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DEPT. COMPUTER SCIENCE

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**Automatic Feature Selection for Website Fingerprinting**

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This report is submitted as part requirement for the BSc Degree in Computer Science at UCL. It is substantially the result of my own work except where explicitly indicated in the text. The report may be freely copied and distributed provided the source is explicitly acknowledged.

## Abstract

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# Chapter 1

## Introduction

### 1.1 The Problem

The internet has become an essential tool for communication for a majority of the population. But privacy has always remained a major concern, which is why nowadays most web content providers are slowly moving away from HTTP to HTTPS. For instance, at the time of writing, around 86% of Google’s traffic is encrypted. This is a significant improvement compared to 2014 where only 50% was sent over HTTPS [13]. However, this encryption only obscures the content of the web page and does not hide what websites you are visiting or in general who you might be communicating with.

Hence, an Internet Service Provider (ISP) can easily obtain a lot of information about a person. This is an especially large concern for people living in oppressive regimes since it allows a government to easily spy on its people and censor whatever websites they would like. To circumvent these issues, several anonymization techniques have been developed. These systems obscure both the content and meta-data, allowing a user to anonymously browse the web. One of the most popular low-latency anonymization networks is called Tor, which relies on a concept called Onion Routing [44].

The list of known attacks against Tor is at the time of writing very limited and most of them rely on very unlikely scenarios such as having access to specific parts of the network (*both entry and exit nodes*) [44]. However, for this report we will make a more reasonable assumption that an attacker is a *local eavesdropper*. By this we mean that the entity only has access to the traffic between the sender and the first anonymization node, like ISPs.

One of the most successful attacks that satisfies these conditions is known as *website fingerprinting* (WFP). It relies on the fact that Tor does not significantly alter the shape of the network traffic [16]. Hence, the attack infers information about the content by analysing the raw traffic. For instance by analysing the packet sizes, the amount of packets and the direction of the traffic, we might be able to deduce which web pages certain users are visiting. Initially, Tor was considered to be secure against this threat but around 2011, some techniques such as the *support vector classifier* (SVC) used by Panchenko et al. started to get recognition rates higher than 50% [31].

However, one of the main problems with majority of the attacks proposed in the research literature, is that they rely on a laborious, time-consuming manual feature engineering processes since most primitive machine learning techniques are only able to process fixed-length vectors as its input. These features are most often picked based on intuition and trial and error processes. But there is no guarantee that these features are the most appropriate ones.

Thus the goal of this paper is to investigate the use of novel deep-learning techniques to automatically extract a fixed-length vector representation from a traffic trace that represents loading a web page. Next, we aim to use these features in existing attacks to see if our model successfully identified the appropriate features.

## 1.2 Aims and Goals

We can subdivide the project up into different aims, each with their own goals:

1. **Aim:** Critically review the effectiveness of current website fingerprinting attacks.

**Goals:**

- Analyse various models that are currently used in fingerprinting attacks.
- Examine how would a small percentage of false positives impacts a potential attack.
- Analyse how the rapid changing nature of some web pages would impact the attack.
- Review if there are any assumptions that are being made that could impact the effectiveness of an attack.

2. **Aim:** Automatically generate features from a trace that represents loading a webpage.

**Goals:**

- Critically compare various different feature generation techniques such as stacked autoencoders, sequence-to-sequence model and bidirectional encoder-decoder models.
- Identify a dataset that is large enough to train a deep-learning model.
- Compare several software libraries to perform fast numerical computation such as Tensorflow, Torch, Theano and Keras.
- Implement the most appropriate feature generation model in one of the previously mentioned software libraries.

3. **Aim:** Train existing models with the automatically generated features and test their performance compared to hand-picked ones.

**Goals:**

- Identify several different models that have previously been successful in various website fingerprinting attacks and implement those models.
- Extract the same hand-picked features from our dataset as mentioned in the respective papers.
- Investigate an appropriate technique for evaluating the results of different models.
- Compare the hand-picked features compared to the automatically generated ones for different models. In addition, we also want to investigate their effectiveness in different threat models. For instance if an adversary wants to identify which specific web pages a user is visiting (*multi-class classification*) or if the adversary just wants to know whether the user visits a web page from a monitored set of web pages (*binary classification*).

## 1.3 Project Overview

As previously mentioned, the project can be split up into three different aims, which is why we also approach it in three different stages:

- First, we examine different existing website fingerprinting models to gain a deeper understanding of the concept.
- Next, we perform more research to contrast different automatic feature selection models and implement the most appropriate model.
- Finally, we compare the effectiveness of hand-picked features with the automatically generated ones by training them on different existing website fingerprinting models.

## 1.4 Report Structure

The general report has a very simple structure. In the following section we further explore similar works and several concepts that are necessary to understand the basics of website fingerprinting and our specific attack. Next, we identify the threat model and design an attack that performs an automatic features generation. Finally, we explore several methods of evaluating the performance of different models with different features and contrast the hand-picked features with the automatically generated ones.

## Chapter 2

# Background Information and Related Work

In the following chapter, we further explore the motivation for undertaking the project, analyse the current state of the project domain and outline the research that forms the basis for the rest of the report.

### 2.1 The Problem

As previously mentioned, the goal of this project is to automate the feature selection process for a website fingerprinting attack. By this we mean that given a specific trace, our model should be able to produce a fixed-length vector that is a close representation of the respective trace. However, before we delve into the details of the attack, we first need to gain a greater understanding of some concepts such as onion routing, website fingerprinting in general and deep learning.

#### 2.1.1 Onion Routing

To preserve privacy, we do not only need to obscure the content of a webpage but also hide who is taking to whom [12]. Tor achieves both of these by making use of a technique, called *onion routing*, which is a very simple protocol that can be divided up into three phases: connection setup, data movement and connection tear-down [12]. We show how it works by taking a simple example of Alice trying to communicate with Bob.

##### 1. Connection Setup:

- Alice's Tor client obtains a list of Tor nodes from a directory server.
- Then Alice picks three random Tor nodes and labels them *one*, *two* and *three*.
- Alice communicates with the first node and shares a symmetric key.
- Next, Alice sends messages to the first node, which are then forwarded to the second node to share another symmetric key to the second node.
- Finally, Alice continues sending messages to the first node, which are forwarded to the second and finally to the third node to share the final symmetric key. What is important here is that we use a secure key-sharing algorithm such that only Alice and the respective node know the keys. Additionally, since all of the traffic is forwarded from the first node, the second and the third nodes do not know the identity of Alice.



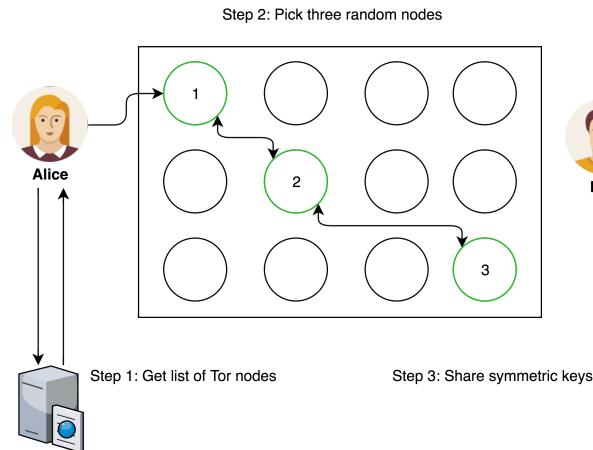


Figure 2.1.1: An example of a connection setup for the onion routing protocol.

## 2. Data Movement:

- Before Alice can send any data, it first needs to encrypt it in different layers. By this we mean that first it encrypts the data (*with Bob's address*) using the shared key from the third node. Next, it encrypts that data again (*with the address of the third node*) using the key from the second node. Finally, as expected, it encrypts the data a final time (*with the address of the second node*) using the shared key from the first node.
- Now Alice is ready to send the data to the first node.
- Once the first node received the data, it decrypts it using the shared key. This reveals the address to the second node. The key is here that the first node cannot see the data nor where it is going, since that is still encrypted.
- Next, the first node forwards the data that it just unencrypted to the second node. Again, this node decrypts the data, revealing the address to the third node but now it doesn't know what the data is, where the final destination is or where it originally came from.
- Lastly, the second node forwards the data to the third node. After encryption, this final node can see the data and where it is going but it does not know where it came from. So it forwards the data to Bob and not a single party should be able to know the data, the final destination and where it originally came from except for Alice and Bob.

Now we know why the protocol is called onion routing because it encrypts the data in multiple layers and at every node, one of the layers of the onion is peeled off [35]. The key is that none of the nodes know the complete path that has been taken.

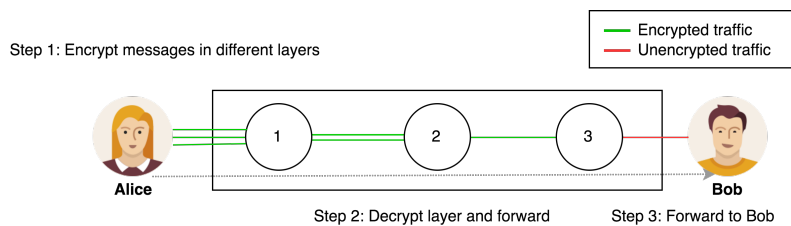


Figure 2.1.2: Sending a message with the onion routing protocol.

3. **Connection Tear-down:** This can be initiated by any of the nodes or the client and the process is very straightforward. Either the client sends a request for a tear-down to the first node to remove any data on the connection (including the shared key), which is then forwarded to the other nodes. Or one of the nodes sends a tear-down message to both the previous node and the next node, which are then forwarded in both directions [12].

Tor generally uses the same circuit for connections that happen within around 10 minutes after creating a circuit. Later requests, however, are given a new circuit [44, 35].

## 2.2 Related Work

### 2.2.1 Website Fingerprinting

Website fingerprinting (WF) is the process of attempting to identify which web pages a specific client visits by analysing their traffic traces. Hence, an attacker is considered to be *local*. By this we mean that they can eavesdrop on the traffic between the client and the first Tor node, as shown in figure 2.2.1. So it can be anyone from a person on the same network to an ISP. The reason as to why the attacker has to be local is because in onion routing systems, it is the only place in the network where you still know the true identity of the client.

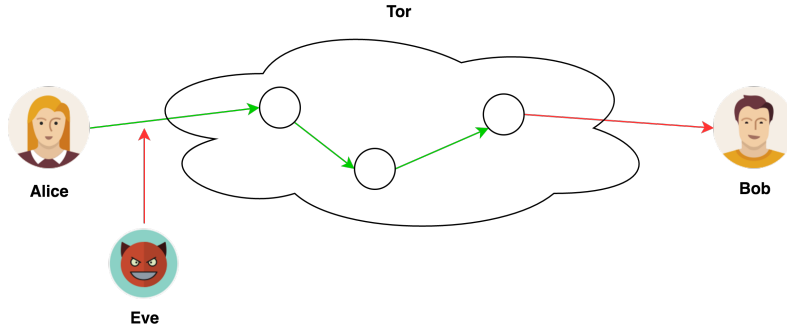


Figure 2.2.1: Threat model for a simple website fingerprinting attack.

In general, the attack is as follows. The attacker first collects a dataset that contains packet traces for loading several web pages. In practice, these web pages are the ones that an attacker is interested in monitoring. From those traces, the attacker extracts a fixed number of features, the so-called *fingerprint*. Next, it uses those fingerprints to train some sort of model (*often a machine learning model*) to find patterns in the data. The attacker observes packet traces, generated by the client, and uses the model to classify which web pages the user is visiting. Therefore, an attack is essentially a classification problem, where you try to map packet traces into web pages.

We should denote that throughout this report, we will be using the word “web pages” rather than “websites”. The reasoning behind this is that a website often consists of multiple HTML documents, which can result in significantly different traffic traces, depending on which document you load.

The term “*website fingerprinting*” was first mentioned by A. Hintz who performed a simple traffic analysis on Safeweb, an encrypting web proxy that tried to protect user’s privacy [18]. Although the attack was very simple, it and other earlier works show the possibility that encrypted channels might still leak information about the URL [18, 51]. Later, in 2009, the first fingerprinting attack against Tor was executed by Herrmann et al. using a *Multinomial Naive Bayes* model. They managed to get a relatively high accuracy

for several protocols, such as *OpenSSL* and *OpenVPN*, on a small set of 775 web pages. However on more specialised anonymization networks, the model only achieved a 2.96% for Tor, which they said was caused by a suboptimal configuration [17].

Around the same period Y. Shi et al. performed an attack that specifically focused on anonymity systems such as Tor [40]. Using a cosine similarity, they achieved around 50% accuracy on an even smaller dataset of only 20 web pages [40]. The first real threat however, was by A. Panchenko et al. who used a *Support Vector Classifier* (SVC) on the same dataset of 775 web pages as Herrmann et al. and got a 54.61% success rate [17, 31].

All of the previously mentioned research, except the one done by Panchenko et al. have only considered a *closed-world scenario*. A closed-world setting means that all of the web pages are known in advance [31]. For instance, when Herrmann et al. trained their model on a dataset of 775 web pages, they made the assumption that a client could only visit one of those web pages and none other. In an *open-world scenario*, the attacker does not know in advance which URLs the victim might visit. The most prominent example of this is when the authorities want to monitor which people try to access a set of censored sites [31]. In order to achieve this, the models need to be trained on both *monitored* and *unmonitored web pages*.

Wang et al. later conducted an attack on Tor using a large open-world dataset [53]. Using a novel *k-Nearest Neighbor* (k-NN) classifier with *weight adjustment* on a very large feature set (*3736 features*). In addition to getting around 90% accuracy, they also significantly reduced the time needed for training [53].

Using a completely different approach, Hayes et al. extract fingerprints using a *random forests* [16]. This novel technique involved storing a *fingerprint* as a set of leaves within that forest. Next, they simply use the *hamming distance* as a distance metric to find the k-nearest neighbors. If all the labels within those *k* instances are the same, the model classifies the new instance as the previously mentioned label. The interesting aspect is that changing the value of *k* allows them to vary the number of *true positives* (TP) and *false positives* (FP) [16].

