

Feature Selection for Brain Computer Interface using Genetic Algorithm

by

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Dear Professor Brünig

In accordance with the requirements of the Degree of Master of Engineering Science in the School of Information Technology and Electrical Engineering, I submit the following thesis entitled

“Feature Selection for Brain Computer Interface using Genetic Algorithm”

The thesis was performed under the supervision of Assoc. Prof. Marcus Gallagher. I declare that the work submitted in the thesis is my own, except as acknowledged in the text and footnotes, and that it has not previously been submitted for a degree at the University of Queensland or any other institution.

Yours sincerely



Timothy Mahoney

**Acknowledgments**

I would like to acknowledge the following people for their support and contribution over the course of this Master of Engineering Science thesis:

**Assoc. Prof. Marcus Gallagher** – as the direct supervisor of this thesis project, Marcus has provided valuable guidance, steering the focus and scope to ensure a well-defined and valuable result.

**Brendan Keane (BSc (Hons I)) and Nicholas Bland (BPsycSc (Hons I)) –** both academics were paramount in the labelling of EEG artefact components for the AAR filter system. Their contribution involved ascertaining what elements of the frequency and spatial distribution of components were consistent with artefactual phenomena. These key features were used by the author to manually label over 3000 individual components for classification training.

# Abstract

Advances in computational power and machine learning techniques have enthused global interest in the development of Brain Computer Interfacing technology. Features relevant to neural event classification include temporal patterns in a variety of frequency bandwidths across many spatially separated electrodes. Selecting an appropriate subset of features from such a large set of candidates could reduce the computational complexity and hardware requirements of such a system, thereby improving the systems feasibility. This thesis aims to satisfy three criteria: use Genetic Algorithm (GA) to establish and heuristically optimal subset of features for classification of motor imagery events relating to left and right hand movement, ensure this subset significantly reduces the number of features and has reasonably low classification error, and the entire process should be theoretically implementable as an online BCI system in that it is automated, computationally efficient and robust.

Motor imagery data from the PhysioNet Database was successfully filtered with an original Automated Artefact Removal implementation which classifies and removes artefact components using independent component analysis and a trained artificial neural network. Features were segmented from the clean data across the 64 channels, into three frequency bands and over six temporal samples, resulting in 738 features. GA and greedy backward elimination feature selectors were implemented as wrappers with an SVM classifier. GA was found to outperform greedy backward elimination in terms of classification error, however GA included over 300 features. We developed two GA alterations to encourage the selection of fewer features: penalised genetic algorithm (PGA) and a hybrid genetic algorithm with greedy backward elimination (GAwGBE). PGA did not perform particularly well despite variations in the penalty rate. GAwGBE provides a reasonable solution of 18.6% classification error with just 52 features (reduced from 738), resulting in a solution including just 28 of the original 64 channels. We state that this solution does satisfy the thesis aims. We conclude that the results of this thesis, as with many related publications before it, are an indication that continued research into this field can bring this technology out of the realm of science-fiction and into reality.

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# List of Acronyms

AAR *automated artefact removal*

ANN *artificial neural network*

BCI *brain computer interface*

ECG *electrocardiography*

EEG *electroencephalography*

EMG *electromyography*

EOG *electrooculorgraphy*

ERD/ERS *event-related de/synchronisation*

GA *genetic algorithm*

GAwGBE *genetic algorithm with greedy backward elimination*

GBE *greedy backward elimination*

GFS *greedy forward selection*

ICA *independent component analysis*

MI *motor imagery*

MM *motor movement*

PGA *penalised genetic algorithm*

SL *surface Laplacian*

SVM *support vector machine*

# Introduction

Advances in computational power and machine learning techniques have enthused global interest in the development of Brain Computer Interfacing (BCI) technology. With applications ranging from motor and sensory prosthetics, to communication and even connection to the internet of things, BCI advancement has the potential to provide significant benefits to society. Recently, Facebook has begun development of a BCI to enable typing via thought [1], engineers at Johns Hopkins University have created a BCI capable of controlling an upper limb prosthesis [2], and even Elon Musk’s latest venture is in BCI technology with his startup company *Neuralink* [3].

Currently however, effectively classifying neural data in an online and non-invasive scenario is no trivial problem. Features relevant to neural events include temporal patterns in a variety of frequency bandwidths across many spatially separated electrodes. Due to the high spatial resolution and frequency bandwidth of non-invasive electroencephalographic (EEG) signals, one hurdle slowing BCI development is determining what data is useful and what can be ignored. Selecting an appropriate subset of features from such a large set of candidates could reduce the computational complexity and hardware requirements of such a system, thereby improving the system’s feasibility.

## Aims and Scope

This project examines various methods of feature selection to produce a heuristically optimal subset of features using the natural selection-based optimisation technique Genetic Algorithm (GA), for the purpose of motor-imagery event classification from non-invasive neural data (EEG). The specific aims can be summarised in three points:

1. Use Genetic Algorithm to establish a heuristically optimal subset of features from the full EEG feature set.
2. The subset should both significantly reduce the number of features *and* result in low classification error
3. The data sampling and processing methods should simulate an online BCI system such that it should be: automated, computationally efficient and robust to slow sampling.

Specifically, after filtering and artefact removal, we extract features consisting of temporal data spanning standard EEG frequency bands from 64-channel EEG recordings. The extracted features are passed to the Genetic Algorithm selector which is implemented as a wrapper, such that the metric used to evaluate a feature set’s fitness is classification error with a Support Vector Machine (SVM).

The dataset being used was created in 2004 by the BCI2000 instrumentation system and contributed to the PhysioNet database [4] [5]. The dataset contains over 1500 one- and two-minute EEG recordings from 109 participants. The system used 64 channel EEG with a sampling rate of 160 Hz. It consisted of 14 experimental runs including two baseline recordings (eyes open, then closed) and three repeats of four tasks. The datasets being considered for this project are of Task 2: the subject *imagines* closing their left or right fists (motor imagery, MI) for a period in response to visual cues. We pose the task as a binary classification problem, with labels corresponding to left (L) or right (R) MI events. The aim of the BCI for this project is to classify these events using the EEG recordings.

This dataset has been used previously for classification problems. Alomari et al. [6] used a different approach with a focus on feature extraction to classify left/right hand MI events. This thesis will use the results given by Alomari et al. as a benchmark to ascertain the level of success of its findings, while providing justification for any contrast in performance.

MATLAB will be the primary software package being used to import, sample, optimise and classify the data, a brief explanation of the key functions and scripts written for and provided with this thesis is available in Appendix A. MATLAB is used because the author is familiar with the software package and there is a substantial pool of supporting documentation available for correctly implementing most, if not all, of the processes involved with this project.

It should be noted that there are assumptions being made in this project. Previous studies have investigated the effects of user specific parameter selection under the motivation that EEG responses can be dependent on the subject [7]. Due to the nature of the dataset, implementing the BCI in this way may produce less meaningful results as the number of experimental runs per person is too small. Lastly, the focus topic for this project is the optimisation methods being implemented, hence the classification algorithms are simply used to evaluate the performance of the features selected and will not be investigated in detail.

Having clearly defined the aims and scope of this document, we will next explore in detail any background material required for understanding the underlying concepts within the thesis. Then the methods used are defined in a natural and intuitive progression, outlining and justifying any design decisions or adaptations made over the course of the project. Briefly we display the results of the implemented methods, followed by an in-depth discussion of their significance. We then compare the risk mitigation and project timeline with those proposed in the initial project proposal, before reaching a conclusion regarding the success of the paper.

# Background

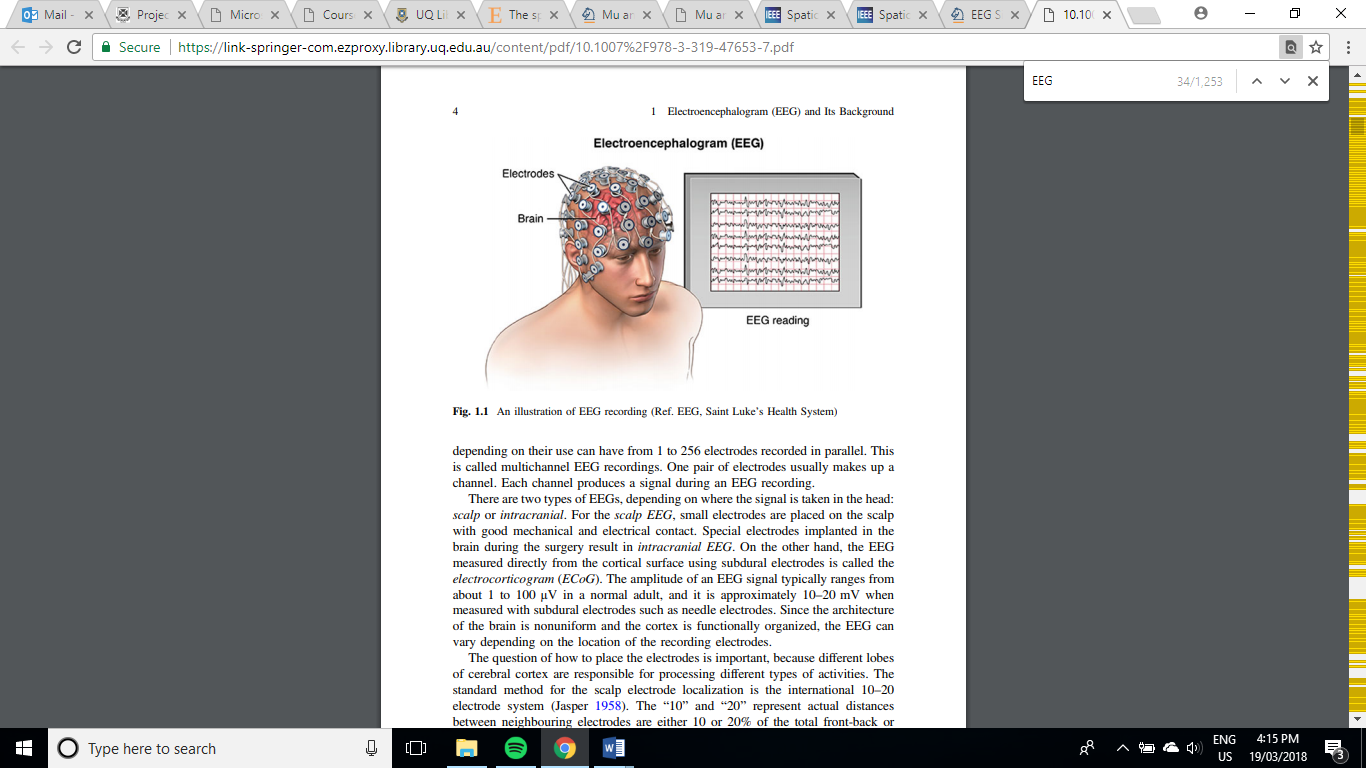
BCI systems are the culmination of knowledge across fields spanning biology, neuroscience, electrical and biomedical signal processing, and machine learning to name a few. The following section covers, in detail, the large amount of background knowledge required for this project due to the multidisciplinary nature of BCI technology.

