.22. Prediction accuracy: 86"  
 10.0.0.3 : "Linux 3.11 - 3.14. Prediction accuracy: 100"

Figure B.4: Example graphing interface output using both scans of Abertay University Hacklab and running the application in dual mode.

Figures B.5 and B.6 below show both halves of the Nessus GUI interface overview. The same Nessus output file from the scan above in B.2 was used. The original representation was split into two Figures due to the size otherwise being too large for this document.

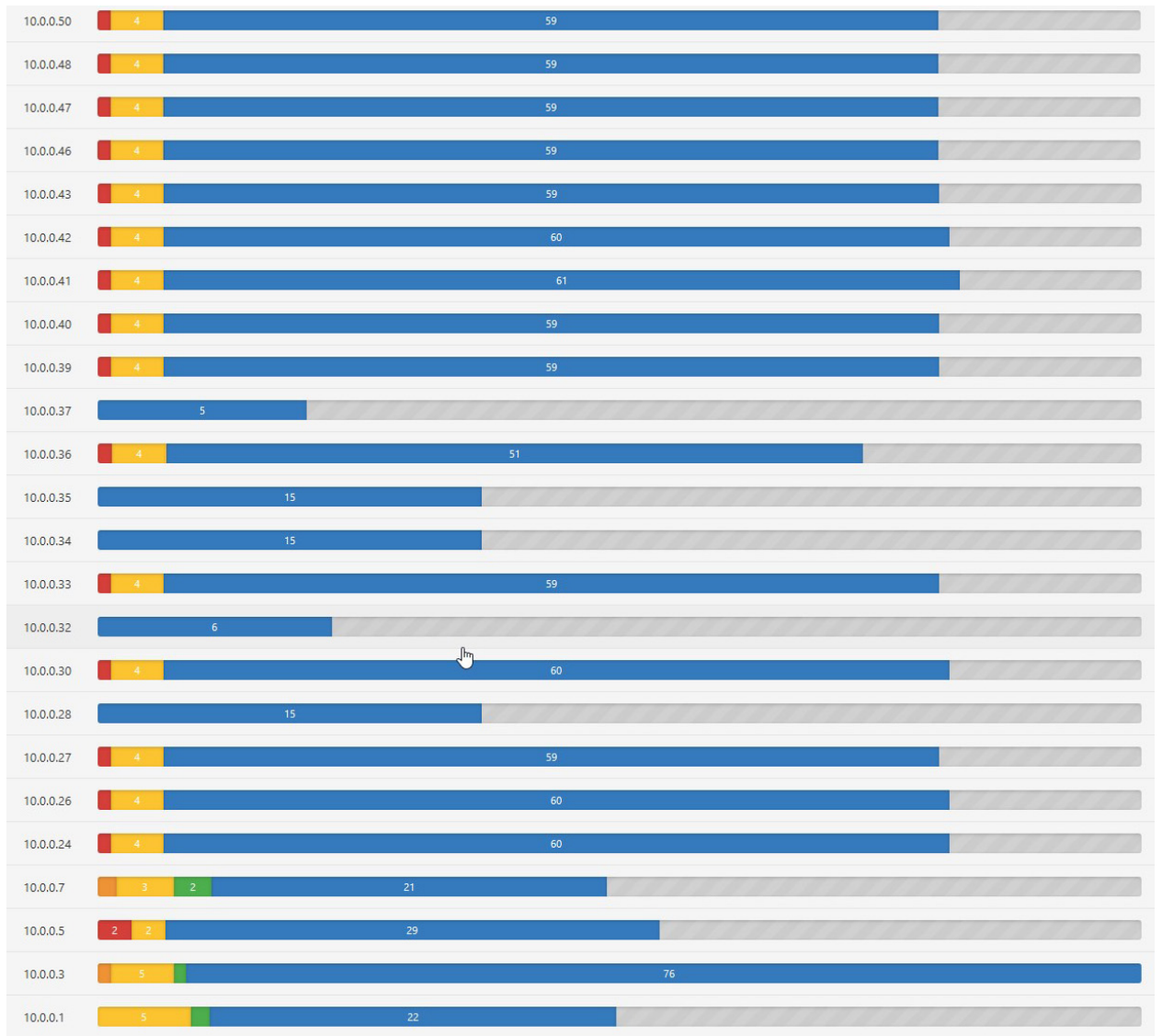


Figure B.5: Nessus interface overview of Abertay University Hacklab network using default policy (First Half).