Finally, Panchenko et al. improved upon their previous attack to create one current state-of-the-art methods. They tested their approach on several datasets, including the one used by Wang et al. on their k-NN attack, where they got around 92% accuracy. In an open-world scenario, on the other hand, they got up to 96% accuracy.

### 2.2.2 Automatic Feature Selection

There are not many works that have examine the use of automatic feature selection techniques in the context of a website fingerprinting attacks. First, Abe et al. study the use of a *stacked autoencoder* with a *softmax classifier* [1]. However, since a stacked autoencoder still requires a fixed-length input, they pad and truncate cells, whose length is shorter or longer than 5000. With it, they manage to achieve a 88% accuracy [1].

V. Rimmer takes a very similar approach, as she also uses a stacked autoencoder with a softmax classifier [37]. But rather than padding and truncating the cells, she transforms the traces into a fixed-length histogram or *wavelength coefficients*. With this, she manages to achieve a 71% accuracy.

### 2.2.3 Defenses

Not only has there been research regarding different attacks but there are also various works that describe possible defenses. First of all, Tor already implements *padding*, which means that all packets are padded to a fixed-sized cell of 512 bytes. Next, in response to the first attack by Panchenko et al., Tor also supported randomized ordering of HTTP pipelines [31, 16, 34]. Finally, on top of these defenses, fingerprinting on Tor is made more difficult by all of the background noise present. This is due to the fact that Tor also sends packets for circuit construction or just *SENDME*'s, which ensure flow control [30]. Although Wang et al. proposed a probabilistic method to remove these [52], they might still make the classification process slightly more difficult.

Lua et al. designed an application-level defense, that was able to successfully defend against a number of classifiers by modifying packet size, timing of packets, web object size, and flow size [34]. This is achieved by splitting individual HTTP requests into multiple partial requests, using extra HTTP traffic as a cover and making use of HTTP pipelining [7]. Although this has been a relatively effective technique to obfuscate traffic, several attacks have proven that this defense only still does not suffice [7, 53].

*BuFLO*, on the other hand, is a *simulatable* defense [53], designed by Dyer et al. that performed packet padding and sending data at a constant rate in both directions [11]. The disadvantage of this method is the high overhead required in order to keep the constant data rate. Some extensions have been described that try to minimize this overhead such as Nithyanand's work that uses existing knowledge of a website traces to maintain a high level of security [25].

More recently, Cherubin et al. developed the first website fingerprinting defense on the server side [8]. This can be particularly interesting for *Tor hidden services* that want to provide, all of their users, the privacy that they require. The attack uses a technique, called *ALPaCA*, which pads the content of a web page and creates new content to conceal specific features on a network level [8].

Finally, there are also some other techniques such as *decoy pages* and *traffic morphing* [55, 31]. Decoy pages or *camouflage* is a very simple technique that involves loading another web page whenever a web page is requested. This process provides more background noise that makes fingerprinting more difficult [31]. Whilst traffic morphing is a slightly more complex technique that changes certain features in the traffic in order to make it appear as if another page is loaded [55].

### 2.2.4 Critical Analysis

Most attacks, that have been described above, are based on a set of different assumptions. Here we list these assumptions and see if they are reasonable.

The first one we examine is the open and closed-world scenarios. One of the main problems with website fingerprinting is the amount of web pages readily available on the web. An open-world scenario tries to solve this issue by only classifying a small amount of web pages and by labelling the other ones as unmonitored. However, machine learning theory states that the bigger the world size, the more data is required. So the small size of the *hypothesis space* compared to our world size, could have a direct impact on the amount of false positives since the more web pages there are, the higher the probability that one of the traces will be very similar to one in the monitored set. Therefore, the false positive rates, described in the previously mentioned papers, might be higher in real life [33].

Nonetheless, even if those false positive rates are accurate, a very small amount of false negatives could have a large impact on the classification. M. Perry shows that if the FP rate is as low as 0.2% and just a 100 page loads, around 18% of the user base would be falsely accused of visiting at least one monitored website. Or after 1000 page loads, this

percentage increases to around 86% [33].

There are also a variety of different factors that are often not considered such as the rapidly changing nature of some web pages. Juarez et al. show that it takes around 9 days for the accuracy to drop from 80% to under 50% [21]. Additionally, the content of some web pages is dynamic and some of the traces will vary, depending on who visits the website, making the classification for a large set of people difficult. Not only is dynamic websites an issue but different users will also be using different versions of the *Tor Browser Bundle* and might load the web page from different locations. This can decrease the accuracy with 70% and 50% respectively [21]. On top of this, we also need to consider multi-tab browsing, where a client might be loading multiple web pages at the same time. Although some papers consider this [15], most assume that the attacker know where the trace of a single page starts and ends.

## 2.3 Deep Learning and Automatic Feature Selection

In the following section, we will give a very short introduction to deep learning and describe some of the deep learning solutions that allow us to perform feature extraction. We start by introducing deep learning by exploring *artificial neural networks* and *stacked autoencoders*. Then we move on to *recurrent neural networks* and *sequence-to-sequence models*. Finally, we describe some of the issues with deep learning, such as the *vanishing gradient problem*. All of these explanations assume some familiarity with neural networks and only aim to give a high-level overview of the most important concepts.

Machine learning models basically take some value as its input and output a value, whilst trying to minimize some sort of error. All of the learning models that we explore below are forms of *supervised learning*. This means that we know the expected output and minimize the error between the actual output and the expected output.

### 2.3.1 Artificial Neural Networks

*Artificial neural networks* consist of a network of nodes, called *neurons*. These neurons are named and modeled after their biological counterparts. One of the simplest ones, is called a *perceptron*, which consists of a set of *binary inputs*, *weights* and an *activation function*. Hence, essentially it weights different evidence by assigning a different weight to every input. Next, the output of the neuron can be calculated as follows:

$$\text{output} = f\left(\sum_i w_i x_i\right)$$

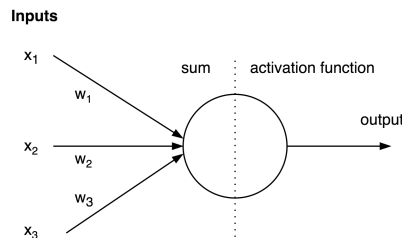


Figure 2.3.1: Model of a perceptron with three inputs.

This function  $f$  represents the *activation function*. Essentially, the activation function expresses the idea that a neuron can ‘fire’ after the sum of the inputs exceeds a certain threshold. There are certain different functions such as the *step function*, *sigmoid function* and the *tanh function*, which are all outlined in figure 2.3.2.

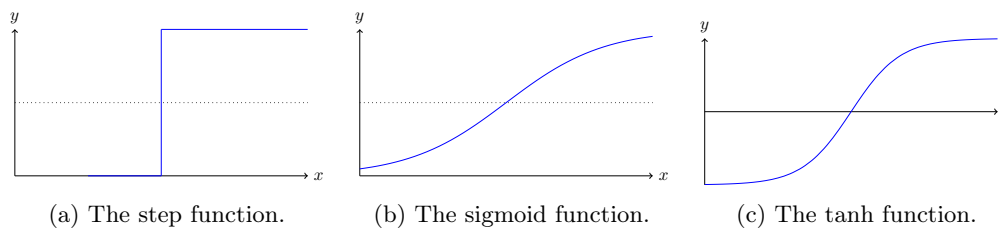


Figure 2.3.2: Examples of different activation functions.

Now to learn a function, it adapts the weight  $w_i$  such that it minimizes the error between the predicted and the expected output. In order to achieve this, we need some manner of quantifying the error, called a *loss function*. The most commonly used ones are the *mean squared error* ( $MSE = (x - x')^2$ ) and *absolute loss* ( $AL = |x - x'|$ ) [24]. There are many other cost functions such as *cross-entropy*, but they are not often used for the models that will be described below.

To build a neural network, several of these perceptrons are connected together. The most standard network is a *multilayer perceptron*, also called a *feedforward neural net*. This specific network, as can be seen in figure 2.3.3, consists of an *input layer*, one or more *hidden layers* and an *output layer*. All of these layers have an arbitrary number of neurons and the connections between these neurons can only go from left to right and can never form a loop. It is known that these kinds of networks can learn to approximate any function [50].

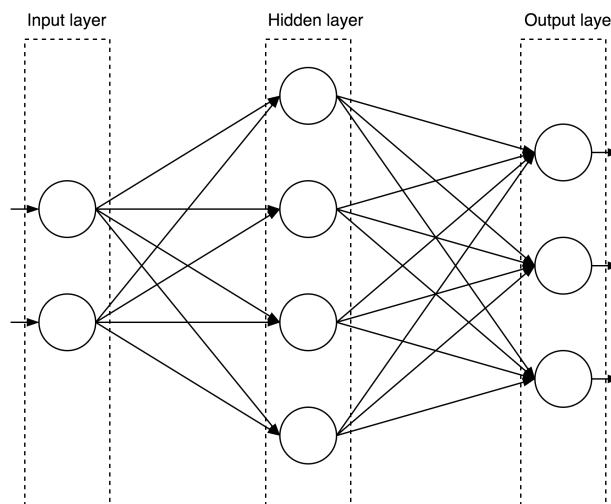


Figure 2.3.3: Example of a feedforward neural network with one hidden layer.

Now that we know how to construct these networks, we still need to manage to assign the appropriate weights to every connection such that the loss function is minimized. These weights can be learnt by using an algorithm, called *backpropagation*. The optimization process usually starts with initializing all of the weights with a random value and then running backpropagation, which is structured as follows [24]:

1. Compute the outputs, given a certain input (*feedforward pass*).
2. Calculate the error vector.
3. Backpropagate the error by computing the differences between the expected output and the actual output, starting at the output layer and working towards the input layer.
4. Compute the partial derivatives for the weights.
5. Adjust the weights by subtracting these derivatives, multiplied by the *learning rate*.

The learning rate is essentially a hyperparameter to the model that defines how fast the network learns. If its value is high, the model learns quickly and if the value is low, the model learns more slowly but the learning process will be more accurate. The propagation process is also often done in *batches*, which means that you calculate the propagations of a fixed amount of input vectors and only once this has finished, the weights are changed. The size of these batches is a hyperparameter of the model.

### 2.3.2 Stacked autoencoder

These feedforward neural net can be used to perform feature selection, by using a network called an *autoencoder*. This network tries to learn the *identity function*  $f(x) \approx x$  when the number of neurons in the hidden layer is smaller than the ones in the input and output layers. Hence, essentially the network is trying to learn how to compress the initial feature vector [2].

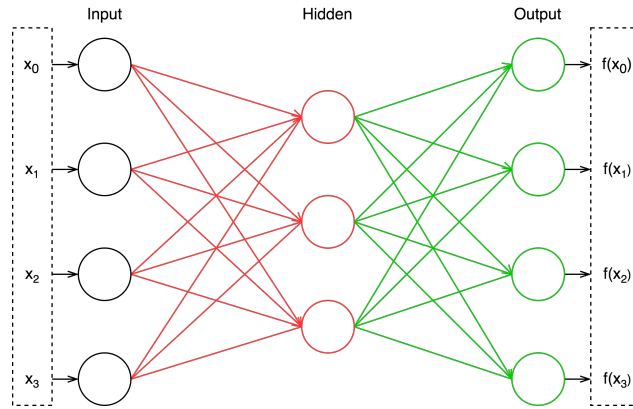


Figure 2.3.4: Structure of a simple autoencoder.

In order to learn an even more compressed representation of the input, multiple layers can be introduced, where each hidden layer contains even less nodes than the previous one. However, the problem with this approach is that the deeper the network, the more difficult it can be learn the appropriate weights [24]. A solution to this problem is a greedy approach where each layer is trained in turn and then stacked on top of each other. By this we mean that we first train the first layer, just as before. Next, the weights of the first layer are used to transform the raw input in a compressed representation. This representation is then used to train the second layer and so on.

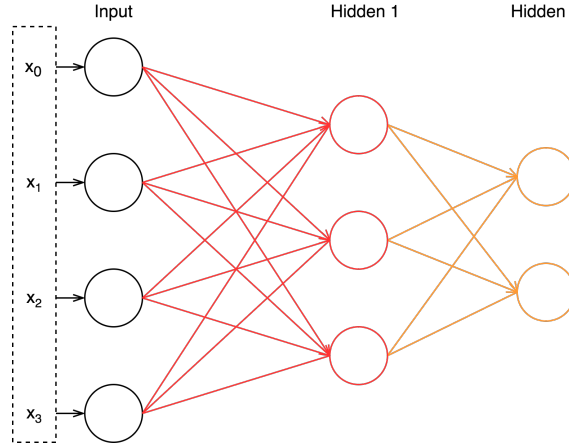


Figure 2.3.5: Stacked autoencoder.

### 2.3.3 Recurrent Neural Networks

Although stacked autoencoders have successfully been used in several cases [37, 54], they still have a couple of drawbacks. First of all, they require a fixed length input, which means that sequential information will need to be preprocessed. Next, they also assume that all of the inputs and outputs are independent of each other [6] whilst this might not be the case for sequential data such as the cell traces. Hence, we will look at *recurrent neural networks* (RNNs), which relaxes some of the restrictions with a feedforward neural net. They allow connections to form loops.