## Electroencephalography

Electroencephalography (EEG) is the recording of action potentials originating from the brain. It is most often used in the study of brain functionality or the diagnoses of neurological disorders [8]. EEG signals are sampled using several small electrodes placed on the scalp, which can detect minute potential differences non-invasively on the surface of the skin [8]. Through proper filtering and signal processing techniques, the summation of electrical activity from local neurons can be isolated and displayed allowing professionals to examine neural activity pertaining to each electrode [8].

The exact placement of the electrodes is a generally subject to the standardised International 10-20 System, whereby the distance between left-right and front-back electrodes is either 10% or 20% of the total skull distance [9]. The electrodes are labelled according to the area of the brain, or lobe, that they’re reading from. These are pre-frontal (Fp), frontal (F), temporal (T), parietal (P), occipital (O) and central (C) as seen in Figure 1 [8] [9]. The number of electrodes in total varies, systems can sample anywhere from 1 to 256 EEG signals in parallel [8].

Figure 1(left) Standard EEG labelling of each lobe. (right) Examples of EEG positioning and recordings [8]



## Cortical Homunculus

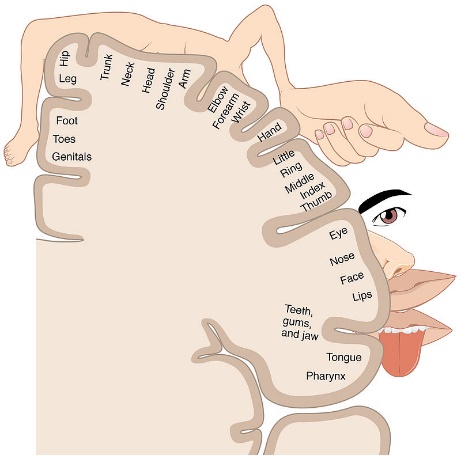
Neural sensory and motor function are generally thought to be confined to certain spatial locations within the brain. A cortical homunculus is an illustration describing this spatial separation as observed from the brains surface along a coronal cross section at either the sensory or motor cortex. Considering the homunculus depicted in Figure 2, we can see the tissue associated with hand control is a relatively large portion, approximately half way down the cortex. While we do aim to find an appropriate and small feature set using primarily GA, for the purposes of this project we can perform some preliminary reduction of the feature space by neglecting the peripheral channels of the EEG as we can, with confidence, say that these channels are unlikely to relate to MI events associated with hand movement.

Figure 2 The motor cortical homunculus displays the brain tissue segments associated with certain motor functions.

## 2.1.2 Frequency Bands

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| Band | Delta | Theta | Mu | Alpha | Beta | Gamma |
| Frequency (Hz) | 0.5 - 4 | 4 – 7 | 7-12 | 11-15 | 15-31 | >31 |

The frequency components of EEG are considered some of the most important criteria for interpreting brain function [8]. The bandwidth of these components is commonly defined as per Table 1.

Table 1 Standard EEG frequency bands

The presence or absence of each component is generally associated with a particular type of brain function or state [8]. For instance, the three bands which are commonly associated with sensory-motor function are alpha, beta and mu waves, and are hence the three frequency bands being used for feature extraction in this project.

## 2.1.3 Event Related De/Synchronisation

There is evidence to suggest that in addition to the amplitude of each bandwidth indicating a neurological event, temporal patterns referred to as event related de/synchronisation (ERD, ERS) between particular bands may correlate to the mind processing information [10] [11]. ERS occurs when there is an observable increase in neural activity due to the synchronisation of neurons firing action potentials at a given frequency band. Likewise, ERD is the phenomena of de-synchronised neurons cancelling each other out, thereby reducing the observable electrical potential of a given frequency band. ERS and ERD events shown to relate to sensory-motor activity last over a period of seconds. In order to consider this information in the feature set for this project, we include features extracted from previous samples as inputs into our classifier.

## 2.1.4 Artefacts

Signals included in EEG recordings that do not originate from neural potentials are referred to as artefacts. They can be introduced from other physiological events such electromyographic signals (EMG) from muscles, electrooculographic signals (EOG) from eye movements or even electrocardiographic signals (ECG) from the heart. They can also exist as noise from poor electrode contact or 50/60Hz mains coupling. EEG signals can easily be corrupted by artefacts as their amplitude is in the range of 10-100μV, hence filtering techniques have been thoroughly explored in previous studies to establish clean signals for professional examination or further processing [8].

Figure 3 Examples of EEG artefacts and their impact on neural signals. Often artefacts exhibit much larger voltage potential amplitude than EEG, thus resulting in poor signal to noise (SNR) ratio if not removed.

Three commonly used artefact removal methods include:

1. Epoch removal – simply manually remove any epoch samples corrupted with artefacts
2. Manual artefact removal – manually remove components likely to be artefacts before reconstructing the signals
3. Automated artefact removal (AAR) – implement automated classification and removal of artefacts

As this project aims to produce a solution which is capable of online classification, implementing AAR is the ideal method of artefact removal. Artefacts are commonly classified by first separating the n-channel EEG into some k components (k≤n), and then considering features of these components such as their temporal pattern, frequency response and spatial distribution. Components are isolated using Independent Component Analysis (ICA).

ICA is often likened to isolating voices in a crowd. Given a number of spatially separated microphones, ICA is able to isolate specific voices (or components) by combining information from each. In the case of EEG data, we have 64 channels all sampling a symphony of neural information and artefacts. Components isolated from this data may be attributed to either one of these groups, if we can identify which, we can remove artefact components before reconstructing the channels. The automated ICA implementation for this project will be discussed in more detail in the *Methods* section.

## Brain Computer Interface

A brain-computer interface (BCI) aims to establish a pathway between neuropsychological signals and an external device [12]. One potential goal of such an interface is to provide a substitute method of communicating or controlling extremities for people with neuromuscular impairments [8] [12]. The ability to classify mental imagination events would allow patients to regain some degree of freedom whether it be through wheelchair or neuroprosthesis control, or communication. With proper training a person could repurpose their neural pathways for any number of applications to improve quality of life. However, for such BCI systems to be considered a useful alternative control method they need to achieve critical performance requirements for robustness, consistency and safety [12].

Mason and Birch described a general framework for BCI systems as a way of standardising ideas and terms for these devices to aid with their development and comparison [13]. The framework consists of 6 steps [13] [8]:

**Brain activity measurement** - the sampling of the raw EEG data

**Pre-processing** – digital to analogue conversion, initial filtering and artefact removal

**Feature extraction** – retrieve meaningful features of the EEG data which can discriminate between mental states being used to control the system

**Classification** – employ some algorithm which can classify mental states based on the extracted features.

**Translation into a command** – relate the result of classification with a response in the target device.

**Feedback** – sensory feedback to the user not only allows them to know that the device is working, but also helps train their brain to use it.

This project will focuses on the areas from pre-processing up to classification.

Commonly used mental responses for controlling a BCI are motor imagery (MI), where the subject imagines moving without performing the movement, and motor movement (MM), which is the brains response to an actual motor movement being made. Of these two tasks, this project will only consider MI as this medium is inherently more suited to applications such as limb replacement.

## Feature Extraction

One of the most critical elements of a functional BCI is effective feature extraction. As discussed in sections 2.1.1, 2.1.2 and 2.1.3, the features being extracted span frequency, space and time dimensions in order to best utilise known phenomena observable in the EEG data relating to MI events. Prior to feature extraction however, the data undergoes only one more process known as Surface Laplacian (SL).

SL is a transformation commonly used to improve spatial resolution of EEG data [14]. When EEG data is sampled, it is measured as a potential difference between each channel and some reference electrode. This means there is often redundant information common to neighbouring electrodes due to their close proximity. SL is a matrix multiplication which effectively decouples the channels from the reference electrode, transforming the recordings into comparative measurements between neighbours. There are several SL techniques, each with varying complexity as described by Carvalhaes and de Barros [14], however the method used in this project only considers the simple Finite Difference Method as described in the reference and presented in Appendix B. This method makes the assumptions that the scalp surface is locally flat and that electrodes are equidistant [14]. These assumptions are considered reasonable for the purposes of this thesis. SL is the final step of feature extraction before employing feature selection.

## Feature Selection

Consider the case where a 128 channel EEG was divided into its 5 frequency bands and the mean of each band for the previous 5 samples was taken as the feature set, resulting in a 3200 long feature set. A feature vector this large introduces several problems in the classification stage, such as [15]:

1. Over complicating the classification model
2. Long training time
3. Slow real time BCI performance
4. Increased risk of overfitting
5. Curse of dimensionality – small ratio of data to number of features

Feature selection attempts to counter these problems by selecting a subset of features without incurring much loss of information [15]. The selection process follows the premise that certain features may correlate with each other (termed: *redundancy*), whereas others may have little to do with classification so including them within the feature vector results in no information gain (*irrelevance*). Feature selection algorithms are essentially a search problem across candidate feature subsets, for some set which is at least heuristically optimal concerning some evaluation score. Score metrics depend on the specific algorithm being applied, the three types are: filter, wrapper and embedded [16]. Filter algorithms select the feature subset independent of the prediction method with metrics such as correlation [17], whereas wrappers assess the subset using the predictor and embedded methods optimise the subset in the process of training the predictor [16]. This project initially considers the use of a filter feature selector before expanding into methods involving wrappers.

## Computational Optimisation

Computational optimisation uses algorithms to minimise or maximise an objective function. In the case of EEG feature selection, the objective function would be the metrics we chose to optimise for (e.g. minimise redundancy, maximise relevance) over the space of feature subsets. If the space of subsets includes every combination of the feature vector proposed earlier (3200 features), there exists over 10963 combinations of features. Brute force search methods over this space are not computationally feasible and as such optimisation problems involving large spaces over many dimensions have led to the creation of novel search algorithms to find *good enough* solutions, these are referred to as *heuristic* algorithms. These methods aim to computationally balance *exploration* and *exploitation* of the search space. Exploration is the wide spread search of the space, often at random, in order to avoid focussing on what could potentially only be local optima, whereas exploitation attempts to converge on the ideal local solution based on information the algorithm has already discovered. This project implements one such heuristic search method based on the evolutionary phenomena: natural selection. The method is Genetic Algorithm (GA).

## Genetic Algorithm

As is the case with many optimisation methods, GA is an algorithm designed to mimic the principals of a naturally occurring system. The method takes the process of evolution, such that the genetic code of the parents is inherited by the child, and simulates a ‘survival of the fittest’ scenario over a series of generations [18]. With respect to this project, the genetic code of the parents/child would be a combination of EEG features and the measure of ‘fitness’ could be the metrics of redundancy and relevance for that particular combination. The method starts with randomly generated parents in the search space. The next generation is computed with a combination of three *genetic operators* [18]:

**Reproduction operator:** poor solutions are neglected, whereas fit solutions are passed down. The number of candidates is preserved by including more children with strong performance to replace the those lost. The resulting candidates are now part of what is referred to as the *mating pool*. Clearly the reproduction operator does not result in any new solutions, they are formed in the remaining two operators. Passing on the best candidates to the next generation directly is a method of preserving the best solution, these best candidates are referred to as *elite children*.