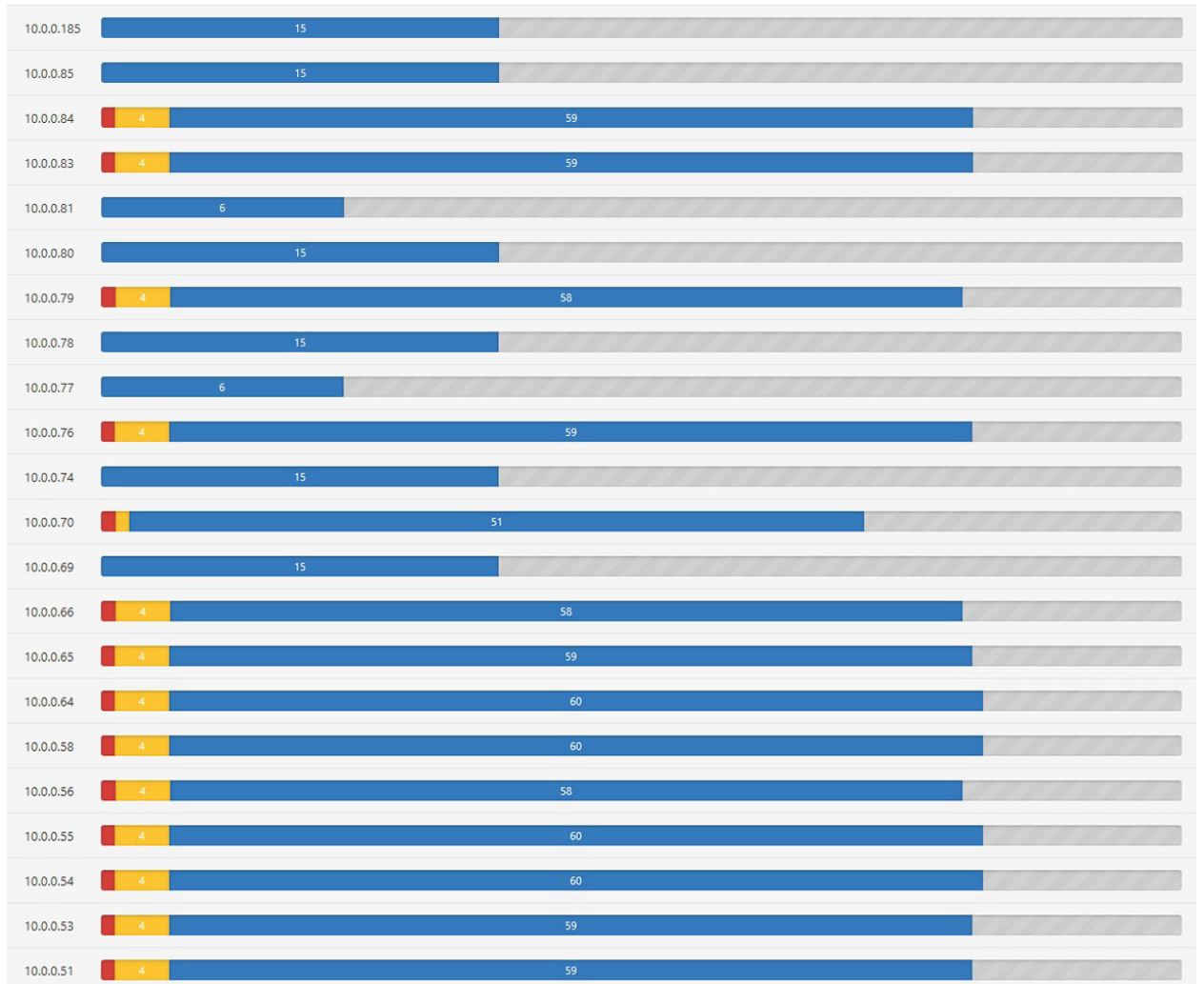


Figure B.6: Nessus interface overview of Abertay University Hacklab network using default policy (Second Half).

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