These loops basically represent that the network can be *unrolled*. This we mean that if we have the same network as in figure 2.3.6 and you have a sequence that has a length of  $n$ , you unroll the network for  $n$  steps. Hence, essentially an RNN has ‘memory’ that affects the outcome of the computation [6]. At every step, you can calculate the output by performing the calculation:

$$\text{result}, h_t = f(h_{t-1}, x_t)$$

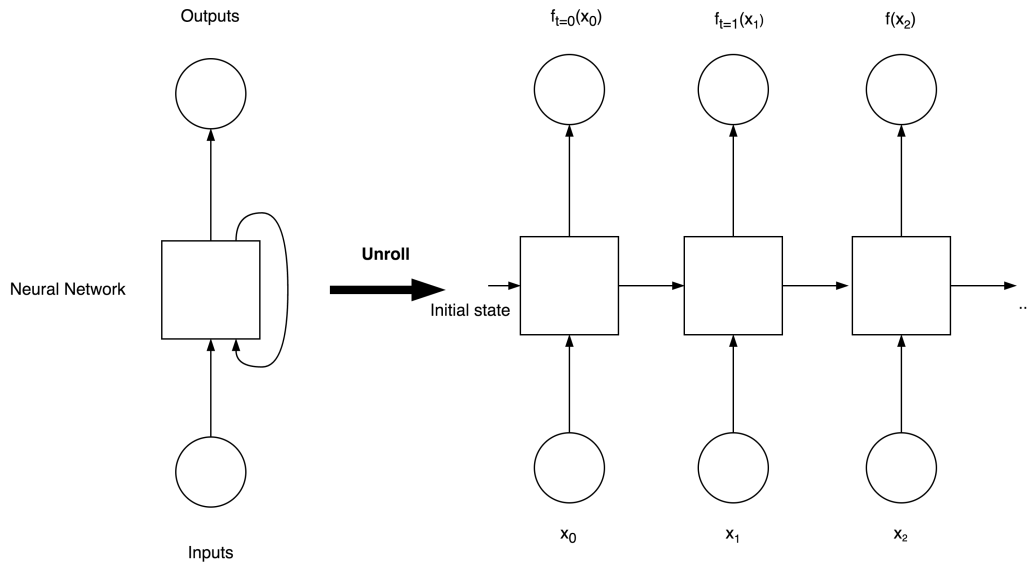


Figure 2.3.6: Structure of a RNN and an example of the unrolling process [6].



Unlike traditional deep networks, every layer in the unrolled network shares the same weights  $W$ . This greatly reduces the amount of parameters the network needs to learn [6].

One of the problems with RNNs are that they struggle to learn long-term dependencies [4]. But some cells, more specifically *long short term memory* (LSTM) and *gated recurring unit* (GRU) cells do not have this problem [19, 28, 9]. Rather than just having a single neural network layer, LSTMs consist of four different layers, that all interact in a novel way [28]. These interactions are outlined in figure 2.3.7. Although a full description is outside the scope of this paper, essentially the top state remains relatively unchanged and can therefore store long-term dependencies. Whilst the bottom state contains more short term information [28, 19].

The *gated recurring unit* (GRU) is a slightly more simple model than an LSTM. The cell combines several of the gates and states to minimize the amount of parameters the model needs to learn. Therefore, GRU allows a model to learn at a faster pace. Whilst LSTM cells have a greater expressive power [19, 28, 9].

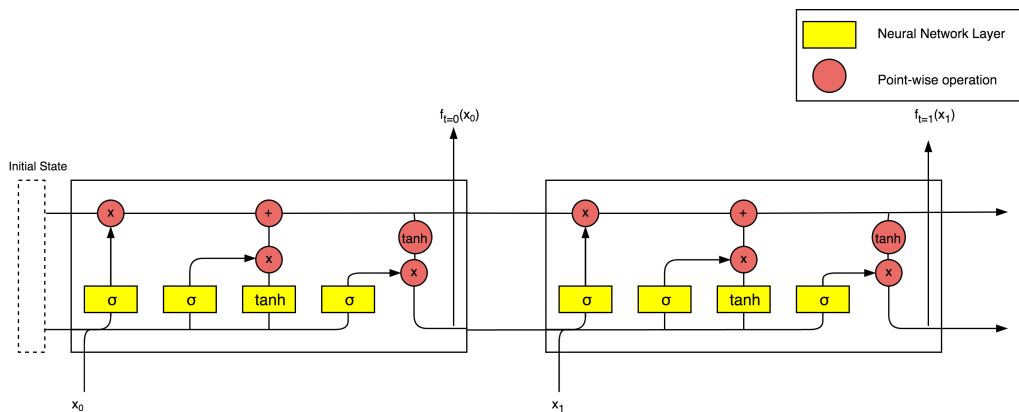


Figure 2.3.7: Interactions within a LSTM cell [28].

Recurrent neural nets cannot make use of the same backpropagation algorithm used by feedforward neural nets since the unrolled layers share the same set of weights. Instead, we use a different optimization method, called *backpropagation through time* (BPTT). BPTT is very similar to standard backpropagation with the key difference of summing up the gradients for the weights at every time step [6].

### 2.3.4 Sequence-to-Sequence Model

One specific type of RNN, that could be used for feature generation, is a *sequence-to-sequence model*, introduced by Cho et al. [9]. Sometimes called a *encoder-decoder model*, it consists of two RNNs where one RNN encodes a sequence into a fixed-length representation and the other decodes those representations into another variable-length sequence [9]. The models have been proven to be successful in several *natural language processing* (NLP) tasks such as translation tasks [9, 42, 39]. However, since the model extracts a fixed-length representation, we might be able to train a sequence-to-sequence model on a copy task, just like an autoencoder, and use that *thought vector* as the extracted features.

The model works as follows, each box in figure 2.3.8 represents an RNN cell. Next, the encoder RNN is unrolled, depending on the length of the input vector. After the encoder is finished, the final state vector is stored in a *thought vector* variable. Next, the decoder is unrolled and run, with the thought vector as its initial state and a start token as the first input. Finally, the output of the previous cell is then used as the input to the next cell until the network produces the end of sequence token [9]. The encoder and the decoder can

share the same weights or, as is more common, use a different set of parameters [39].

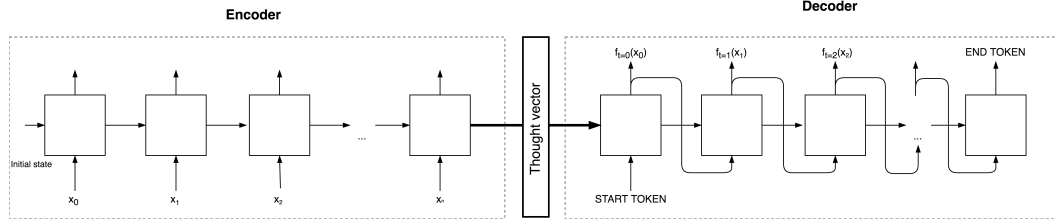


Figure 2.3.8: Structure of a sequence-to-sequence model [5].

If this model is now trained on a copy task, where it tries to reproduce the input sequence from a thought vector, it means that just like an autoencoder, the network has learned to compress any variable-length sequence into a fixed-length representation. The size of this thought vector, however, is another hyperparameter to the model.

To allow a sequence-to-sequence to learn more complex representations, Sutskever et al. experimented with multilayered LSTM cells [42]. They showed that this model was very successful in a English to French translation task and managed to cope with long term dependencies. Additionally, they also showed that reversing the sentences made the learning process easier. Sutskever et al. do not have a complete explanation for this phenomenon but they believe that it is due to the many short-term dependencies in the dataset [42].

In addition to reversing the traces, Oshri et al. introduced the idea of using a *bidirectional RNN encoder* [29]. The idea behind this kind of encoder is based on the fact that the output at time  $t$  might not only depend on past information but also on future information. To combat this issue, the encoder consists of two layers, the forward and the backward layer, stacked on top of each other. Next, the output is computed, based on the hidden state of both layers [6]. Finally, to extract the thought vector, the last state of both layers are extracted and averaged.

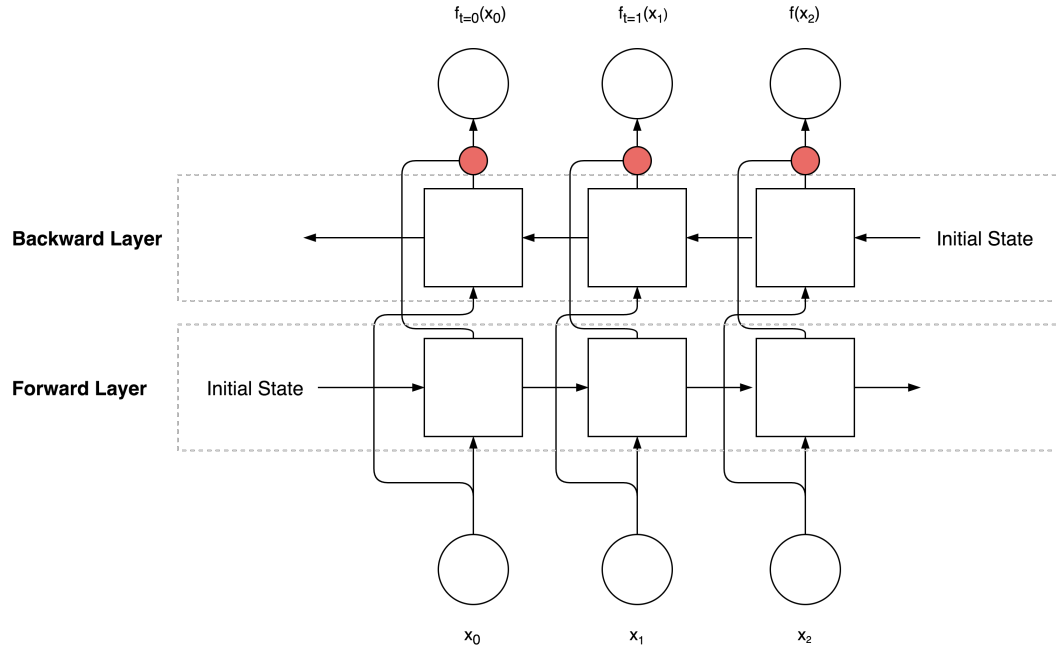


Figure 2.3.9: Structure of a bidirectional RNN [6].

This kind of model sounds more appropriate than one with a simple encoder. However, Oshri et al. argue that the performance gain is limited, whilst introducing a lot more parameters [29].

Some research by Bahdanau et al. has been to improve the performance of encoder-decoder models by introducing *attention mechanisms* [3]. This mechanism allows the decoder to have a more direct access to the input, thereby relieving the encoder by having to embed all the information in a fixed-length vector. Although this technique might work well for translation tasks, our work focuses on the feature extraction process. Therefore, we will not perform any analysis on sequence-to-sequence models with an attention mechanism.

At the time of writing, there hasn't been much research regarding using sequence-to-sequence models for feature extraction. In fact, currently they are most often used for translation and other NLP tasks. Hence, they are most often used as a classification task where the model classifies which word is likely to be next [39, 9, 29, 42].

### 2.3.5 The Vanishing Gradient Problem

Deep learning suffers from an problem, called the *vanishing gradient problem* [24]. This problem represents the fact that neurons in early layers tend to learn a lot slower than those in the last layers. The vanishing gradient problem is also the reason why some RNNs are not able to learn long-term dependencies. This is due to the fact that the derivatives of both tanh and sigmoid activation functions approach zero near both ends. Thus when one of the neurons is saturated, which means that the value of the gradient gets close to zero, it drives the gradients of previous neurons to zero as well [6]. Hence, since those gradients are close to zero, they vanish after a couple of time steps. As RNNs tend to be a lot deeper than traditional feed-forward networks, this problems tends to be a lot more common [6].

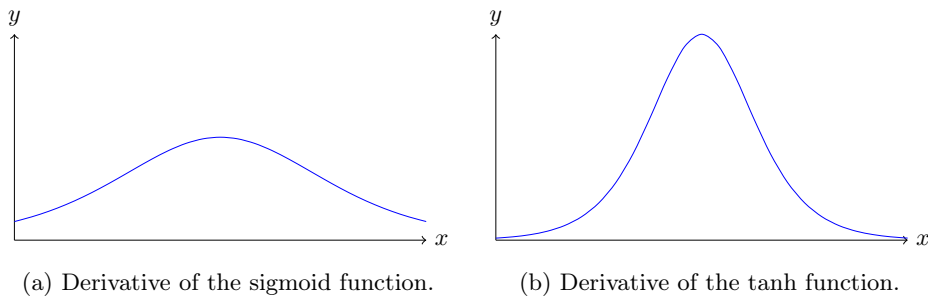


Figure 2.3.10: Derivatives of activation functions.

One solution to this problem is to use a ReLU activation function, as the derivative is either 0 or 1 [6]. But both LSTM and GRU cells were especially designed to solve this issue in RNNs and therefore we will be using those throughout the rest of this paper [28, 19, 9].

### 2.3.6 Regularization

Neural networks can easily *overfit*. By this we mean that the model learns noise in the training data, rather than learning to generalize. For instance, as can be seen in figure 2.3.11, the model is generalize when it fits a straight line but overfits otherwise. The reason why neural nets often overfit is because of the large amount of free variables [24].



Figure 2.3.11: Example of overfitting.

Overfitting can be reduced by performing *regularization* without the need to decrease the size of the network. The most popular techniques are *L1*, *L2*, *dropout layers* and *batch normalization* [24]. Both L1 and L2 regularization work by penalizing large weights and thus making the functions less complex. They do this in a different fashion, for instance L1 adds the sum of the absolute values of the weights to the cost function [24]. Whilst L2 has a different *regularization parameter*, which generally performs better in practice [24].

Dropout, on the other hand, doesn't modify the cost function but it changes the network. Essentially, it ignores several neurons while performing an iteration of the back-propagation process. After that iteration, it picks a set of different neurons to ignore and repeats the process. The effect of this is that the network basically consists of an average of various different networks. Each of those neurons will overfit in a different way so the average should reduce the total amount of overfitting [24].