**Crossover Operator:** there are a few variations of this operator, but generally two candidates (the parents) from the mating pool are chosen randomly and a subsection of their genetic structure is switched to form children. This operation supplies new candidates with genes that have been previously shown to perform well in the reproduction stage, in the hope that they may perform better than their parents.

**Mutation Operator:** the remaining codes are now subject to mutation, where there exists a small probability that one or more of their genes may be altered. This introduces exploration into the algorithm, allowing it to traverse previously unseen areas of the search space in pursuit of better optima.

The scores of each candidate are computed after these operations in order to inform the algorithm how best to manipulate the next generation. Bit strings pertaining to high scores are carried through to next generations, whereas poor solutions are not. New generations are computed continuously until some stopping criteria is met, for example a certain number of generations or a high score threshold. The process is described quite well by Deb, K. who uses the example of minimising materials used to produce a cylinder of set volume as a function of diameter and height [18]. It is worth noting that this algorithm in no way encourages a significant reduction in the number of features selected, only that a high scoring solution be found. Creatively altering or adding to the method is required to achieve this, which is further discussed in the *Methods* section.

## Classification Algorithms

Classifiers are used in this project for AAR and MI event classification. The scope largely consists of investigation into optimisation techniques however, as such exploring and tuning of machine learning aspects of the project are relatively brief. The classifiers used in the project include an Artificial Neural Network (ANN) and a Support Vector Machine (SVM). These methods were chosen as they have been used previously with EEG classification [19] [20] [21] and MATLAB documentation on their implementation is extensive.

## Artificial Neural Network

An ANN mimics the principals of neural pathways in the brain. A single neuron takes weighted inputs from one or more neurons in a previous layer, where it can then signal other neurons in subsequent layers. The weights between the neurons are updated as learning progresses, tuning the network to accept data features as inputs and predict its class based off supervised learning from a training set. A wide variety of network topologies exist, however this project will only be considering simple feedforward networks with a small number of layers. Hyperparameters consist of the number of layers, the number of neurons in each layer, the training function, the performance metric and the stopping criteria.

## Support Vector Machine

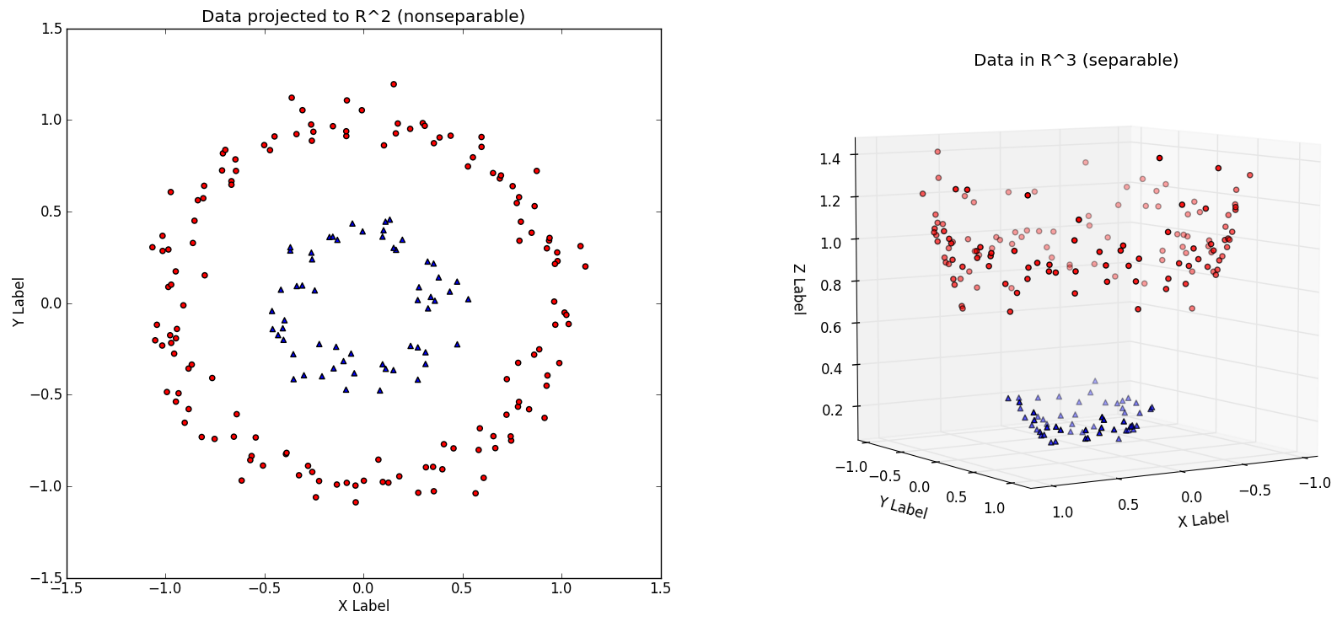
Support Vector Machines are simple discriminant classifiers which construct an optimal hyperplane in the sample space between classes from a training set. The resulting hyperplane is then the discriminant used to classify new data points. Usually, the best hyperplane is taken as the one that maximises the separation of the classes. In cases where the data is not linearly separable, SVM can still work quite well if the data is extended to a higher number of dimensions via some kernel function. Kernel functions are used to map the features to some higher dimension where they are seen as separable and hence can be classified by the SVM, an example of this is shown in Figure 4.

Figure 4 An example of projecting a non-linearly separable 2-dimensional classification problem into a higher 3-dimensional space. In the 3D space the two classes can clearly be separated by a plane.

SVM parameters include the kernel function scale (gamma) and the soft margin penalty scalar (*C*). Variations in gamma relate to the gradient of the kernel function, if the data for a given class is sparse then gamma should be small to reduce the slope of the kernel function describing the data. The soft margin scalar *C* relates to how much the classifier penalises data found outside the soft margin for its class, a larger *C* results in a more strictly trained SVM.

## Previous Related Study

*EEG Classification with Advanced Feature Extraction and Machine Learning* by Alomari et al. uses the same dataset proposed for use with this project [6] [4] [5]. They propose a system which focuses on feature extraction, rather than selection. The features include Energy, Power and Mean metrics for ERD and ERS events of the beta and alpha bands, as well as Movement Related Cortical Potentials (MRCP). The feature set is evaluated using an ANN and SVM. The results show high classification accuracy: 97.1% using SVM and 86.5 using ANN. One limitation of the paper is the length of the epochs used; each sample ends over five seconds after the beginning of the motor event. This means that there would be a classification lag of over five seconds, which is not ideal for a responsive BCI. The paper also makes assumption regarding the sampling frequency of the BCI. Each sample used to train and test the SVM is partitioned as to line up consistently, in reality a BCI may not sample fast enough to perfectly capture the ERS/ERD. Regardless of these limitations, this paper sets a benchmark for this project to aim for and may provide insight into the importance of feature extraction vs. feature selection. Comparison may also highlight limitations in the approaches of both papers despite using the same dataset.

# Methods

The following section contains the methods used and decisions made during the project period. Any results displayed in this section are done so to validate the project before moving to subsequent methods, or to justify an alteration in the approach. Results pertaining to the feature selection methods employed are held out and displayed in the *Results* section. The data is segmented into two second epochs sampled at 2Hz for the entirety of the process in order to simulate an online BCI system as per the third aim of this thesis.

## 3.1 Initial Filtering

Before anything else, we filter the data. We apply a bandpass filter between 0.5-80Hz to eliminate any baseline drift and high frequency noise. We also apply a 60Hz notch filter to reduce mains coupling (USA frequency). Having performed basic filtering, the data is ready for artefact removal.

## 3.2 Automated Artefact Removal

The AAR pathway is illustrated in full in Figure 5. The following sections will discuss each operation chronologically and in detail.

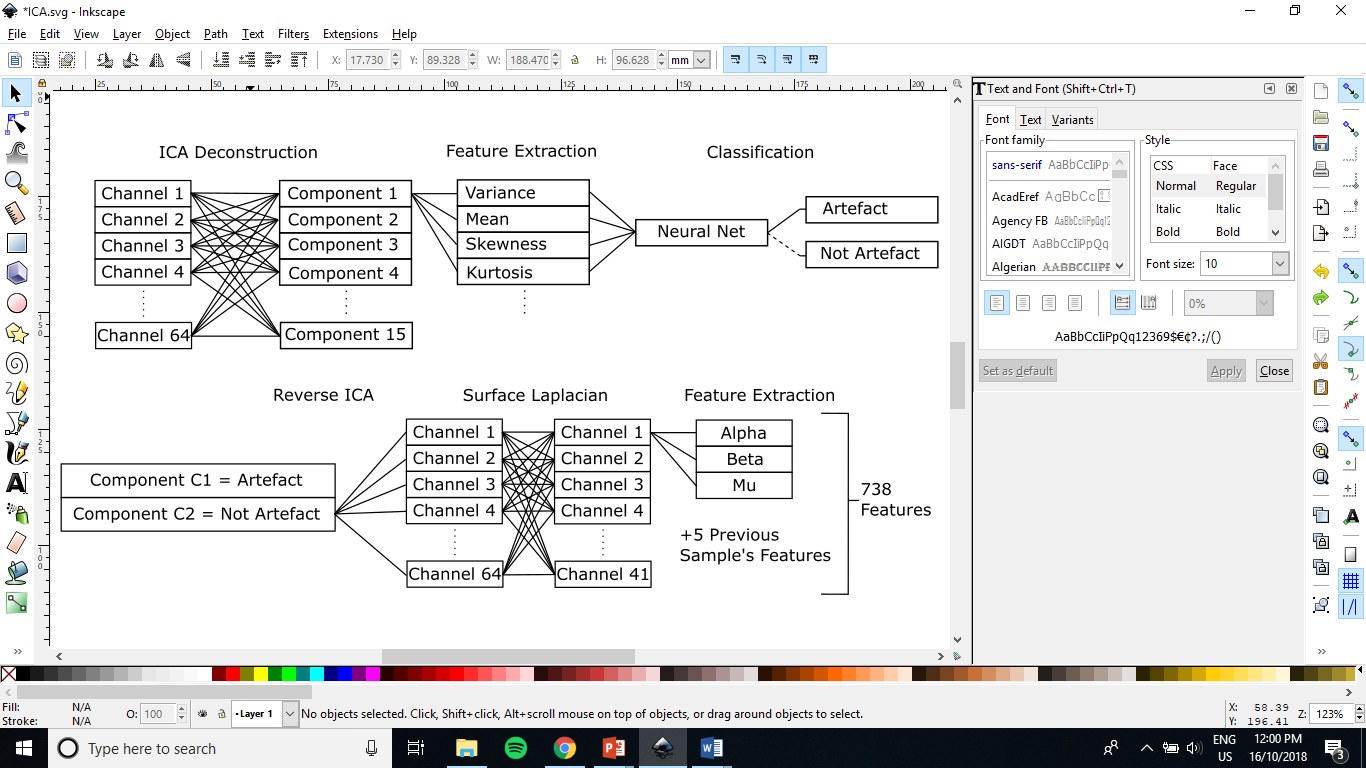


Figure 5 The progression of the data post initial filtering, but prior to classification. This diagram can be separated into three key areas: AAR (covering ICA decomposition up to Reverse ICA), surface Laplacian transformation and feature extraction.