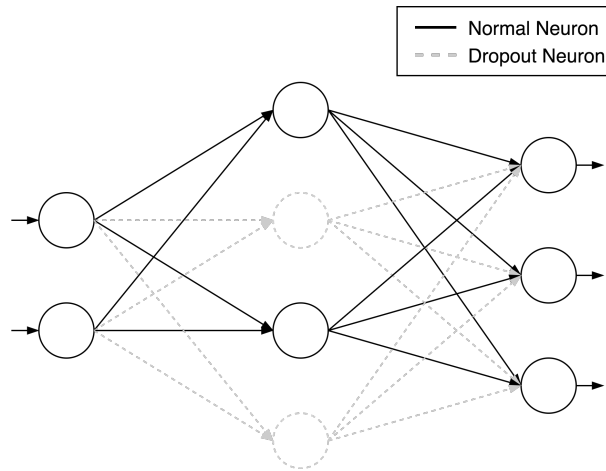


Figure 2.3.12: Feedforward network with dropout.

Finally, batch normalization is a relatively new technique that also changes the network slightly by making the normalization process part of the network and performing it for every minibatch [20]. Usually this wouldn't be considered a regularization technique. However, it has been proven that it acts as a regularizer, and sometimes eliminates the need for dropout layers [20].

## 2.4 Software Libraries

There are various different numerical computation libraries to efficiently implement deep learning algorithms. However, the most common ones are *Tensorflow*, *Theano*, *Keras* and *Torch* [43, 45, 22, 47]. The first three all provide their API in Python, whilst torch only be used in Lua. Although a python API, called *Pytorch* has been open-sourced very recently. All of the above allow you to use *cuDNN* to perform GPU-accelerated computations.

Tensorflow has a C++ backend and its low-level API is provides more than enough flexibility to implement a sequence-to-sequence model. They do also provide a high-level API that include different RNN cells and even a sequence-to-sequence model. However, this model has been specifically designed for NLP tasks and therefore does not support the computations that we are trying to achieve [43]. Theano is very similar to Tensorflow but it does not provide all of the high-level tools [45].

Keras, on the other hand, runs on top of either Tensorflow or Theano as a back-end. Hence, the library provides high-level functions to create different models, rather than the low-level APIs provided by both Tensorflow and Theano. Although this library would allow us to quickly prototype a sequence-to-sequence model, it does not provide the low-level access that we might need in order to be able to tune certain hyperparameters [22]. Finally, Pytorch is also a very attractive option but the current release is still a beta version [47].

After a careful analysis, Tensorflow provides us with the highest flexibility whilst still providing tools to easily perform certain calculations with their high-level API. Hence, the decision to use Tensorflow for the implementation of our main model. On top of Tensorflow, we will also be using sklearn, which provides us with a large amount of machine learning models to perform some of our testing, and numpy for fast data preprocessing. Both of which were chosen since they are in python, therefore allowing easy code-reuse for different modules.

## 2.5 Data Sets

In order to collect data, a web crawler needs to visit a large set of website over Tor where the traffic is recorded for every page visited. This web crawler can emulate user browsing or just visit websites such as the one constructed by Panchenko et al. [30] or simply visit web pages in *Alexa's Top 10,000*, which contains the most commonly visited pages. Or real data can be collected from users but due to privacy concerns, this data is often hard to get by.

After a web-crawler has been set up, data is collected through a TCP dump. This data is then often preprocessed and converted into *Tor cells*. The reason for this is because Tor pads packets to a fixed-length (512 bytes) [44] hence these cells are a simple representation of the traffic. Next, probabilistic techniques are used to remove SENDMEs [52]. After this processing, a cell looks like a list of tuples, each containing a time value and a direction, which represent whether it was an incoming or outgoing packet.

Timestamp	Direction
0.0	OUT
0.0630009174347	OUT
0.575006008148	IN
0.575006008148	IN
0.691473960876	IN
0.719605922699	OUT

Table 2.1: Extract of a cell trace [14]

All of these individual Tor cells are then stored within different files. The names of these files will have the following format for monitored pages `<page_id>-<instance_id>.cell` and `<page_id>.cell` for unmonitored pages.

There are several data sets readily available that have already been preprocessing. Some of the largest ones are the one used by Wang et al. [53], Greschbach et al. [14]. Wang et al.'s dataset contain traces for a 100 monitored websites with 90 instances each and 8400 unmonitored sites, all from Alexa's top 10,000. Greschbach et al. collected an even larger dataset with 100 samples of each website in Alexa's top 9,000 and one sample for 909,000 unmonitored sites [14]. Both of these datasets are definitely large enough to train our sequence-to-sequence model.

## Chapter 3

# Threat Model and Problem Statement

In the following section, we further describe the adversary models from which we deduce a list of requirements.

### 3.1 Threat Model

We consider an adversary, who wants to perform a website fingerprinting attack against Tor. Tor, specifically, since it has become one of the most widely used internet privacy tools. As specified in figure 2.2.1, the adversary is a *local eavesdropper*. Hence, the attacker passively collects encrypted web traffic between the client and the first tor node, or the *entry guard*. This is achieved by either monitoring the link itself or a compromised entry node. Next, it performs an analysis on that data to classify which specific web pages the client is visiting.

This analysis can be performed with several different goals. The first one is to identify whether or not a user visits a web page from a set of monitored web pages. Thus the attack is essentially a *binary classification problem*, where you label a web page as *monitored* or *unmonitored*. Or the adversary might want to know which specific web pages a user visits, or a *multiclass classification problem*.

Within this adversary model we do make various assumptions. First of all, the adversary is not interested in blocking Tor traffic nor in modifying any traffic. Next, the adversary is able to replicate the conditions under which the user browses the internet such as download speeds, OS and TBB. On top of this, the adversary can also determine the beginning and the end of a user session on a web page and that the attacker has enough resources to collect traffic and train a deep learning model. Finally, we also make the adversary is unable to decrypt the traffic and thus only has access to metadata such as traffic direction, length and timing of packets.

### 3.2 Problem Statement

At the time of writing, most WF works make use of machine learning techniques that require a fixed-length input. Even though a traffic trace consists of a variable-length sequence of packets. Therefore, these works often rely on a laborious, time-consuming process to extract fixed-length representations, or *fingerprints*. But there is no guarantee that these fingerprints are the most appropriate ones. On top of that, the previously mentioned process often requires domain-specific knowledge, making the attack even more difficult. Thus, here we investigate the use of automatic feature generation techniques to extract features automatically, without the need for any domain-specific knowledge.

Hence, our main contribution is the creation of a new models, capable of learning fixed-length fingerprints from variable-length traces. This means that we will not be focusing on creating a new attack, but rather re-using existing attacks with these automatically-generated features. Next, we will contrast the performance of these different models and note which ones seem to be the most appropriate for the threat model described above.

## Chapter 4

# Attack Design

In this section, we describe the design of our automatic feature generation model, outline the attack strategy and explain certain design decisions. Most of this section will be split into two different sections. First we describe the feature generation process and then the overall attack.

### 4.1 Stacked Autoencoder

A stacked autoencoder takes a fixed-length input and tries to learn how to compress that vector. Thus if we were to use it for our fingerprinting extraction model, we will need to manner to map the variable-length traces into a fixed-length one. There are various different manners of doing this. One of the most naive ones is to find the longest length trace and pad all of the other traces up to that length. However, in Greschbach et al.'s dataset, this length is around 250,000 [14], which means that our network will need to be incredibly deep.

Instead, we pick the average length of the traces, which is around 3,000 and cut or pad traces which are longer or shorter. On top of that, we deal with the fact that each packet is represented by a tuple, by just multiplying the time by the direction. Therefore, all of the outgoing packets are positive and the incoming ones will be negative.

### 4.2 Sequence-to-Sequence Model

As described in section 2.3.4, a sequence-to-sequence model is able to learn how to construct a fixed-length representation from a variable-length sequence. However, here we outline the exact structure of the model and show the implementation in Tensorflow.

One of the parameters, which has a large impact on the accuracy of the reconstruction process is the *amount of hidden states* in the RNN cells. Every neural network layers within a cell has a given amount of neurons. Hence, if for example the amount of hidden neurons is set to 100, our state at every cell is represented by vector of length 200 (since the state is represented by two vectors of length 100). The higher this number, the easier it should be to learn a representation, as the compression factor is lower. But we also need to consider the fact that the higher the amount of hidden neurons, the more variables the model needs to learn.

Each value in a trace can be represented by a vector of length two (timestamp and direction), as seen in table 2.1. Therefore, if the amount of hidden cells is not equal to two, we will also need to *project* the input and output to the necessary dimensions.



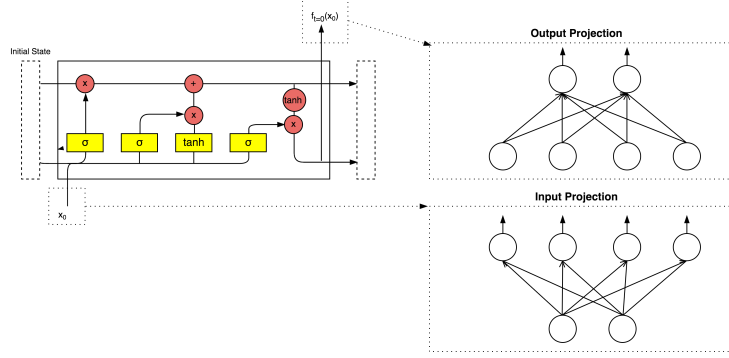


Figure 4.2.1: Example of projection within a LSTM cell with 4 hidden states.

Some of the traces can be particularly long and therefore the network needs to be unrolled to extreme lengths [14]. In fact, given memory constraints, this issue can become a major problem. This can be solved by cutting the traces after a couple seconds since it has been shown that the first part of a trace carries more information than the latter.

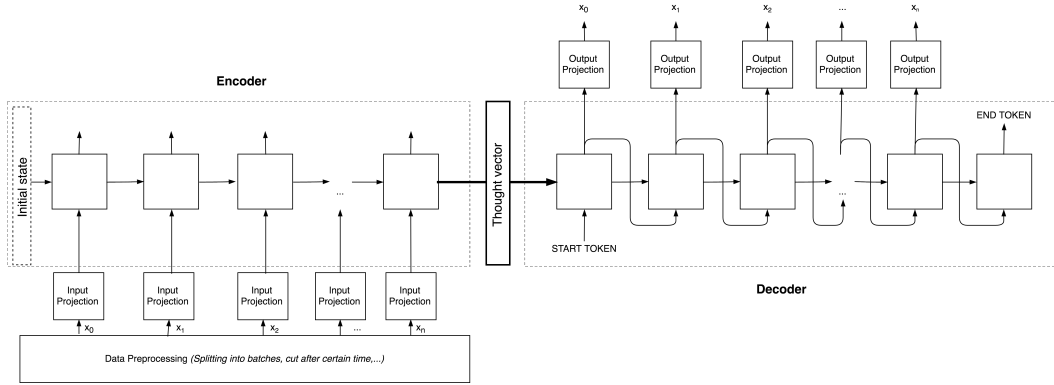


Figure 4.2.2: Overall structure of our sequence-to-sequence model.

Based on the above, we can construct our *computational graph* in Tensorflow as outlined in figure 4.2.2.

### 4.3 Attack Strategy

Here we consider an adversary that relies on deep learning to extract fingerprints for a website fingerprinting attack. This adversary can have two different goals in mind, as previously stated in section 3.1. In this work, the adversary relies on a sequence-to-sequence to extract the fingerprint from Tor cells for both goals. The full attack, however, can be split up into four different stages: *data collection*, *fingerprint extraction training*, *classifier training* and *the attack*.

#### Data Collection

1. Choose web pages that the attacker wishes to monitor.
2. Collect traffic for a set of monitored sites and unmonitored sites.
3. Convert the raw TCP data into Tor cells.
4. Remove SENDMEs and other noise.

## Fingerprint Extraction Training

5. Further process the data into batches and perform any other preprocessing required by the model such as cutting or padding the traces.
6. Prepare the fingerprint extraction model.
7. Train the fingerprint extraction model on a copy task for monitored and some un-monitored web pages.
8. Extract fingerprints from data by using the trained model.

## Classifier Training

9. Given a classifier, train it using the extracted fingerprints.
10. Measure performance of classifier.

## The Attack

11. Passively capture traffic from Tor users.
12. Pre-process the collected data.
13. Extract fingerprints using the trained fingerprint extraction model.
14. Classification via the trained classifier.

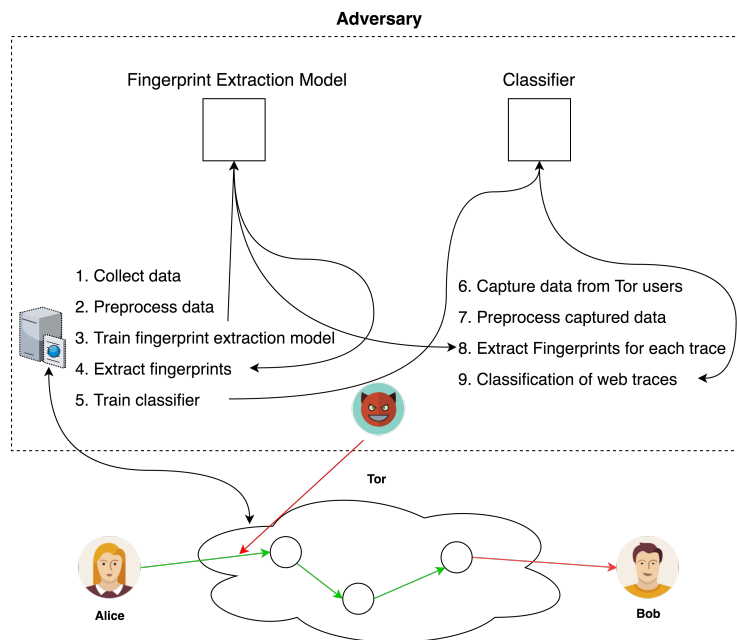


Figure 4.3.1: Attack strategy.