## Independent Component Analysis

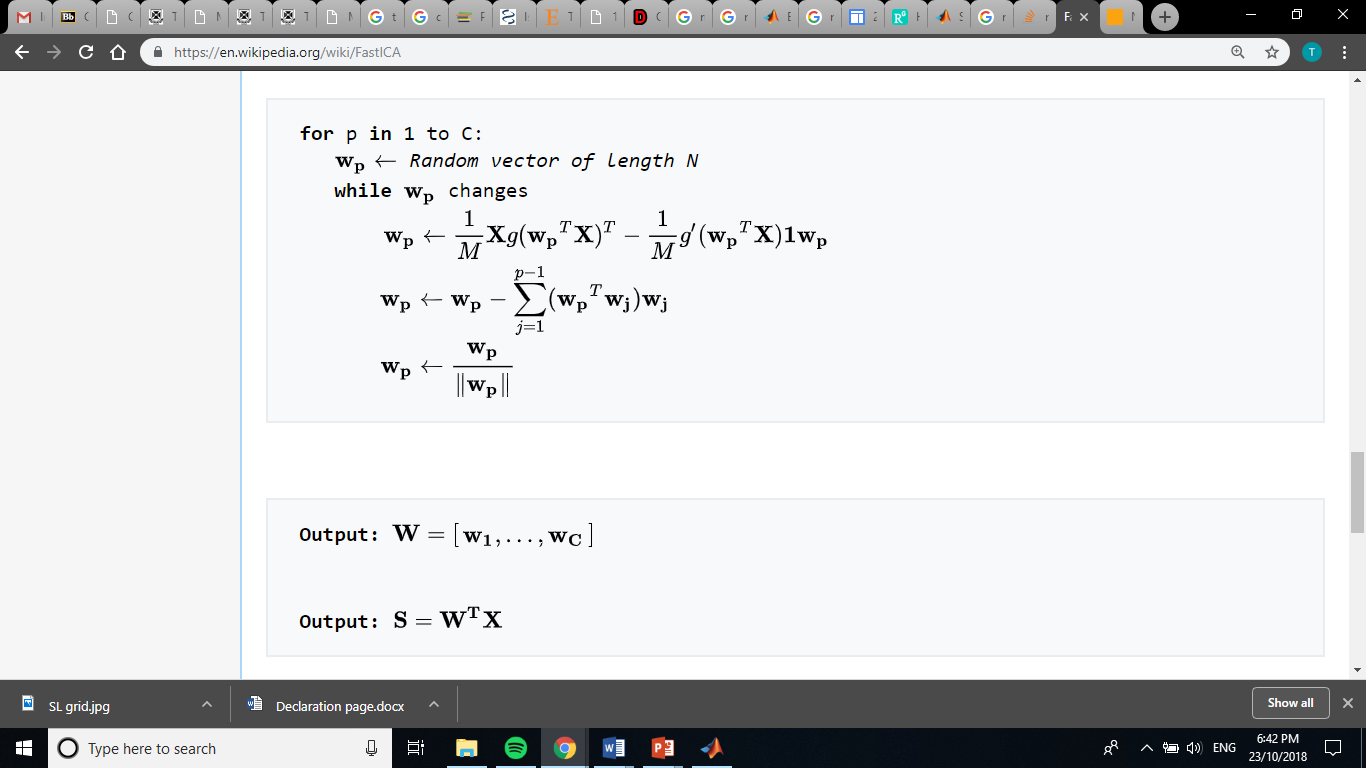
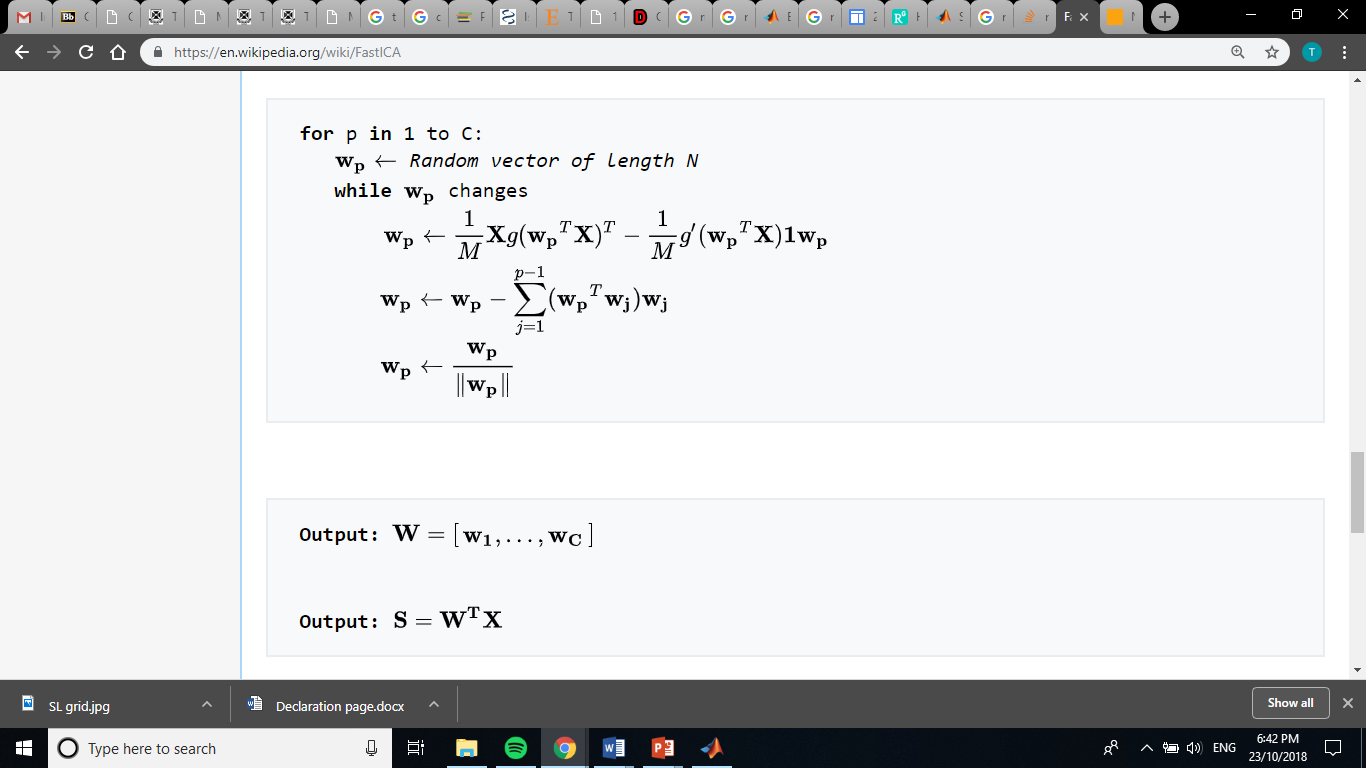
ICA is applied to the data using the FastICA algorithm [22]. First the data must be centred and whitened. Centring the data implies setting the mean of each channel across all samples to zero:

Whitening the data involves transforming the centred data into an uncorrelated set with variance equal to one. This can be achieved through eigenvalue decomposition of the covariance matrix of the centred data, such that:

Where *E* is the matrix of eigen vectors and *D* is the diagonal matrix of eigenvalues.

With our nicely centred and whitened data, we can now apply the FastICA algorithm as shown in Figure 6.

Figure 6 Pseudo code describing the FastICA algorithm. [22]



Where:

*C* is the number of components

*M* is the number of samples

*N* is the number of features

*W* is the component weight vector or unmixing matrix

*S* is the component matrix

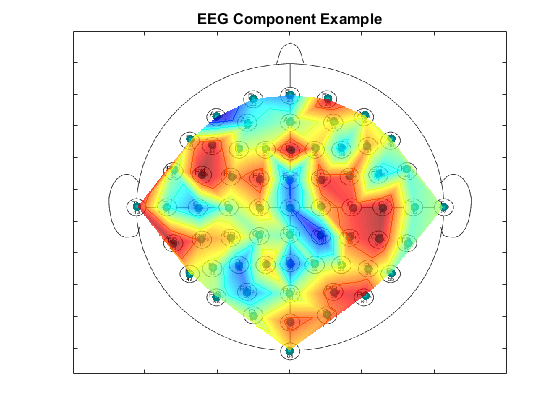
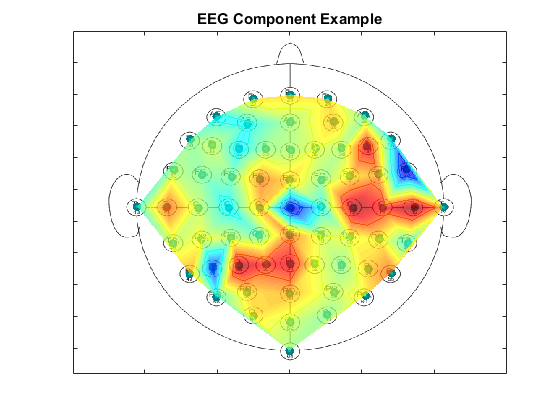
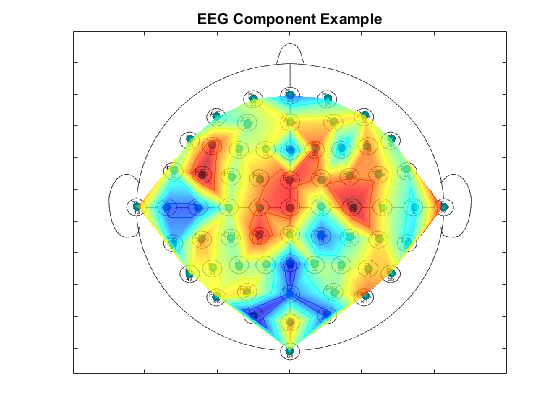
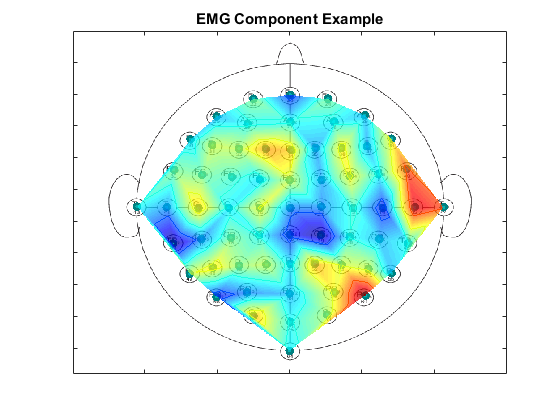
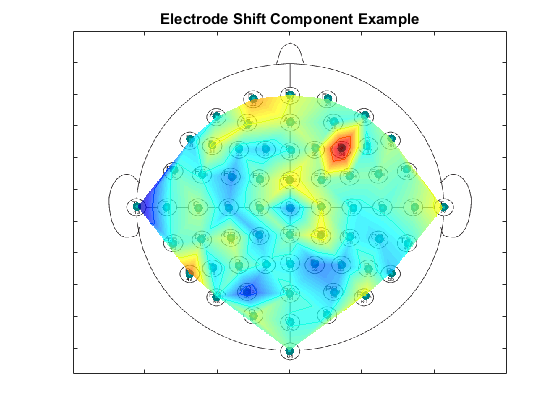
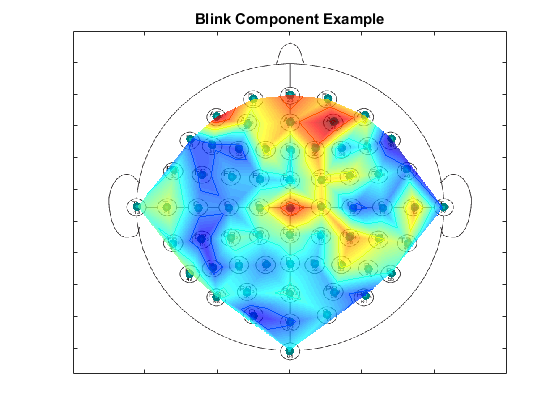
This algorithm is quite simple to implement in MATLAB. In the implementation used for this project, we consider the first 15 (*C*=15) components. The resulting *S* matrix gives us our component set. The next step is to extract features from these components to pass to a classifier. We can speculate what approach to take for feature extraction by examining what experts look for when manually labelling artefact components.

Two PhD candidates studying psychology and neuroscience at the University of Queensland, Brendan Keane and Nicholas Bland, have extensive experience analysing EEG data and identifying artefact components with ICA. Both were consulted to help identify signature attributes of an artefact component for the purposes of this project. Attributes indicating an artefact were summarised as:

1. Components which are spatially condensed to one, or few sparse channels (electrode shift)
2. Components with a frequency response showing peaks at high frequencies (>80Hz) or at USA mains frequency (60Hz)
3. Components with activity localised at peripheral electrodes (EMG) or frontal electrodes (EOG, blinking or eye movement)
4. Components spread over a strong majority if not all the channels (noise)

In the absence of all these observations, a component was labelled as *not* an artefact. Using these guidelines, 3000 components were manually labelled to be used as training data for a classifier to implement AAR. Examples of these components are displayed in Figure 7.

Figure 7 Examples of labelled EEG components. The top three all exhibit spatial patterns consistent with meaningful EEG data. The lower three are typical representations of three artefact types (from left to right): EMG stemming from the right side of the subject, electrode shift from a single channel and EOG artefact from blinking or possible eye movement.



## 3.2.2 Component Feature Extraction

Features were extracted from the components to provide the classifier with the tools to establish similar guidelines. The features are listed in Table 2.

Table 2 List of features extracted from EEG components computed through ICA and the rationale for their inclusion.

|  |  |
| --- | --- |
| Feature | Reasoning |
| Spatial mean of all channels | These features are tailored to inform the classifier of spatially observable artefacts such as those in described by guidelines I, III and IV. The classifier should learn that the amplitude of these features, and variations in comparative amplitudes of some these features can indicate that a component is or is not likely to be an artefact. |
| Spatial mean of peripheral channels |
| Spatial mean of central channels |
| Spatial mean of front channels |
| Spatial Variance |
| Spatial Kurtosis |
| Spatial Skew |
| Frequency response mean 0-80Hz | Features concerning the frequency response are aimed at arming the classifier with knowledge to identify components with high frequency or mains coupling noise, as described by guideline II. |
| Frequency response mean 59-61Hz |
| Frequency response mean >80Hz |

Feature-feature plots displaying the separation of the labelled data are shown in Figure 8. The trends observable support the decision to extract such features for classification. The classifier implemented at the centre of the AAR algorithm was an ANN.

Figure 8 Separation of labelled components when investigating feature-feature relationships of classes: artefact (A) and non-artefact (N). The data shows general trends suggesting that it is indeed somewhat classifiable given the depicted features.



## Artificial Neural Network

The ANN was initialised, trained and tested using code generated from the MATLAB neural network toolbox. The data was separated into train/validate/test datasets (70/15/15%). The network training function used was MATLABs inbuilt Bayesian Regularisation Backpropagation (trainbr). The stopping criteria consisted of the first to occur between: 10,000 epochs, gradient of < 1e-7 or 15 consecutive validation fails. The learning rate was set to 1e-4. The performance metric used is mean squared error. The layout of the network (ie. Number of hidden layers and perceptrons per layer) was tuned using a simple randomised search algorithm:

1. Randomise the number of perceptrons in a three-hidden-layer network between 0 and 8 per layer
2. Train and test the network
3. Keep the network if it produces the *best* *performance* thus far
4. Repeat from step 1.

It should be noted that while the performance metric used by the MATLAB toolbox for training the network was mean squared error, the metric used by the above algorithm in comparing two given networks was the mean between the sensitivity and specificity of the network. Sensitivity and specificity provide information regarding the capability of a classifier should the two classes have different prevalence within the data. In the case of artefact/non-artefact classification, 77% of the samples are labelled as non-artefacts. A classifier could easily achieve 77% accuracy in the trivial case where it classifies *every* sample as a non-artefact, thus classification accuracy alone is not an appropriate performance metric. Carefully considering the metric used has forced this simple algorithm to select networks which may sacrifice successful classification in the more prevalent class to improve classification in the other. This algorithm was left to run for several hours before settling for the current best layout, as shown in Figure 9.

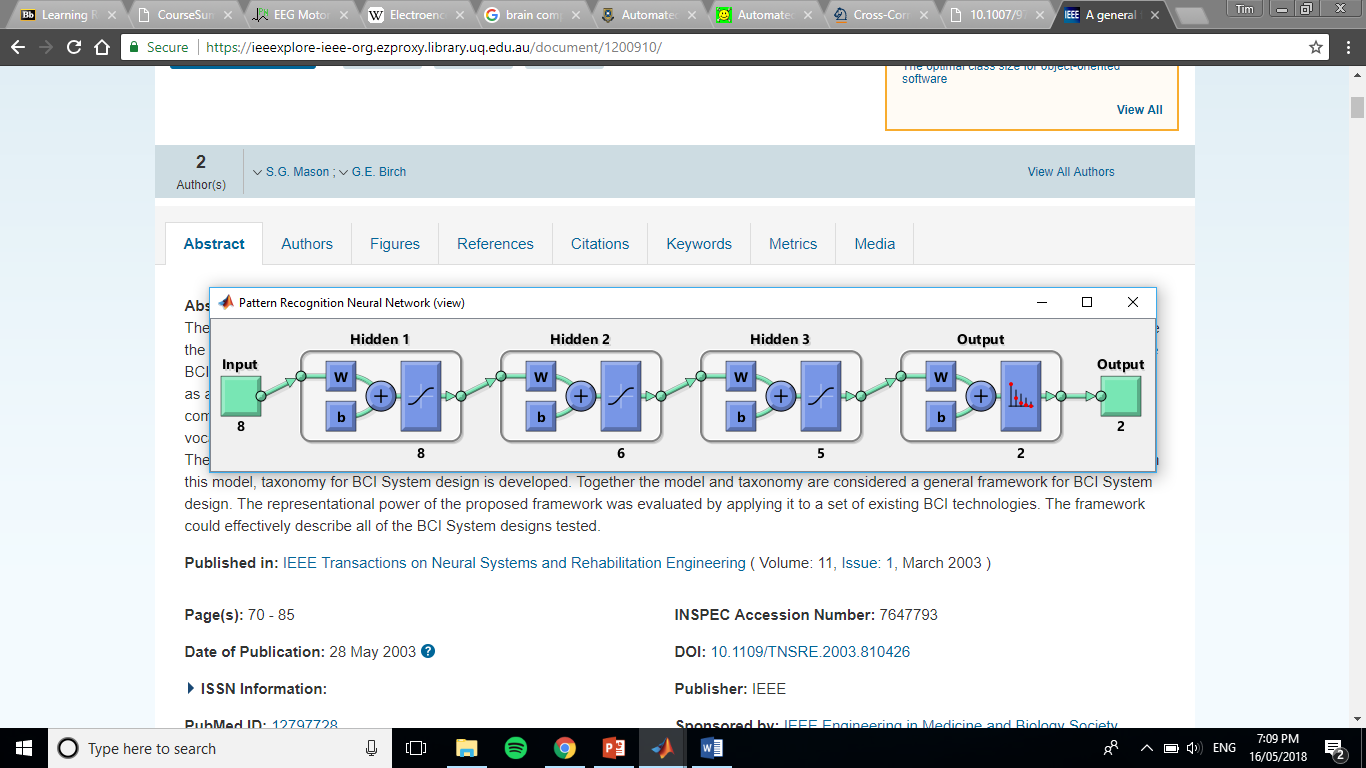


Figure 9 The architecture of the final ANN for use in AAR.

Figure 10 displays the confusion matrix for the network. The network produces reasonable performance (sens=95%, spec=80%). It is approximately 15% better at classifying non-artefact components (Class 1) than artefact components (Class 2). These results are considered reasonable as we would prefer to include data at the risk of it containing artefact, rather than exclude data which could potentially be useful to the BCI. An example of pre and post-AAR filtered data can be viewed in Appendix C.

Figure 10 Confusion matrix for AAR ANN. Class 1 corresponds to a non-artefact, class 2 an artefact..

## Channel Reconstruction

After classifying artefact components, the process of reconstructing the 64 EEG channels from the 15 detected components minus any artefacts is quite simple. We set the rows in the *S* matrix (*component matrix* as per Figure 6) to zero to create what we denote as *Sfiltered*, and recompute the mixed data *X* with the inverse of *WT*, as such:

The resulting *Xfiltered* matrix is our artefact removed 64-channel data, thus concluding the AAR algorithm.

## Surface Laplacian

With our filtered data we apply SL. Applying SL is a simple matrix multiplication, such that:

However, developing the *L* matrix requires applying the Finite Difference Method as shown in Appendix B and by Carvalhaes and de Barros [14] for each channel being included in the final feature set. We only consider the central 41 channels and ignore the peripheral 23, thus *L* becomes a 41x64 matrix. Each row is relating a channel to a scaled combination of only its close neighbours, hence *L* is mostly zeros. The MATLAB script .m file containing the *L* matrix initialisation is included with this thesis (formLS.m). Having performed the SL transformation, we are now ready to extract our features for classifying MI events.

## Feature Extraction

As discussed previously in the *Background* section (2.1.1, 2.1.2 and 2.1.3) we can utilise prior knowledge regarding MI neural activity to limit our feature search space in the time, space and frequency domains. We have already reduced the spatial dimension of the data by performing SL and only considering the central 41 channels. Regarding EEG frequency bands, we are only concerned with those typically relating to MM/MI neural events, ie. mu, alpha and beta. These frequency bands are attained using 10th order Butterworth bandpass filters at the thresholds listed in Table 3. Including temporal patterns within the set allows for the capture of ERD/ERS events which have been previously shown to relate to MM/MI activity. These patterns generally occur over a period of seconds. Thus, we include 2.5 seconds worth of data in each feature set, updated and shifted along with every new sample at 2Hz. If our feature set is given by:

Then each is a single and unique combination of the frequency, time and space dimensions. In this case, (41x3x6). Table 3 illustrates the exact ordering of the feature string.