### 4.3.1 Data Collection

As previously mentioned, the data collection process first requires the adversary to choose a set of  $n$  websites to monitor. Next, the adversary crawls these pages a total of  $i$  iterations to ensure that a classifier has enough data to generalize. As suggested by Wang et al. when performing page loads, browser caching should be disabled since Tor does not allow caching to disk and therefore the browser cache is cleared every time it is restarted [52]. After the collection, the TCP data is converted into Tor cells and probabilistic algorithms are used by Wang et al. [52] to remove SENDMEs. This data can be processed further, however we chose not to since the model should be able to learn how to perform this processing.

Most of our analysis will be done using the dataset provided by Greschbach et al. [14], which can be used for open-world analysis since it provides us with 100 samples for Alexa’s top 9000 websites and one sample of one sample for 909,000 unmonitored sites. For the rest of this paper, we will refer to this dataset as **GRESCHBACH**. Next, to see how our model performs on a dataset, whose data is recorded at a different time and under different circumstances, we will also be using the dataset provided by Wang et al. [53], which we will call **WANG14** [30]. This set is slightly smaller with 100 monitored websites with 90 instances each and 8400 unmonitored sites.

### 4.3.2 Fingerprint Extraction Training

In order to truly evaluate the model, we need to split the data up into a training and validation set. We do not train the model on any data in the validation set but instead use it to see how well the model performs on unseen data. For this split, we use a *stratified shuffle split*, meaning that we shuffle the data and then perform the split, whilst preserving the class distributions. On top of training the feature extractor with monitored pages, we also train it on unmonitored pages, as it needs to be able to extract features effectively from both sets.

Whilst training and extracting the fingerprints, *mini-batch processing* will always be used. This will allow us to gain a performance boost and perhaps even have a faster convergence.

When dividing the data up into these batches, we also need to determine how big they will be. The bigger they are, the larger the performance gain will be but the lower the accuracy might be. Additionally, the size of the batches also depend on the amount of available memory since we cannot have the VM run out of memory whilst training.

On top of determining the batch size, the individual models require different preprocessing steps and tuning of different parameters to get a performance gain.

### Stacked Autoencoder

As previously mentioned, after dividing the data up into batches, we either need to cut or pad the traces such that they all are of a fixed-length. Next, we know that the first layer needs to have the same amount of neurons than the length of the input. But after that, there are a variety of different architectures that need to be chosen. Some of which are outlined below:

- How many hidden layers we want. The more there are, the more complex the functions are that the model can learn but the harder it is to train.
- The amount of hidden neurons in each layer. This number should gradually decrease to the number of features we would like to extract.
- The activation function of the neurons. The most popular ones being *sigmoid*, *ReLU* and an *atan*.

Now that the model has been constructed, there are still several learning parameters that need to be tuned:

- The optimizer to use (*adam*, *gradient descent* or *RMSProp*) [43].
- Learning rate ( $\gamma$ ) for the previously chosen optimizer.
- Amount of traces within a single mini-batch ( $b$ ).
- Cost, or loss function ( $f$ ) to minimize (*mean squared error* (MSE), *absolute loss* (AL) or *cross-entropy*) [43].

### Sequence-to-Sequence Model

Each batch can either be presented in *batch-major* or *time-major* form. Although time-major is slightly more efficient [43], we opt for a batch-major form, since it makes the fingerprint extraction easier. Next, after the data has been divided into mini-batches, we perform some further processing such as cutting the traces after several seconds. Finally, since all of the traces within a batch need to be of the same length, padding is performed as a final preprocessing step.

After collecting and fully preprocessing the data, the adversary can start to construct the sequence-to-sequence model. However, in order to do so, there are a variety of different architectures that need to be chosen. Some of which that we will consider are outlined below:

- Which sort of RNN cells to use. This can either be a GRU or an LSTM cell. We could also potentially investigate the usefulness of multilayered RNN cells but it would make the network even deeper.
- Using a bidirectional encoder to ensure that the output at time  $t$  is not only affected by past information but also on future information.
- The amount of hidden states within a RNN cell, which affects the length of the fingerprints.

The adversary would like to have the amount of learned features to be small enough. Since the bigger the amount of features, the more training data is required, due to the *curse of dimensionality*. However, if the amount of hidden states within a cell is too low, it might not be able to effectively recreate a trace from a fingerprint, due to the lack of data. Hence, the adversary will need to consider this tradeoff when choosing the model architecture.

Now that the model has been constructed, the adversary still has to choose various learning parameters such as:

- The optimizer to use (*adam*, *gradient descent* or *RMSProp*) [43].
- Learning rate ( $\gamma$ ) for the previously chosen optimizer.
- Amount of traces within a single mini-batch ( $b$ ).
- Cost, or loss function ( $f$ ) to minimize (*mean squared error* (MSE), *absolute loss* (AL) or *cross-entropy*) [43].
- Whether or not to reverse the traces. This parameter does not make a difference when using a bidirectional encoder.
- After how much time the traces are cut.

After these parameters have been tuned, the computational graph can be constructed in Tensorflow and the training can be started using the training data. When this training has been completed, fingerprints can be extracted for data all the data in the test set.

### 4.3.3 Classifier Training

After the adversary has extracted the fingerprints for websites within the test set, they need to construct a classifier. This classifier can then use these fingerprints to learn how to classify web pages. Most works so far rely on some sort of *supervised machine learning* techniques such as *support vector classifiers* (SVC), *k-nearest neighbours* (kNN), *random forests* (RF) or *naive bayes* (NB) [31, 30, 53, 16, 15]. All of these algorithms rely on different techniques but an explanation of their inner workings is outside the scope of this paper. Instead, we will consider them as *black box* models. This means that all we know is that we can apply a `fit` function to the models, which causes them to learn how to classify the fingerprints and a `predict` function, which predict the class of given inputs. However, in reality, an adversary will have to carefully consider which models to use and tune the hyperparameters of that specific model to get the best possible performance.

To measure the performance of our black-box models, we use a similar technique as we did in the previous section. We split our test set up into two more sets, a *classifier training set* and a *classifier test set*. But since training a classifier, requires less time, we can use another technique, called *stratified k-fold validation*. Here we split our original test set up into  $k$ , mutually exclusive, folds. Next, one of the folds is chosen to be the classifier test set and all of the other folds form the classifier training set. This process is repeated for  $k$  iterations, where for each iteration, a different fold is chosen to be a test set. However, what is special about stratified k-fold validation is that the class distributions are preserved within the folds.

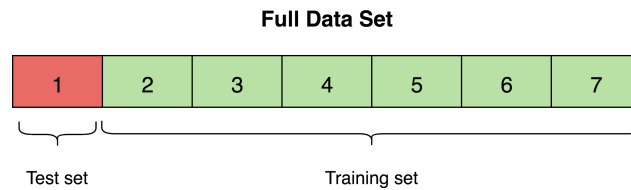


Figure 4.3.2: Example of one iterations in a k-fold validation ( $k = 7$ ).

For each iteration of the validation process, several statistics can be recorded and then averaged over all iterations. This process ensures that every data point will be in the test set at least once and therefore giving us an accurate measure of these statistics.

Some of the statistics that are used for the performance measure of models, are outlined below within the context of a website fingerprinting attacks:

- **True Positive Rate (TPR)** is the probability that a monitored page is classified as the correct monitored page [16].
- **False Positive Rate (FPR)** is the probability that an unmonitored page is incorrectly classified as a monitored page [16]
- **Bayesian Detection Rate (BDR)** is the probability that a page corresponds to the correct monitored page, given that the classifier recognized it as that monitored page [16]. This can be calculated as follows:

$$\text{BDR} = \frac{\text{TPR} \times \Pr(M)}{\text{TPR} \times \Pr(M) + \text{FPR} \times \Pr(U)}$$

where

$$\Pr(M) = \frac{|\text{Monitored}|}{|\text{Total Pages}|}, \quad \Pr(U) = 1 - \Pr(M)$$

This measure essentially indicates the practical feasibility of the attack, as the adversary is mainly concerned with this specific measure [16].

- **Accuracy (A)** is the percentage of correctly classified instances. Although it can be used as a rough indicator, it will not be used in the final conclusions because of the *accuracy paradox*, which arises due to class imbalance.
- **F1-Score (F1)** measures the harmonic mean between precision and recall [38].

$$\begin{aligned} \text{F1} &= 2 \times \frac{\text{precision} \times \text{recall}}{\text{precision} + \text{recall}} \\ &= \frac{2 \times TP}{2 \times TP + FP + FN} \end{aligned}$$

where

$$\text{recall} = \frac{TP}{TP + FN}, \quad \text{precision} = \frac{TP}{TP + FP}$$

This measure is particularly useful since it is not affected by class imbalance.

Rather than using another classifier, we could also potentially add a *softmax layer* on top of our encoder, like V. Rimmer uses on top of her stacked autoencoder [37]. We could also use this idea for both our autoencoder and the sequence-to-sequence model.

This process would involve first training the sequence-to-sequence model, then stacking the softmax layer on top of each unrolled cell in the encoder and using it for classification. What would be interesting about this approach is that the adversary can analyse how the probability of a trace being classified changes, as it analyses packets from the trace. However, this is outside the scope of this paper, as we are focusing on the feature extraction process.

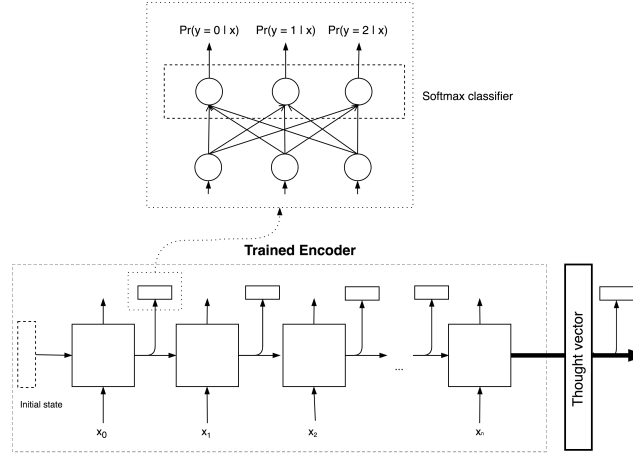


Figure 4.3.3: Example of encoder with softmax layer for 3 web pages.

#### 4.3.4 The Attack

Finally, after the adversary has trained the required models, the real WF attack can start. First, the adversary starts capturing web traffic data between the user and the first Tor node, as shown in figure 2.2.1. Next, the data is processed for the fingerprint extraction process, as described in section 4.3.2. After all processing has been done, fingerprints are extracted using the previously trained model. Finally, those fingerprints are used as features for a classifier, which classifies which web pages the adversary is visiting.

The time between data collection for training and performing the WF has to be kept as small as possible since Juarez et al.’s experiments show that website’s content changes greatly over time, therefore affecting the accuracy of the attack [21].

### 4.4 Code Structure

In the work, we will not be conducting the final stage of a WF attack. Instead, we will be reporting the results on the test sets. All of this is reflected in the overall structure of the code, which consists of four main components, as can be seen in figure 4.4.1.

All of the data that is used within this work has already been preprocessed, as described in section 4.3.1. The `feature_generation` module contains all of the code to further preprocess the data, train the fingerprint extraction models defined and to extract the fingerprints using the respective model.

All of the models that will be used for the attack section are defined in the `attacks` module. We tried to pick a variety of models used, to measure how our fingerprints work on different models. The logic to actually run the models is defined in the `run_models` module, which also does some data preprocessing and defines the logic for the stratified k fold validation and the different scoring methods.

Finally, there is also the `feature_extraction` module, whose use will be explained later.

As can be seen, all of the code is written in Python, due to the wide availability of machine learning tools. Except for the `kNN.go` file, in the `feature_extraction` module, which is written in *Golang* to gain a performance boost [36]. `../report.bib`

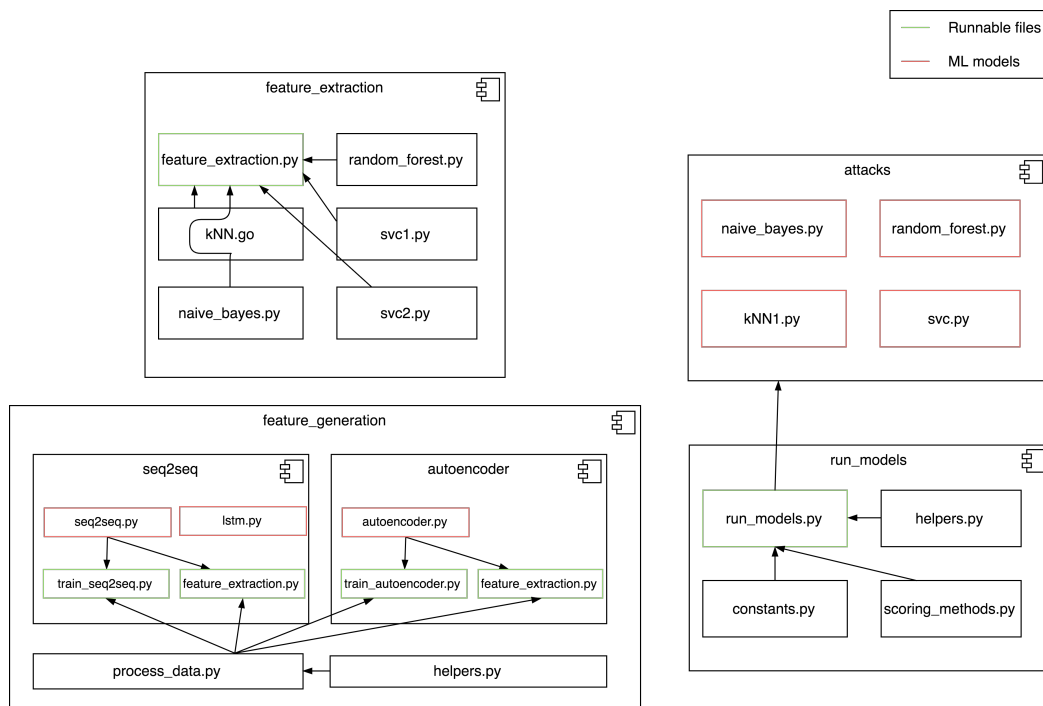


Figure 4.4.1: Diagram of how different components are related.