Table 3 The structure of the feature binary string. Each of the 738 features maps to a unique frequency band, spatial channel and temporal delay.

|  |  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Frequency | Mu | | | | | | | Alpha | | Beta | |
| Channel | 1 | | | | 2 | … | 41 | 1 | … | | 41 |
| Time | t | t+0.5 | … | t+2.5 | t | t+2.5 | t | t+2.5 |
| Feature |  |  |  |  |  |  |  |

Before extracting these features however, we carefully consider how to segment the data. We have continuous recordings (160Hz) which we are subsampling in 2Hz blocks. We also have labelled data providing the exact starting time of the subject being given the cue for an MI event. This presents two options:

1. We specifically segment a temporal block about the exact time of the event (cherry picking) resulting in more consistent data for classification or
2. Sample the entire recording at 2Hz (as a BCI might in an online scenario) and choose epochs about the event, regardless of potentially leading or lagging.

While option I appears tempting, it does not reflect the capability of an online BCI. Segmenting the data in this way implies that the BCI can filter, perform AAR, extract features and classify new data at the same sampling frequency as the EEG hardware (160Hz in this case). This is not feasible with the proposed methods. Thus, we choose option II, which could result in the event leading/lagging the sample as much as 0.25 seconds, potentially causing inconsistencies between the same features for different samples. A BCI system which could perform well despite such a limitation would be a robust solution, satisfying one of the aims of this thesis. It is worth noting that at this point in the process, we leave out 20% of the data to use as a test set for classification later. We cannot use the entire dataset to first optimise our feature set, and then reuse the set for training a classifier as the system has already seen the labels, hence leading to overfitting.

## Feature Selection

Feature selection involves reducing the number of features to utilise relevance and minimise redundancy in the feature set. The problem can be modelled as binary optimisation across *n* dimensions such that our feature set :

Where *n* is the total number of features. To search this unconventional space, we can employ the heuristic search method Genetic Algorithm.

## Genetic Algorithm Implementation

The general framework of GA is relatively constant for most applications: some form of randomised initialisation followed by selection, crossover and mutation operations. However, beyond this there are several degrees of freedom regarding hyperparameters and specific techniques used. Most variable criteria either present a trade off or are more applicable to certain types of problems than others. For instance, having a large candidate pool spans the search space more per generation, but takes longer to compute. K-point crossover is likely to preserve neighbouring genes through generations whereas uniform crossover will arbitrarily pass down from either parent at each gene. Too little mutation will slow the algorithm in terms of exploring potentially unseen features, but too much will undo the exploitative action of crossover and degrade the algorithm essentially into random search. These are all examples of design decisions which influence the performance and efficiency of a GA implementation. Through preliminary experimentation, we determined the specific GA method below to be appropriate for this feature selection problem.

1. Eight candidate sets are sampled at random from the feature space to create the *candidate* pool.
2. Evaluate *candidate* pool scores.
3. Automatically include the two highest scoring candidates (elite children) in the new *candidate* pool
4. Create a *mating* pool which includes two copies of the highest scoring candidate and one of each of the next two highest scoring candidates.
5. Select six sets of two different parents from the *mating* pool at random and perform uniform crossover to produce the remaining six candidates for the new *candidate* pool.
6. Apply mutation with a probability of 0.01 to every candidate except for the best.
7. Repeat from step 2 until reaching a stopping criterion of 1000 generations.

Reasoning for each design choice is provided in Table 4.

Table 4 An in-depth explanation of each design choice for the GA implementation used in this thesis.

|  |  |
| --- | --- |
| Design Choice | Reasoning |
| Eight candidate generation | Eight appeared to be a healthy medium. Eight candidates span the space quite well allowing mutation to search a reasonable number of feature changes per generation whilst also including a good number of crossover combinations. One generation using a wrapper takes approximately four seconds (fewer using filter). |
| Two elite children | Including elite children solidifies high performing candidates into future generations lest they be improved upon. Future generations are encouraged to exploit and build on these feature sets through crossover. |
| The mating pool | Four candidates: two of the best, one of the second, one of the third. This architecture seemed to carry combinations to next generations from high performing candidates without resulting in homogeneity. |
| Two different crossover parents | Obviously different parenthood was enforced to discourage exact clones appearing in subsequent generations. Generations containing copies of candidates (especially high performing candidates) negate the benefits of the crossover operation. |
| Uniform crossover (as opposed to the more common k-point crossover) | There appears to be no reason to group neighbouring feature strings together during crossover. The inclusion or exclusion of each feature is seen as equally as important as any other feature on its own. For this reason, it seems more appropriate to pass on genes with equal probability from each parent at the gene to gene level. |
| Mutation probability of 1% | This parameter was decided through experimentation. Considering a string length of 738, 1% mutation appeared to provide enough variation for searching new solutions through the generations. |
| 1000 generation stopping criterion | This parameter is simply a time constraint. 1000 generations takes a little over an hour to compute. More generations could lead to a better solution with the cost of more computation time. In general, after several hundred generations, candidates with improved performance were far less likely to be found. |

Initially, GA was implemented as a filter feature selector

## Filter Implementation

As discussed in the *Background* section (2.3), filter methods select feature sets using some fitness metric independent of a classifier. These methods are generally faster than wrapper methods, however may result in lower classification performance as the set is not specifically selected for the classifier. In this project, we consider Correlation Feature Selection, which aims to select features with high correlation to the labels (*relevance*) and poor correlation with each other (*redundancy*) [17]. The score balances relevance and redundancy as such:

Where *k* is the number of features, is the mean correlation between the features and the class labels and is the mean inter-correlation between features. We aim to maximise this function as we require features which correlate strongly with the class labels, and poorly with each other.

We can exhaustively evaluate the correlation between features and the labels as well intercorrelation between features as this computation is not too taxing. Given that we have precomputed this information, calculating the score of any feature set is very fast. Before considering GA, we first consider the effectiveness of a simple algorithm to provide a benchmark: Greedy Forward Selection (GFS). The GFS algorithm is as follows:

1. Start with an empty feature set
2. Compute fitness of new *candidate* sets given the addition of a single feature.
3. Next set is the set from *candidates* with the best computed fitness
4. Repeat from 2 until all features are included in the set
5. Select the feature set which best balances fitness and number of included features

The score given by this algorithm over its progression is illustrated in Figure 11. We simply choose the feature set with the highest score as it also yields a reasonably low number of features: 57 reduced down from the original 738.

Figure 11 The progression of the GFS algorithm. The blue circle signifies the chosen point to cease the algorithm due to a peak in score and relatively low number of features.

The correlation score at this point is 0.2936. It is worth noting that this score is not bound between 0 and 1 as with correlation. In fact, its upper limit is dependent on the number of features included, as per the score equation. Given we decide to include 57 features, the Correlation Score space would appear as shown in Figure 12, for varying mean relevance and redundancy correlations.



Figure 12 The score space given we decide to include 57 features. This figure illustrates the score plane should we achieve a range of redundancy and relevance values from the worst-case scenario (redundancy = 1, relevance = 0) to the best-case (redundancy = 0, relevance = 1). The position marked with the red asterisk is the solution given by GFS.

In the case of a perfect solution (,) we achieve a score of 7.5498 (). The best solution found using GFS is marked on this surface by the red asterisk and is far from the optimal solution. The algorithm was able to select a set with low mean redundancy, but also low mean relevance (, ). This poor score could indicate one of two things: that the features poorly represent the classes, given that the correlation score is a ‘good’ measure of this relationship, or that GFS is not an appropriate search algorithm for this problem. At this point, we hope for the latter, and that implementing GA to search for a better feature set can prove this.

Unfortunately, GA is not able to perform as well as GFS. Figure 13 displays the gradual increase in correlation score over the computed generations. Even having increased the stopping criteria to 10,000 (from 1000), GA was unable approach the best score found through GFS. A summary of the results observed using the two methods is shown in Table 5.

Figure 13 The progression of the GA filter feature selector.

Table 5 A summary of the results observed from the filter implementations of GFS and GA. Classification error is given by an SVM, tuned as described in Section 3.5.3.

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
|  | Number of Features (*k*) | Redundancy (*rff*) | Relevance (*rcf*) | Correlation Score | Classification Error |
| GFS | 57 | 0.0364 | 0.0670 | 0.2936 | 0.4011 |
| GA | 249 | 0.0353 | 0.0407 | 0.2054 | 0.3671 |

While redundancy between the features is quite low for both methods, relevance is still far too low to assert any significant relationship between the features and the classes. Despite optimising for correlation, the classification error yielded by each feature set when passed to a tuned SVM[[1]](#footnote-1) is quite poor. In addition, the set found with GFS resulted in a higher error when passed to a classifier than the set found using GA, despite a better correlation score. It seems a higher correlation score does not necessarily indicate better classification performance in this case. Perhaps a filter feature selector with a correlation metric is not an appropriate method for this particular problem. Taking this poor performance into consideration, we decide to change our approach and develop a wrapper feature selector with the hope of reducing classification error by tuning our feature set to a specific classifier.

## Wrapper Implementation

Unlike filter feature selection, a wrapper relies on the performance of some classifier as the metric for some set of features. The classifier being used for this feature selector is an SVM. Tuning the SVM involves appropriately selecting parameters gamma and *C* (as discussed in *Background* Section 2.3). The process for tuning for these parameters is suggested by MATLAB documentation [23] and is outlined below:

* 1. Use MATLABs fitsvm function with default parameters to determine initial gamma.
  2. Adjust gamma and *C* parameters (KernelParameters.KernelScale and KernelParameters.BoxConstraint in MATLAB) through 11 values. Gamma consists of the default gamma found in step 1 scaled by 1e-5 to 1e5, increasing by a factor of 10. *C* is adjusted by starting with 1e-5 and increasing by a factor of 10 to 1e5. At each iteration we perform a 10-fold cross validation and retrieve the 10-fold loss.
  3. The best parameters of the 121 combinations are those which yields the lowest error.

The feature set used for this tuning was the set found previously using the correlation filter with GA. The parameters found to give the lowest error were gamma≈37.5 and *C*=1000. The kernel function used was a gaussian kernel. Having tuned the classifier, we are now ready to implement a wrapper feature selector.

To thoroughly investigate the potential of GA as a feature selector wrapper, we develop four methods:

1. **Greedy Backward Elimination (GBE)** – similar to GFS, GBE begins with a full set of features and greedily eliminates features which result in the lowest reduction in performance, as the name suggests. The progression gives us the freedom to choose at what number of eliminated features we believe the performance of the set to be satisfactory. However, as it is a greedy algorithm, we expect more sophisticated methods to perform better due to the complexity of the problem. GBE is used as a benchmark to compare our three remaining methods involving GA.
2. **Genetic Algorithm** – implemented in the exact same manner as described in *Methods* Section 3.1.1 with classification error being the feature set performance metric. It is worth noting that GA does not attempt to reduce the number of features in the set, which is one of the key aims of this thesis, hence the remaining methods attempt to enthuse feature reduction.
3. **Penalised Genetic Algorithm (PGA)** – we adjust the score metric to include a penalty term proportional to the number of features included in the feature subset, as per the equation below:

If our candidate set is given by:

Then PGA evaluates the error score of the set with the function:

Where *CE* is the classification error given by the train SVM and *R* is some scalar.

The value of *R* is representative of how much we penalise the set for including any given number of features. In fact, the equation implies that should we increase the number of features in the set by one, we would add *R* to the score. Thus, should we want to include some new feature in any candidate set, its inclusion should reduce *CE* by at least a value of *R* for it to be considered a worthy addition. Large values of *R* imply that we strongly oppose large feature sets and care little for classification error in comparison, whereas the opposite is true for small *R*. Selecting an appropriate value is key to the outcome of GA.

* 1. **Genetic Algorithm with Greedy Backward Elimination (GAwGBE)** – first GA is used to find a suitable feature set (as with method 2). The resulting feature set is then the starting point for GBE. This hybrid method aims to utilise the heuristic search of GA and then reduce the set using GBE, with the added benefit of allowing us to select how many features to eliminate before we cause too much error in classification. To the authors knowledge, this variation of GA has not been presented in any previous works, much less in the context of this thesis.

The performance of these methods is presented in the *Results* section.

# Results

The following section presents the results of the four wrapper feature selection methods outlined in *Methods* Section 3.5.3. Data is separated into a training and test set prior to feature selection so as not to overfit the classifier. Classification error against the test data for all methods is provided in Section 4.4.

## 4.1 Greedy Backward Elimination

Figure 14 illustrates the progression of GBE. As features are eliminated, we can observe the classification error reducing. This is likely due to many features being included in the set which do not relate to the classes, and hence the trained classifier poorly generalises the classes in terms of these features. As such features are eliminated the classification error reduces to below 24% before increasing again as the feature set becomes too small to appropriately separate the classes. We choose the point marked by a red circle as the *best* solution provided by the algorithm as it has the lowest classification error and a significant number of eliminated features. Despite a reasonable solution, we hope to do better with the remaining, more sophisticated methods.

Figure 14 The progression of GBE implemented as a wrapper. The red circle signifies the chosen point to cease the elimination as it exhibits a low error and low number of included features.



## Genetic Algorithm

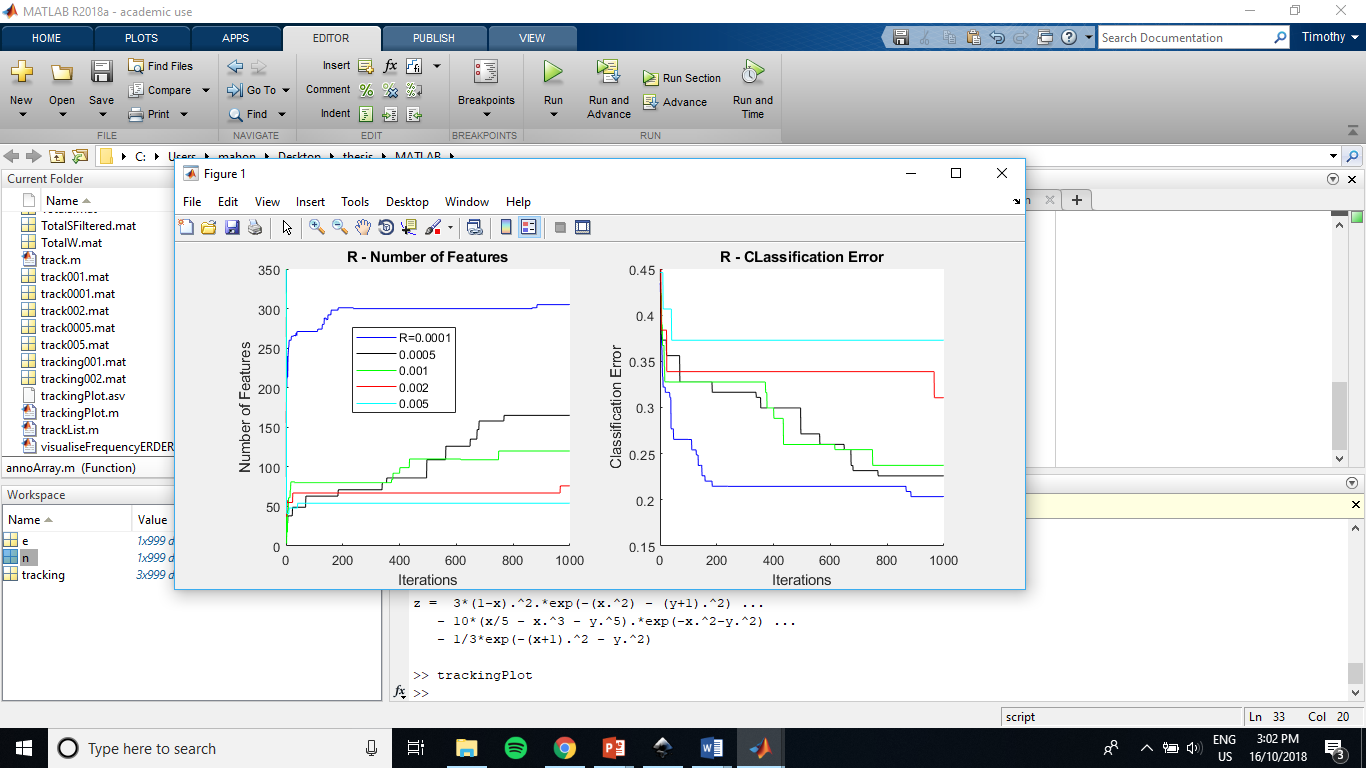
Figure 15 displays the classification error and number of features given by the best candidate found using GA up to its stopping criteria of 1000 generations. We were able to achieve a lower classification error than the set provided by GBE, however the number of features in the set is over 300. We can also see that the reduction of classification error slows considerably past the first 100 generations, indicating that better solutions become increasingly sparse beyond <25% classification error. With the remaining two methods, our goals are to improve upon this solution by reducing the size of the set but maintaining reasonable classification error.

Figure 15 The progression of the GA wrapper method. The blue line indicates the number of features as per the left axis, whereas the red line indicates the classification as per the right axis.

## 4.3 Penalised Genetic Algorithm

Penalised Genetic Algorithm offers limited improvement upon standard GA. Figure 16 depicts the classification error and number of features for various values of *R*. It is clear that *R* behaves in the way described in *Methods* Section 3.5.3. As *R* increases, we can see the number of features in the candidate set drops over the generations, however this comes at a clear cost of classification error. Our best solution in terms of classification error is given by our smallest *R* (=0.001) with an error≈0.2 but over 300 features. Whereas when *R*=0.005 we have a set with almost 50 features but an error>0.35. Of those trialled, there appears to be no overwhelming *R* values which produces a solution significantly better in terms of both of these parameters, however for the sake of comparison we consider the case where *R*=0.001 to be a reasonable medium.

Figure 16 The progression of PGA. The left figure depicts the number of features in the best candidate solution over each generation. The right shows the classification error. Each line shows the progression for a GA with a scoring penalty coefficient R equal to the number given by the legend.



## 4.4 Genetic Algorithm with Greedy Backward Elimination

Genetic Algorithm followed by Greedy Backward Elimination produced the best solution of the four methods in terms of both classification error and number of features. Figure 17 illustrates the progression of the two algorithms. GA finds a set with a reasonable error but high number of features, as shown previously in the results of method 2 (Section 4.2). The addition of GBE then produces a linear decrease in the number of features while maintaining, and for the majority of the progression *improving*, the error rate of the SVM. Indeed, in this case GBE was able to eliminate over 200 features from the set found with GA before error starts to increase. The combination of these two algorithms appears to have performed better than both alone, and the proposed altered method PGA.



Figure 17 The progression of the GAwGBE algorithm. The first stage in white is the GA stage. The second stage in grey is the GBE stage continuation.

## 4.5 Summary

Table 6 displays a summary of the performance of the methods for ease of comparison. We include the PGA score for the best candidate found using each method given *R*=0.001.

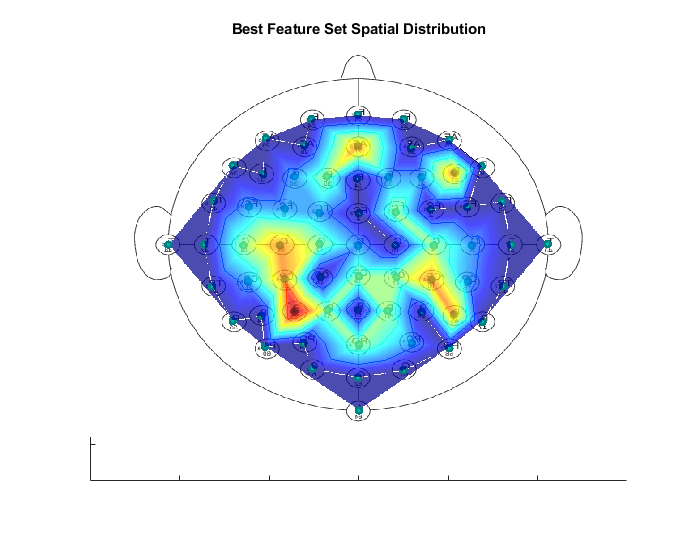
Table 6 A summary of the Wrapper implementation results, showcasing classification error evaluated using an SVM. We include a score metric calculated using the PGA score formula with R=0.001.

|  |  |  |  |
| --- | --- | --- | --- |
|  | Classification Error (Test Data) | Number of Features | Score (R=0.001) |
| GBE | 0.2350 | 80 | 0.3150 |
| GA | 0.2091 | 328 | 0.5371 |
| Penalised GA | 0.2410 | 131 | 0.3720 |
| GAwGBE | 0.1864 | 52 | 0.2384 |

## 4.6 The Feature Set

While no predictions or preferences are in place regarding what specific features are included within the reduced set, it may be interesting to examine their distribution across the time, space and frequency domains and briefly comment on any apparent trends. This distribution is depicted in Figure 18.

Figure 18 The distribution of the final feature set over the frequency (18.1 top-left), time (18.2 top-right) and space (18.3 bottom) domains. More features being included from a given channel correspond to a ‘hotter’ colour in the spatial distribution figure.



Regarding the frequency distribution, all the included bands (mu, alpha and beta) are often related to MI events and hence it is no surprise that all are included to some degree. While beta band being significantly more present in the set than the other two bands is interesting, any conclusions which could be drawn from this are not the focus of this thesis and are hence outside of its scope. The wide spread of temporal features presents a limitation for the use of this feature set as a BCI system: temporal information required for classification exists as far as 2.5 seconds after the MI event. This results in a lag in the classification. Reducing this lag could be an area of further research, as discussed later in Section 5.6. The spatial distribution as seen in Figure 18.3 depicts a certain likeness to what one might expect given the mirrored channels approximately clustered about the *hand* section near the motor cortex. Recalling the cortical homunculus discussed in *Background* section 2.1.1, we might expect to see channels localised around the motor cortex (approximately central horizontal band) and slightly offset from the vertical centre, which is marginally similar to what we see in the figure. Stipulating further than this is, again, not the focus of this thesis. It is important to note that the number of channels has been reduced from the original 64 down to 28, satisfying one of the aims of this thesis by reducing hardware requirements for such a BCI system.