## Chapter 5

# Evaluation and Testing

In the following section, we outline how we will be evaluating the suggested model. Next, we will perform that evaluation and present the final results.

### 5.1 Experimental Setup

Since our deep model requires a large amount of computation, we like to make use of parallelization. Hence, all of our experiments that involve the sequence-to-sequence model have been run on an *Amazon EC2 p2.xlarge* instance. This VM has a *NVIDIA K80 GPU* with 12 GiB of GPU memory. This instance was then setup with *CUDA 8* and *cuDNN v5.1* [43, 27].

The rest of the experiments are run on a 2016 MacBook Pro, with a 2.9GHz Intel Core i5 and 8GB of RAM, running MacOS 10.12. To make sure that the same Python environment is used on both these machines, we consistently use *Python 3.6* and a *virtual environment* for the python dependencies.

As previously mentioned, the main dataset that will be used is **GRESCHBACH** but we will also be using some of the data in the **WANG14** dataset to see how the model performs on data that was recorded under different circumstances. From both these datasets, we will only be using the preprocessed Tor cells and not the raw TCP traffic data.

For all of the experiments that will be conducted below, we only consider an *open-world scenario*. This means that the test set will contain unmonitored pages that the sequence-to-sequence model or the classifier have never seen before. For this to work, we train the models on a large set of monitored web pages but also on a small percentage of unmonitored web pages such the classifiers can distinguish between both.

### 5.2 Evaluation Techniques

There are several different manners in how we can evaluate the feature selection process. First of all, we could analyse the training and test error, as the model learns. If the training curve suddenly drops, the learning rate might be too high. Or if the space between both the training and the test error increases, the model will clearly be overfitting.

However, these graphs only show us how well the model is at reproducing the trace from a fingerprint but now how well it performs in a WF attack. For this we need to train a classifier and see how well it performs by using the metrics described in section 4.3.3.

To be able to compare these fingerprints with hand-picked ones, we could run the classifier with hand-picked features and with the automatically generated ones. These hand-picked features are often chosen by experts and after a careful analysis of what the most appropriate features are. Hence, if the classifier with our fingerprints were to get similar results or even outperform the classifiers with the hand-picked features, we know that the sequence-to-sequence model has been successful. For these results to be accurate, we do not change any parameters within the classifiers. Thus everything, except for the features, stays fixed.

For the classifiers, we pick a small set of five existing models. We aim to pick models that have had an influence on the WF field whilst also having a variety of different classifiers.

This set includes the two *support vector classifiers* (SVCs) used by Panchenko et al. [31, 30], the k-fingerprinting attack, which relies on a *random forest* (RF) used by Hayes et al. [16] and finally the *k-nearest neighbours* (kNN) classifier used by Wang et al. [53].

For all of these models, we extract the exact same features as outlined in the respective papers and compare the performance of our generated features compared to the hand-picked ones. The code for this feature extraction process can be found in the `feature_extraction` module. After analysing all of these features, the most important ones seem to be [31, 30, 16, 53]:

- Total number of packets.
- Number of incoming packets.
- Number of outgoing packets.
- Percentage of incoming and outgoing packets.
- Concentration of packets.

We also aim to use the exact same hyperparameters described in the respective papers. More specifically:

- **SVC** [31] - a *radial basis function* (RBF) kernel with  $C = 2^{17}$  and  $\gamma = 2^{-19}$ .
- **SVC** [30] - uses the same hyperparameters as in the previous SVC but with different features.
- **RF** [16] - shows that the best accuracy/time tradeoff is made when  $k = 3$  and  $num\_trees = 20$ .
- **kNN** [53] - also shows that the best accuracy/time tradeoff is made when  $k = 2$  and  $k_{reco} = 5$ .

However, we do need to note that this might have an impact on the performance because these parameters have been specifically tuned for the hand-picked features and not for our fingerprints.

## 5.3 Evaluation

### 5.3.1 Learning Parameter Tuning

As mentioned in section 4.3.2, there are a couple design decisions that need to be made regarding different architectures and learning parameters for the sequence-to-sequence model. We first try to aim to get the appropriate values for the learning parameters within a simple encoder and decoder with LSTM cells and 120 hidden states.

First, we start by varying the mini-batch sizes from 20 to 400 in steps of 20. The higher the batch size, the longer it takes before making a weight update and the lower the value, the more noise in the training data. For instance, as can be seen in figure 5.3.1, there is clearly a trend of the training error decreasing over time. However, since the batch size is low for the first case, there is a higher probability of having a batch where the training error is high for all of the samples within that batch. Hence, the data will look very noisy. Whilst the greater the batch size, the less noise there is in the data and therefore the easier it is to spot trends.

Preferably we would like to have an even greater batch size than 200 but due to memory constraints, the model runs out of memory when the amount of hidden states is greater than 100. Thus through the rest of the report we will use a mini-batch size of 200.

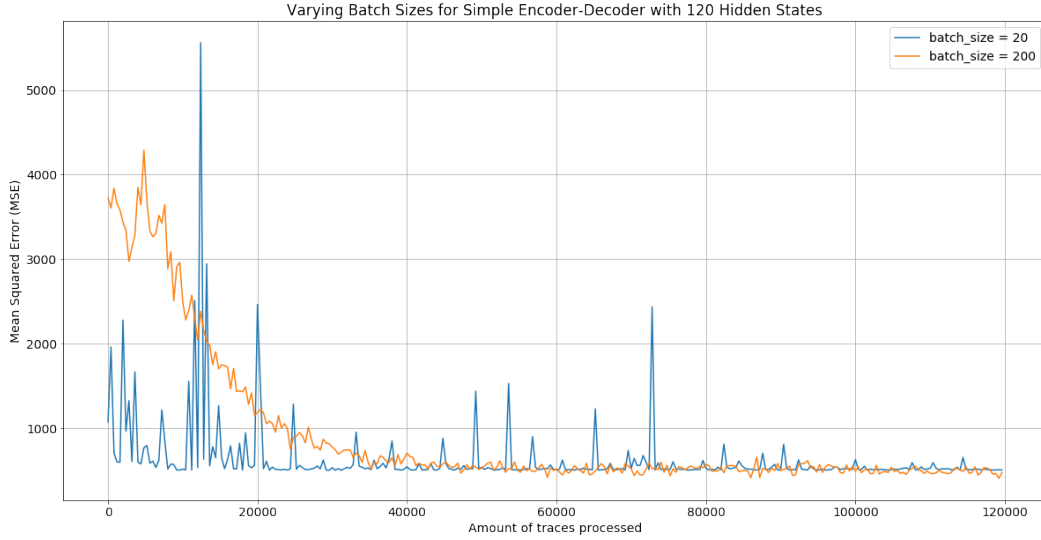


Figure 5.3.1: Scaled MSE over traces processed for different batch sizes.

Next, we vary the learning rate  $\gamma$  from 0.01 to 0.000001 with various optimizers (*adam*, *gradient descent* or *RMSProp*) and loss functions (*mean squared error (MSE)* or *absolute loss (AL)*). For this, we aim to get a learning curve that gradually decreases, the more traces the model processes and eventually slows down once the model reaches a minimum.

After trying a wide variety of different permutations, an *adam optimizer* continuously demonstrated better results. We expected this since adam optimizers are computationally efficient, require a relatively little amount of memory and tend to perform very well with problems that have a large amount of parameters [23], which is ideal since our network can be unrolled to a very large lengths.

Next, we also note that the best quality of data compression was achieved with a *MSE loss function* and a learning rate of 0.000002. Hence, we set  $\lambda = 0.000002$ ,  $b = 200$  and use an adam optimizer with MSE for the rest of our experiments.

## Reversing Traces

### Cutting Traces

### 5.3.2 Architecture Tuning

Now that we have made a decision on which learning parameters to use, we can start changing the architecture of the sequence-to-sequence model to see which ones yield the best results. We do need to note that we might slightly deviate from the previously chosen learning parameters for different models due to memory constraints. But when we do, we will clearly state that this is the case.

#### Hidden States

We first start by examining the amount of hidden states in the network. These directly affects the size of the fingerprints that will be extracted. We do not want these values to be too large since this makes the network more complex and requires more computation. On the other hand, we do not want these values to be too small either since the state vector

might be too small for the network to learn anything useful.

Moreover, we also need to consider the classifiers. The more features we introduce, the more time and data they require to learn the classification task. Whilst if the amount of features is too low, the classifier might not be able to learn how to effectively classify any of the web pages. Hence, we base the amount of hidden states on the amount of features used in previous WF attacks.

Model	Features
SVC [31]	305
SVC [30]	104
RF [16]	150
kNN [53]	3737

Table 5.1: Amount of features for existing attacks.

As can be seen in table 5.1, most attacks tend to have between 100 and 300 features if we do not consider Wang et al.’s attack. Thus we vary the amount of hidden states between 60 to 140 in steps of 20 to see which ones yield the most appropriate results.

For these experiments we train a sequence-to-sequence model with a unidirectional encoder, LSTM cells and without cutting or reversing the traces. The training data consists 120,000 monitored and unmonitored web pages, which are shuffled to avoid overfitting on any specific web page. We only train the model for one epoch, as we seem to have enough data for the model to converge within that epoch. Hence, every sample that the model sees in the figure below is one that it has never seen before. So we can easily determine if the model overfits on specific data.

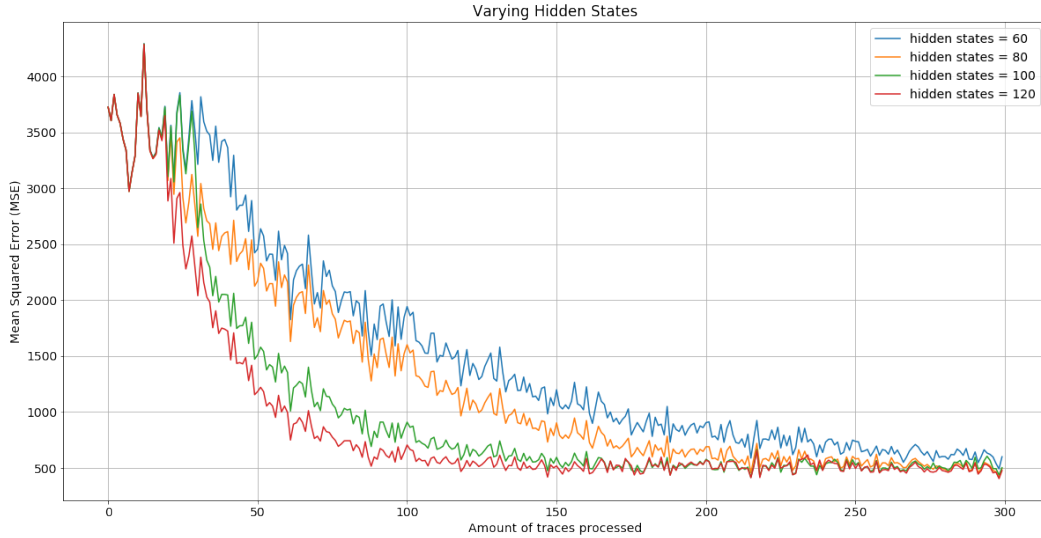


Figure 5.3.2: MSE over the amount of traces processed for varying hidden states.

Figure 5.3.2 clearly shows us that the smaller the amount of hidden states, the faster the network seems to learn the reconstruction task. This was expected because the less hidden states, the less complex the internal representation of the network is. On the other hand, the higher the amount of states, the lower the final error seems to be. Since we aim to compromise between computational complexity and the time it takes to train the model, around 100 hidden states seems to be the most appropriate since it manages to a MSE that is very close to the networks with a higher amount of hidden states.

## Bidirectional Encoder

Rather than using a normal encoder, a bidirectional might be more appropriate for this task. The reasoning behind this is that for a normal encoder, the output at every cell depends on the past. However, there might also be important information in the future that could in fact impact the output at that cell. Thus a bidirectional encoder parses the data in both directions, which means that the output of every cell depends on both the past and the future.

The problem with this method is that it essentially introduces double the amount of free parameters as a normal encoder. Hence, when we perform the analysis, we need to consider whether or not we should take more time waiting to train this model or whether the performance gain is not worth it.

For these experiments, we consider a smaller range of hidden state values from 80 to 120 in steps of 20. Again, for all of these we will be using LSTM cells without cutting or reversing the traces and all of the learning parameters described above and the training set will remain exactly the same.

## LSTM or GRU Cells

So far we've only experimented with LSTM cells since they are the most commonly used ones. However, GRU cells have been introduced more recently. They essentially combine several of the gates and states within LSTM cells to minimize the amount of free parameters and therefore making the model less complex. Consequently, the model should also be able to learn fixed-length representations faster than LSTM cells. But LSTM cells still have a greater expressive power, meaning that they could learn more complex functions.

Here, we train a sequence-to-sequence model with both a unidirectional and bidirectional encoder. These will both have GRU cells with 100 and *TODO : Fillin* hidden states respectively.

Furthermore, we recreate the exact same training conditions as previously. This means that we will not be cutting the traces nor reversing them whilst training on the same dataset of 120,000 web pages with the previously chosen learning parameters.

### 5.3.3 Classifier Performance

We have previously analysed the model's performance based on how well it copies the original input from a fingerprint. But to examine how well our model performs at automatically generating fingerprints, we compare its performance on different existing classifiers with hand-picked features. This means that we choose a set of existing WF attacks and recreate them. Next we run the exact same attack but with the automatically generated features and various scoring statistics.

Note that our results might be slightly lower than in their respective papers since we do not aim to recreate the full attack. Rather than optimizing different hyperparameters, we aim to use these classifiers and the hand-picked features as an indicator as to how well the sequence-to-sequence model performs.

We expect that the automatically generated features will perform worse than the hand-picked ones due to the complexity of the task. To effectively compress a variable-length trace into a fixed-length representation is a very difficult to achieve task and due to the depth of the network, we expect that it got stuck in a local minima. However, we still hope to show that it is in fact possible to automate this feature selection process.

As mentioned in section 3.1, there are two main threat models that we need to consider. The first one is a binary classification task, where the adversary wants to see

whether or not a user is visiting any webpages within a given set. The other threat model involves the adversary having a set of monitored pages, and it wants to know which specific pages the user is visiting in that set. Hence, it is a multiclass classification problem.

Although there are different techniques for evaluating binary and multiclass classification models, we will only use the scoring statistics outlined in section 4.3.3. This allows us for easy comparisons between the different threat models. We do expect that the binary classification models will perform better than the multiclass ones due to the larger amount of options available.

Aforementioned, we have already selected a total of four different existing attacks. We will refer to the first SVC attack by Panchenko et al. [31] as **svc1** and the second one [30] as **svc2**. Whilst we refer the k-fingerprinting attack by Hayes et al. [16] as **RF** and finally the attack by Wang et al. [53] as **kNN**.

## Binary Classification

We first start by analysing the simplest threat model, namely binary classification. For all of the models below, we aimed to extract the exact same hand-picked features as were described in the respective papers to the best of our knowledge.

For training these models, we use an extract from the **GRESCHBACH** dataset with a total of 100 monitored web pages with 70 instances each and 5000 unmonitored web pages. We then split this set into a training and validation set using a stratified split. The training set will contain 90% of the monitored web pages whilst we vary the amount of unmonitored pages to see how the models perform.

After the set is split up into a training and validation set, we perform a *stratified k-fold validation* with  $k = 3$  on the training set. Then finally we train the classifiers on all of the training data and evaluate them on the test set.

The results for the k-fold validation on the training set for the hand-picked features are outlined in table 5.2. Here, we used a total of 10% of the unmonitored data for training.

Model	Accuracy	BDR	TPR	FPR	F1
<b>svc1</b>	$0.91 \pm 0.003$	$0.99 \pm 0.001$	$0.97 \pm 0.001$	$0.07 \pm 0.002$	$0.90 \pm 0.005$
<b>svc2</b>	$0.91 \pm 0.008$	$0.99 \pm 0.001$	$0.95 \pm 0.003$	$0.06 \pm 0.004$	$0.90 \pm 0.008$
<b>RF</b>	$0.93 \pm 0.003$	$0.99 \pm 0.001$	$0.97 \pm 0.006$	$0.05 \pm 0.003$	$0.92 \pm 0.005$
<b>kNN</b>	$0.90 \pm 0.007$	$0.970.003$	$0.94 \pm 0.004$	$0.09 \pm 0.002$	$0.890.004$

Table 5.2: Performance statistics hand-picked features on a binary classification task with k-fold validation.

As expected, the results on the individual folds is relatively high. However, now we will measure their performance when training on the full training set and evaluating on the validation set, whilst changing the amount of unmonitored pages we train the model on. Since bayesian detection rate ultimately determines the feasibility of an attack, we will use that measure for comparison.

## Multiclass Classification

### Different Circumstances

Beside analysing how the sequence-to-sequence model performs on data that was recorded under the same circumstances, it would be interesting to examine how it performs on other data. It has already been shown that the performance of the classifiers is greatly impacted by the network, time and a different TBB version. But that doesn't necessarily mean that

<b>Model</b>	<b>Accuracy</b>	<b>BDR</b>	<b>TPR</b>	<b>FPR</b>	<b>F1</b>
<b>svc1</b>	$0.73 \pm 0.005$	$0.81 \pm 0.007$	$0.81 \pm 0.010$	$0.22 \pm 0.015$	$0.73 \pm 0.005$
<b>svc2</b>	$0.74 \pm 0.013$	$0.84 \pm 0.006$	$0.76 \pm 0.016$	$0.16 \pm 0.009$	$0.74 \pm 0.013$
<b>RF</b>	$0.83 \pm 0.014$	$0.93 \pm 0.014$	$0.80 \pm 0.007$	$0.06 \pm 0.01$	$0.83 \pm 0.014$
<b>kNN</b>					

Table 5.3: Performance statistics hand-picked features on a multiclass classification task.

our fingerprint extraction model is impacted similarly.

If our sequence-to-sequence model is not impacted by these flaws, an adversary would only need to train the model once and then it could continue to use it and only retrain the classifiers. But if it were to be impacted, the retraining process would become significantly slower.

To test this premise, we use a model that we previously trained on the same 120,000 web pages within the **GRESCHBACH** dataset. Next, we extract the fingerprints from the Tor cells within the **WANG14** dataset using this model, train a set of classifiers on these fingerprints and note down their performance.

## 5.4 Unit Tests

On top of evaluating the results, we also needed to ensure that the results were in fact the the code behaves as we expect it to. For this we use unit tests. Some of the code, such as the sequence-to-sequence model is difficult to test but we can still test all of the preprocessing to see if that is correct. For this we use Python’s standard `unittest` module [49]. The reason for this choice is that it is flexible and the standard unit testing framework, which means it is commonly used.

On top of unit tests, *Travis* was also used [48]. Travis is a popular tool, that has an easy integration with Github, for continuous integration. Therefore, every time a commit is pushed to the remote repository, Travis runs all of the tests automatically. If one of the tests fails, Travis then automatically notifies all the contributors.

Finally, to check if our tests cover our entire codebase, *codecov* is used [10]. This tool automatically checks how much of the codebase all of the unit tests cover. At the time of writing, the coverage is 93%. The bits that aren’t covered by unit tests, such as the Tensorflow implementation of the sequence-to-sequence model, have been carefully examined to see if they behave as expected by using the Tensorflow debugger [43].

# Chapter 6

## Conclusion

### 6.1 Future Works

This work shows that the classifiers currently still seem to perform better with hand-picked features rather than automatically generated ones but there is still much room for future improvements. Here we consider several different manners in how we can improve or extend this work. Although we definitely will not cover all the different possible extensions, we try to list the most interesting ones.

As previously mentioned in section 4.3.3, we could add a *softmax layer* on top of a decoder in a trained sequence-to-sequence model. Not only would this allow us to perform the classification with the sequence-to-sequence model, but it would also allow us to analyse how evidence affects the classification. Since you would technically only need one softmax layer, after the fingerprint has been extracted. But having one after every cell, allows us to see how different packets change the prediction of our model. This could then be used as a tool for analysis which features the model actually extracts.

There have also been a variety of different defenses, some of which have been outlined in section 2.2.3. Some works have examined the effectiveness of their attack, when these defenses were in fact used [16, 53]. It would be interesting to see if the sequence-to-sequence model might still be able to effectively extract fingerprints, even with these defenses deployed. This could include both training the model on data where the defense was deployed or training it on normal data and analysing whether it can still extract the fingerprints if the defense is deployed during the attack stage.

Juarez et al. have already shown that WF attacks suffer from the rapid changing nature of the content of web pages [21]. Thus on top of analysing how defenses impact the attack, we could also potentially analyse how the performance of the fingerprint extraction process is affected over time. We have already shown that the model is still successful when extracting fingerprints from other datasets. However, this is not fully show that the sequence-to-sequence model is not affected by content changes within web pages. This could be fully examined by collecting our own data over a period of time and see how the performance of a trained sequence-to-sequence model changes. If the performance is not affected, we could save a large amount of time retraining the fingerprint extractor.

We could also potentially research the possibility that training our model with data collected over time and under different circumstances would also make the model more robust. Since technically, the more different training instances it sees, the better it should get at identifying features. Additionally, we could also investigate how well the sequence-to-sequence model performs when collecting features when given more realistic user behavior. Hence, rather than visiting one page and waiting a certain amount of time before loading the next one, the data can be more realistic such as where the user has multiple tabs open at the same time.

On top of training the model with more realistic browsing data, we could also evaluate its performance for *Tor hidden services*. This is a protocol for Tor users to hide their location while offering a service such as hosting a website [46]. There is already evidence that these services can be classifier using a WF attack [16] but it would be interesting to see how our model would perform on this data.



Rather than extending this work by using more of different kinds of data, we could also improve the sequence-to-sequence model. Currently, one of the main weaknesses is that the traces can be very long, which in turn makes our model very deep. We solved this issue here by cutting the traces after a certain amount of time since most the first part of the trace carries more information than the latter. However, this might not be the ideal solution. There might be another solution or perhaps even another model that does not have this weakness but still manages to map variable-length sequences into a fixed-length representation.

Due to time constraints and limited Tensorflow support, we also did not examine the use of regularization. There are several works that show that several techniques such as *L2 regularization*, *dropout layers* and *batch normalization* are promising techniques [20, 24]. Hence, it could potentially help to reduce overfitting within the sequence-to-sequence model.

## 6.2 Final Thoughts

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# Appendix A

## System Manual

Here, we outline the technical details of the project such that the development of the system could be continued by a third party.

First, we will look at the tools required to run the system then we will look at the overall system and how specific modules connect together. Finally, we explain the specific components and outline how you can contribute to this project.

All of the code is hosted in a private repository on Github. If given access, it can be cloned from the following url (<https://github.com/AxelGoetz/website-fingerprinting.git>).

### 1 Tools Required and Installation Instructions

Most of the project has been written in Python, more specifically, everything has been designed and tested with version 3.6. Our Travis build has also been configured to run all of the tests with Python 3.5, which means that we still get notified if anything breaks under a specific Python version. Some of the code has also been written in Golang, those parts have been adapted from an implementation of Wang et al.'s kNN attack for performance reasons [53, 36].

Next, after the correct version of Python and Golang are installed, we also require a *virtual environment* to manage all of our python dependencies. If you have `pip` on your system, this can easily installed as follows:

```
pip install virtualenv
```

After the virtual environment is installed, go to the project directory and run the following commands to first create the environment and then activate it.

```
virtualenv venv  
source venv/bin/activate
```

At any time, you can go out of the virtual environment by running `deactivate`. After you are in the environment, we need to install a list of dependencies, which can be found in the `requirements.txt` file. This can be done by running:

```
pip install -r requirements.txt
```

This will install a list of dependencies but the main ones are *Tensorflow v1.1*, *numpy*, *sklearn* and *scipy*. There are a couple others but they are not used for major components.

If you plan to use the GPU support on Tensorflow, there are a couple more steps that need to be taken such as installing *CUDA 8* and *cuDNN v5.1*. These instructions can be found on the Tensorflow site<sup>1</sup>. Our project relied on an Amazon EC2 p2.xlarge with one NVIDIA K80 GPU but the code can be run on any GPU card with CUDA compute capability 3.0 or higher [43].

---

<sup>1</sup><https://www.tensorflow.org/install>

In order to run the experiments, the datasets will have to be downloaded next. The main dataset used is GRESCHBACH<sup>2</sup> but we also use WANG14<sup>3</sup>. These should be put in a data folder in the main directory where the directory containing the GRESCHBACH data, should be renamed to `cells` and the WANG14 data to `WANG14_cells`.

## 2 System Overview

Figure 4.4.1 already shows the overall structure of the and how different components interact. Next, in section 4.4 we also explain what the individual sections do. Hence, here we describe the same using a *directory tree*.

```

/
├── attacks - Contains the code for all of the classifiers
├── data
│   ├── cells - All of the individual cell files from the GRESCHBACH dataset
├── feature_extraction - All of the code to extract the hand-picked features
│   from different models
├── feature_generation - Perform automatic feature generation
│   ├── autoencoder - Implementation of the autoencoder, how to run it and how
│   │   to extract features
│   │   ├── autoencoder - Autoencoder class
│   │   ├── feature_extraction - Used to extract features after model has been
│   │   │   trained
│   │   └── train_autoencoder - Runnable file for training
│   └── seq2seq - Implementation of the sequence-to-sequence model, how to run
│       it and how to extract features
│       ├── seq2seq - Sequence-to-sequence class
│       ├── feature_extraction - Used to extract features after model has been
│       │   trained
│       └── train_seq2seq - Runnable file for training
├── report - The latex and pdf files for this report
├── run_models - Provides the infrastructure to run a specific attack in the
│   attacks directory
├── tests - The unit tests to test everything
├── gitignore
├── travis
├── LICENSE
├── README - More information on the project and how to install/run everything
└── requirements - The python packages required

```

<sup>2</sup><https://nymity.ch/tor-dns/#data>

<sup>3</sup><https://cs.uwaterloo.ca/~t55wang/wf.html>

### 3 Components

Here how the individual components work and how to potentially extend them.

#### Attacks

This components implements all of the classifiers that are used to compare the hand-picked features with the automatically generated ones. Each of these classes needs to implement the interface, which has a `fit` and `predict` method. This is similar to the machine learning models provided in *sklearn* [38]. The fit method takes an  $X$  and  $y$  input and changes its internal structure such that it minimizes the error. The predict method on the other hand just takes  $X$  as its input and returns a vector  $y$  of what it predicts the input represents.

The models should also have a `is_multiclass` flag, which represents the fact whether the classifier performs a binary or multiclass classification task.

#### Feature Extraction

All of the logic to extract the hand-picked features for different attacks is in this module. If a new attack is added, a new file should be created in this directory, with the code to extract all the required features from a given trace. Next, this function needs to be added to the list in the `feature_extraction.py` file. This file essentially goes over all the traces within the `data/cells` directory and stores the extracted features within the `data` directory.

#### Feature Generation

This is one of the main components of this work, containing the code to automatically generate features. Currently, we have two models defined, namely a sequence-to-sequence model, which can be found in `seq2seq` and an autoencoder in `autoencoder`. In these directories, we first have a file, containing a class that defines a model and then two more files, which are executables, used to train and extract the features (`train_<model_name>.py` and `feature_extraction.py`).

When running the training files, after every epoch, the infrastructure will save the computational graph such that training can be continued even if it is interrupted. The name of these files depend on the Tensorflow version and the model you are training but generally look like `<model_name>_model.meta`.

Finally, all of the logic to actually extract the features is in the `feature_extraction.py` files, which again stores the fingerprints in the `data` directory.

If you wish to implement a new model, add a new directory within the `feature_generation` folder with the model name. Then create three new files, which contain the model definition, training and extracting code. The model just needs to implement two main functions such that it can be used by the rest of the infrastructure. These are `train_on_copy_task` and `get_vector_representations`, which both take some data as their input and either train the model or extract features.

## Run Models

All of the infrastructure to actually run the classifiers is defined within this module. First, it defines the logic to preprocess all of the necessary data and perform the *k-fold validation*. Next, it also defines all of the scoring methods and how to actually run the models. The main logic is actually in the `run_models.py` file, which allows the user to tune certain parameters such that they can run different models.

Whenever a new classifier is added to the `attacks` directory, it should also be added within the `run_models.py` file such that a user knows that it can be run.

## Unit Tests

For unit testing, we use the standard `unittest` module, which is very simple to use. To create more tests, create a new file in the `tests` directory, which starts with test and then the name of what you will be testing. Next, create a class within that file, which extends the `unittest.TestCase` class and add the methods of what you would like to test.

Then to run all of the unit tests within the `tests` directory, simply run:

```
python -m unittest discover
```

## 4 Contributing

There are a variety of different extensions possible, some of which are outlined in section 6.1. If you decide to implement one of these, we use a very standard git workflow so if you would like to contribute, follow these steps:

1. Fork the project repo.
2. Create a branch with the name of the particular improvement/extension that you will be working on.
3. After you are done, make sure that you run all of the tests and check if everything still works.
4. Submit a pull request from your branch to the master branch and make sure all of the tests pass on Travis.

If you discover an issue or want to work on an extension, please create an issue on our Github issues page to let people know that you are working on this particular extension.

## Style Guide

When contributing, please do note that we try to adhere to the *PEP 8*<sup>4</sup> style guide for our python code [32]. This is done for consistency reasons and readability of the code.

Although it is not necessary to adhere to all of the guidelines, we strongly suggest for you to adhere to the basics. If the code within a pull request is unreadable or does not adhere to any of the standards, there is a very strong chance that it will not be merged.

---

<sup>4</sup><https://www.python.org/dev/peps/pep-0008/>



# Appendix B

## User Manual

In the following section, we explain how the provided code can be used to run the experiments.

Firstly, in order to run the experiments, we require data. Any dataset you want can be used, as long as each cell is represented as a Tor cell, where the SENDMEs have been removed (the same format as in table 2.1). However, for this specific project we will be using two datasets, namely GRESCHBACH<sup>1</sup> and WANG14<sup>2</sup>. Both of these need to be placed in the `data` directory.

After this, you need to install and initialize a virtual environment and install the necessary dependencies as outlined in appendix A.

Next, there are four main scripts that need to be executed to perform the WF attack:

1. `train_<model_name>.py`
2. `feature_generation.py`
3. `feature_extraction.py`
4. `run_models.py`

All of these scripts take several command-line parameters, which are outlined below. These could also be displayed if you run the scripts as follows:

```
python <script_name>.py --help
```

### Train Model

As previously mentioned, we first need to train the fingerprint extraction model. This can be done by executing either of the commands below in the virtual environment, depending on which model you would like to train:

```
python feature_generation/seq2seq/train_seq2seq.py
python feature_generation/autoencoder/train_autoencoder.py
```

Both of these take several command line parameters in order to change the behavior of the models. All of these are outlined below:

---

<sup>1</sup><https://nymity.ch/tor-dns/#data>

<sup>2</sup><https://cs.uwaterloo.ca/~t55wang/wf.html>

Parameter	Type	Default	Description
batch_size	Integer	100	The size of each mini-batch.
bidirectional	Boolean	False	If true, the model will use a bidirectional encoder and a normal one otherwise.
encoder_hidden_states	Integer	120	The amount of hidden states in each RNN cell. The size of the fingerprints depends on this value. $len(fingerprint) = 2 \times encoder\_hidden\_states$
cell_type	String	LSTM	Which specific type of cell to use. Currently only support LSTM and GRU.
reverse_traces	Boolean	False	If true, reverses the traces and leaves them untouched otherwise. This should not be used when <b>bidirectional</b> is true.
max_time_diff	Float	Infinite	The maximum time difference ( <i>in seconds</i> ) after which you start cutting the traces. For instance, if set to 1, all of the traces will be cut after one second.
extension	String	.cell	The extension of the Tor cell files. We expect that they are in the following format <code>&lt;webpage_id&gt;-&lt;instance&gt;.&lt;extension&gt;</code> .
learning_rate	Float	0.000002	The learning rate used whilst training.

Table B.1: Parameters for the `train_seq2seq.py` file.

Parameter	Type	Default	Description
batch_size	Integer	100	The size of each mini-batch.
extension	String	.cell	The extension of the Tor cell files. We expect that they are in the following format <code>&lt;webpage_id&gt;-&lt;instance&gt;.&lt;extension&gt;</code> .
learning_rate	Float	0.0001	The learning rate used whilst training.
activation_func	String	sigmoid	The activation function used for the neurons.
layers	List	[1500, 500, 100]	The sizes of the respective layers in the encoder and decoder.

Table B.2: Parameters for the `train_autoencoder.py` file.

For example to run a simple encoder-decoder with GRU cells, a batch size of 200 and 200 hidden states, run the following command:

```
python feature_generation/seq2seq/train_seq2seq.py --batch_size
    ↪ 200 --encoder_hidden_states 200 --cell_type "GRU"
```

After running this, a couple files should be created in the main directory. First of all `loss_track.pkl`, which is a pickled file, containing the object that represents the loss over time. Next, there should also be a couple `<model_name>_model` files with different extensions, which contain the saved computational graph. Finally, it also creates a `X_test` and `y_test` file in the `data` directory. These contains the paths and the labels to the files, which were not used for training.

## Feature Generation

After the fingerprint extraction has been trained, the features need to be extracted, which can be achieved with the `feature_generation.py` module. Since we do not want to perform any testing on the same data as we trained the model on, it only extracts fingerprints from the traces in the `X_test` file.

Next, these features are stored in either the `data/seq2seq_cells` or `data/ae_cells` directory with a `.cellf` extension.

Most of the flags here are the same as in the previous section. Hence, we will only list the new arguments.

Parameter	Type	Default	Description
<code>graph_file</code>	String	<code>&lt;model_name&gt;_model</code>	The name of where you saved the graph. You should not need to change this, except if you change the graph name in the code.

Table B.3: Extra parameters for the `feature_generation.py` file.

For instance, to extract features from the model that we previously trained, we can run:

```
python feature_generation/seq2seq/train_seq2seq.py --batch_size
  ↪ 200 --encoder_hidden_states 200 --cell_type "GRU"
```

## Feature Extraction

We then want to compare these automatically generated features with the hand-picked ones. These can be extracted using the `feature_extraction.py` script, which again has several parameters. After this script is done running, the features should be stored in the appropriate folders within the `data` directory. Again, all of the files with the extracted features will have the `cellf` extension.

Parameter	Type	Default	Description
<code>all_files</code>	Boolean	False	If true, it generates features for all cells and otherwise just the ones in the <code>X_test</code> file.
<code>extension</code>	String	<code>.cell</code>	Represents the extension of the cell files.

Table B.4: Parameters for the `feature_extraction.py` file.

For example:

```
python feature_extraction.py --extension ".cells"
```

## Run Models

Finally, we can run the classifiers on all the extracted features using the `run_model.py` script. After finishing the k fold validation, the model then prints out the different scoring statistics.

Parameter	Type	Default	Description
model	String	kNN	Which model to run, the options are <i>kNN</i> , <i>random_forest</i> , <i>svc1</i> and <i>svc2</i> .
dir_name	String	Name of handpicked directory	If specified, it will use the features in the given directory, otherwise the standard ones for a specific model.
is_multiclass	Boolean	True	If true, trains the classifier on a multiclass task and binary otherwise.
extension	String	.cell	Represents the extension of the cell files.

Table B.5: Parameters for the `run_model.py` file.

For example, to run a random forest model with automatically generated fingerprints from the sequence-to-sequence model on a multiclass problem:

```
python run_model.py --model "random_forest" --dir_name "  
    ↪ seq2seq_cells"
```

# Appendix C

## Project Plan

### 1 Problem Statement

Anonymity networks like Tor use what is called onion routing where each layer in the onion represent a new layer of encryption. This allows Tor users to freely browse the web without an ISP, government, or anyone else that might be able to sniff the traffic before the first Tor node to see which websites or services the user is accessing. However even with various layers of encryption, an attacker might still be able to infer which web page a client is browsing by performing a *website fingerprinting attack*. The attack often uses machine learning to identify several trends in the network traffic such as the number of packets per second, their size, etc. But most of these attacks rely on a trial and error process of picking the features. Hence, there is no guarantee that the features used are the most appropriate ones or even any good at all. Therefore this project will analyse the use deep learning techniques such as stacked autoencoders to automatically identify features and test their effectiveness compared to the hand-picked ones in various different models.

### 2 Aims and Objectives

The aims and objectives for this project are as follows:

1. **Aim:** Critically review the effectiveness of current website fingerprinting attacks.  
**Objectives:** 1. Analyse various models that are currently used in fingerprinting attacks. 2. See if there are any flaws in the reasoning or the experimentation. 3. Examine how would a small percentage of false positives impacts a potential attack. 4. Analyse how the rapid changing nature of some web pages would impact the attack.
2. **Aim:** Generate features automatically using deep learning techniques for a website fingerprinting attack.  
**Objectives:** 1. Examine different deep learning feature selection methods such as stack autoencoders. 2. Pick the most appropriate method for a website fingerprinting attack. 3. Collect the necessary data to train the feature selection method. This includes a dataset that is collected over a short period of time (days) and another one that would be collected over an extended period of time (weeks). 4. Extract a set of features using this data. 5. Compare these features to existing hand-picked ones.

3. **Aim:** Train existing models with the automatically generated features and test their effectiveness compared to hand-picked ones.

**Objectives:** 1. Identify various models that could be used to test the new features with. 2. Implement the models. 3. Identify various sets of hand-picked features to compare the automatically generated features with. 4. Train those models using both hand-crafted features and the generated features. 5. Compare the effectiveness of the generated features to hand-picked ones in those models in a closed-world environment. 6. Compare their effectiveness in an open-world scenario. 7. Analyse if the feature selection technique can find persistent features that are spread across a period of time and study if this helps with the classification over time. 8. Compare the effectiveness of the automatically generated features when a user uses various common defenses against website fingerprinting like camouflage. 9. Test the attack on tor hidden services as opposed to websites. 10. Investigate an appropriate technique for evaluating the result. 11. Analyse which features tend to be the most informative (highest entropy).

### 3 Deliverables

The project aims to produce the following deliverables:

1. Summary of website fingerprinting attacks. This includes any related work and an analysis how effective a fingerprinting attack could be (see Aim 1).
2. An analysis of the most appropriate automatic feature generation model.
3. A dataset to train both the feature generation model and the models used for the website fingerprinting attack.
4. Fully documented source code for generating the features and the models used for the attack.
5. A strategy for testing the models.
6. An analysis of using the generated features compared to hand-picked one using different models. This includes how the feature generation process might be able to identify persistent features that are spread across a period of time.

## **4 Work Plan**

### **Project start to end October (4 Weeks)**

- Research current website fingerprinting attacks.
- Research various method for automatic feature selection.

### **Mid-October to mid-November (4 Weeks)**

- Refine aims and objectives.
- Further research into using automatic feature selection for website fingerprinting attacks.

### **November (4 weeks)**

- Collect necessary data.
- Initial experimentation with feature selection.

### **End November to mid-January (8 weeks)**

- Implement feature selection.
- Implementation of various models used for attacks.
- Research on how to evaluate the effectiveness of a model.
- Work on the Interim Report.

### **Mid-January to mid-February (4 weeks)**

- Perform tests and evaluate the performance.

### **Mid-February to end of March (6 weeks)**

- Work on Final Report.

# Appendix D

## Interim Report

### 1 Current Progress

We have updated some requirements that were mentioned in the project plan. Rather than first collecting a large dataset of website traces over a long period of time, we decided to start with existing datasets to perform some experiments. Then later, if we get the opportunity to do so, we can still collect our own data.

Given the nature of the challenge, a majority of the time will be spend on researching models that perform automatic feature selection. After careful examination, there are only two couple deep learning methods that seem to be appropriate, a *RNN Encoder Decoder* or a *Bidirectional RNN Encoder*. We also considered using a *Stacked Autoencoder* but it did not seem to be appropriate as it requires a fixed-length vector as an input. Hence, if we were to use it, we either have to pad or compress the traces, which are both not elegant solutions. In addition to simply training these models on the data, we might perform *denoising* on the them, which essentially means learning the models to distinguish uncorrupted data from corrupted data. This final step allows us to fine tune the encoders to get a consistent performance even if some of the data is noisy.

Although we have started experimenting with some of these models, they have not been fully implemented. However we are still on schedule for doing so.

Finally, we have also selected a set of previously successful website fingerprinting attacks. An environment has been set up, some of these models have been implemented and the infrastructure is in place to extract a set of hand-picked features from the raw data. This will allow us to quickly perform a performance comparison with the manually engineered features and the automatically selected ones.

### 2 Remaining Work

The first priority is to fully implement a *RNN Encoder Decoder* or a *Bidirectional RNN Encoder* and fine tune it. Next, we have to perform the analysis on how appropriate our automatically engineered features are compared to hand-picked ones. So this includes finishing the implementation of existing models, and the infrastructure to extract the features.

Next, we need to pick a set of criteria that we will use to compare the predictive power of several models. Using these criteria we will then have to perform a thorough analysis of how the automatically engineered features compared to the hand-picked ones.

Finally, after all of the above has been completed, we will focus on finished writing the final report. If there is still time remaining, we might still try to collect our own data over an extended period of time