# Discussion

This section aims to critically analyse the results presented, particularly focusing on the effectiveness of the wrapper methods as they are considered the prominent centre of this thesis. We critically analyse differences between the results of this thesis and those given by Alomari’s *et al*. with the same dataset. Beyond discussion of the results, we include a reflection on the effectiveness of risk management methods employed as per the original project proposal, as well as the project plan and any variations to it over the course of this thesis.

## Filter Feature Selector Performance Review

The filter feature selection method performed poorly. We observed not only that both GFS and GA could not find a feature set with significant relevance to the classes, but also that when it came to classification error, the set found using GA produced lower error despite having a lower correlation score. These observations are not in line with what we might expect but can be explained quite simply: correlation is not a suitable metric to evaluate this particular problem. Correlation appears to poorly represent the fitness of a given feature because the measure being used in this case (Pearson Correlation Coefficient) is the strength of a *linear* relationship between two variables. While it is difficult to illustrate the relationship between each feature and their labels given the number of dimensions, the inherent complexity of the MI event classification problem implies that this relationship is indeed non-linear. Hence employing a linear metric to asses our feature set’s fitness seems to be an inappropriate approach, a conclusion which is supported through the experimental results. The one advantage the filter method had over wrappers was the significantly reduced computation time, however this benefit is null considering its poor performance. A wrapper method including a classifier capable of separating non-linear classes, such as an SVM, provides a more suitable solution.

## 5.2 Wrapper Feature Selector Performance Review

First let’s consider the basic methods: GBE and standard GA. GBE performed as expected, providing the freedom to choose when to cease the elimination and accept the classification error of the reduced set. The method provided us with far superior classification performance than the filter implementations (error=0.23 vs 0.40/0.37), supporting the choice to wrapper feature selection. The error it produced was low, but high enough still that we hoped for further improvement through more sophisticated heuristic search methods. Lower error was indeed found with standard GA (0.21 vs. 0.23). This confirmed two things: the solution found by GBE was not globally optimal with respect to classification error, and that our particular GA implementation is capable of finding better solutions. The former point may have been stipulated prior to this discovery, but its confirmation means that the introduction of sophisticated search algorithms such as GA to solve this feature selection problem is justified. The latter point, that the GA algorithm design as described in *Methods* Section 3.5.1 is indeed valid, attests to the success of this thesis in applying such an optimisation technique in this context. This is a key milestone in the project, however one aim remains to be included in the optimisation method, and that is encoding the reduction of features.

This encoding was achieved in two distinct ways by the remaining two wrapper feature selection methods. Recall PGA introduced a penalty term into the score metric, and GAwBGE selected a starting set for BGE using GA. Of these two methods, GAwBGE was able to achieve lower classification error *and* a lower number of features. We can stipulate as to why this has occurred. Regarding PGA, consider the case where our generation includes a candidate with a reasonable error and number of features. This candidate is chosen as a parent for crossover and undergoes mutation, producing a child candidate. One of the following four scenarios may occur:

1. The child candidate includes more features and results in a higher error score
2. The child candidate includes more features and results in a lower error score
3. The child candidate includes fewer features and results in a higher error score
4. The child candidate includes fewer features and results in a lower error score

and recall the equation for PGA score,

We know that the score is proportionally related to the number of features in the set. Thus, with the inclusion of more features, it is more likely that we will result in a higher (worse) score. The reverse is also true. With this, we could say that scenarios 1 and 4 are more likely than 2 and 3, as 2 and 3 would require the child candidate to improve their score by a value more than the number of features gained multiplied by *R*, and vice versa if features are lost. Next consider that during the operations of crossover and mutation, there is no preference concerning the passing on, or manipulation of, a one bit over a zero bit. Considering mutation especially, as the algorithm reduces the feature size down due to penalisation in the score calculation, it is more likely to increase the number of features in the next generation (scenario 1) due to large population of zeros in the bit string. These operations increase the feature set without knowledge of the effect this has on the score, thus searching candidates which are likely to result in poor solutions and slowing the progress of the algorithm. Altering these operations in order to encourage a reduced number of features in subsequent generations could be an interesting area of further research, as is discussed further in Section 5.6. We believe that given enough time, PGA would result in more reasonable solutions, however without more informed operations this method does not appear as useful as GAwGBE.

In comparison, GAwGBE performed quite well. We can see that GA provides a decent classification error for a starting point for GBE. The number of features which can be eliminated with GBE without resulting in increased error is substantial (approximately 270 of 328). We consider the classification error of 18% to be significant considering the complexity of the problem. Recalling the aims of this thesis, the proposed method of GAwGBE successfully provides a suitable solution such that it:

1. Uses Genetic Algorithm to establish a heuristically optimal subset of features from the full EEG feature set.
2. The subset both significantly reduces the number of features (from 738 to 52 features, 64 to 28 channels) *and* result in low classification error (18.6%)
3. The data sampling and processing methods simulate an online BCI system such that it is automated, computationally efficient and robust to slow sampling. This is done by significantly down sampling the data and implementing AAR prior to classification.

Despite satisfying the aims of this project, the solution could be improved. A classification error of 18.6% is far from ideal for a reliable BCI system. Indeed, better performance has been achieved previously using other methods on the same dataset.

## 5.3 Previous Studies: A Comparison

The results found in this and Alomari’s *et al*. study are summarised in Table 7.

Table 7 A comparison in classification accuracy between the best presented method used in this paper and the best result from Alomari et al. Both methods utilise an SVM classifier however the feature extraction/selection approaches are fundamentally different.

|  |  |  |
| --- | --- | --- |
|  | GAwGBE | Alomari’s *et al*. Feature Extraction |
| Classification Accuracy | 74.8% | 97% |

There is an obvious performance difference between the two techniques. There are many plausible justifications for this difference which can be subdivided into two areas: the approach and the signal processing.

First, we will consider variations in the approach. It may be the case that the features extracted using the method proposed by Alomari *et al*. are indeed superior features for motor-imagery classification. The features used include the power, mean and energy of the extracted ERS/ERD phenomena. As stated previously, there is evidence to suggest that these signals are associated with motor-imagery tasks [24]. Utilising knowledge regarding what specific channels and frequencies to examine, and how to process and extract relevant information for these signals may have led to the high performance of Alomari’s *et al*. solution. Compared to searching a wide space of generic feature combinations split across position, frequency and time delays, to locate a viable set as was the approach in this study, Alomari’s *et al*. approach appears more focussed and informed. However, there are other discrepancies between the approaches which have likely also contributed to the division.

Relative lacking performance in the proposed solution may also be attributed to the signal processing techniques used to prepare the data. Part of the aim of this study is to replicate online BCI sampling where the system is required to sample, filter, automatically detect and remove artefacts, extract features and classify events in multichannel EEG data fast enough to provide intuitive and responsive interfacing. To allow for all this processing, a sample rate of 2Hz is assumed as a ‘poor scenario’ simulation. A slow sampling frequency introduces problems for classification. Due to the frequency of the neural signal’s EEG is capturing, an event occurring at t=0 will appear quite different from one lagging by 0.25s and captured at the next sample t=0.5s. As a result, the classifier is required to learn a broad selection of potential feature patterns to associate with left or right labels. In comparison, Alomari *et al*. segments samples specifically surrounding events to establish consistent temporal features associated with the ERS/ERD events they attempt to extract, thereby reducing any time delay. Alomari’s *et al*. approach would be feasible given the BCI system is able to sample and process data at the rate the data was originally captured, i.e. 160Hz.

In addition to sampling discrepancies, the actual data selected for use from the PhysioNet dataset by each study is different. We chose to use a single recording of all 109 subjects for training and an additional recording of the first 20 subjects for testing. Alomari *et al*. use three recordings of six subjects in total, resulting in 18 two-minute recordings in comparison to the 129 used for this study. These sample sets are inconsistent and could influence the classification accuracy. In general, a larger number of samples provides a more generalised snapshot of the phenomena being classified. In this case however, generalising the classification problem could have made it harder for a classifier to learn the data. This is because there is evidence to suggest that features for neural event classification may be subject specific [7]. Hence it is possible that Alomari’s *et al*. classifier has learnt to classify events with more accuracy due to both the smaller number of subjects and the repeated samples taken from those same subjects reinforcing its training. Concerning filtering the data, both studies use a similar initial bandpass filter and notch for mains, although Alomari *et al*. state using a 50Hz notch rather than 60Hz despite the data being collected in the USA. Both studies used AAR, however Alomari’s *et al*. uses MATLABs EEGLAB software with an inbuilt AAR function, whereas this study trained a neural network to recognise and remove artefacts. Despite these different approaches, both methods are essentially employing ICA analysis therefore any variations in the data post-filtering is likely negligible. Whilst the proposed solution given by this thesis has not outperformed methods presented in previous studies, it has still produced a viable alternative to advanced feature extraction in this context.

## 5.4 Risk Mitigation Reflection

A risk analysis was completed prior to the commencement of this thesis. The following section will reiterate the three potential risks uncovered and report on the effectiveness of any mitigations employed. The risk level was assessed as per Table 8.

Table 8 The risk assessment matrix, mapping risk likelihood and severity to a risk factor of low, moderate, high or extreme in terms of its threat to the completion of this thesis.

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
|  | Severity | | | | |
| Likelihood |  | Negligible | Marginal | Critical | Catastrophic |
| Rare | Low | Low | Moderate | High |
| Unlikely | Low | Low | Moderate | Extreme |
| Possible | Low | Moderate | High | Extreme |
| Likely | Moderate | High | High | Extreme |
| Certain | High | High | Extreme | Extreme |

## Breach of Copyright

Copyright laws could be breached by including datasets and results from other studies in this project.

**Likelihood:** Rare

**Severity:** Marginal

**Risk level:** Low

**Original Mitigations:** All ideas and data used within this report which are not owned by the author will be referenced correctly in accordance with the University of Queensland referencing and plagiarism policy as well as per request by the original owner/s. In addition, datasets will only be collected from reputable sources.

**Success:** All documents and presentations pertaining to the completion of this thesis were indeed referenced as stated above. The contribution given by the thesis supervisor Assoc. Prof. Marcus Gallagher and the two PhD Candidates Brendan Keane and Nicholas Bland were acknowledge in the preliminaries of this paper. No breach of copyright occurred.

## Data Loss

Unforeseen circumstances such as damage to or loss of the authors computer, data or other records could lead to loss of project progress. As the project progresses, the severity of this risk increases to a critical level.

**Likelihood:** Unlikely

**Severity:** Critical

**Risk level:** Moderate

**Original Mitigations:** All data and electronic records will be committed regularly to a git repository dedicated to this thesis project. This includes but is not limited to: any MATLAB script, function files or data sets, electronic notes and assessment items. Care will be taken to minimise risk of malware being downloaded. Computers being used to work on the project will not be left unattended while thesis material is able to be accessed.

**Success:** All data and files were regularly committed to a git repository as suggested. No data of any kind was lost during the course of this thesis.

## Underestimation of Time Requirement

This project includes many steps which require careful implementation and validation before advancing to the next stage, as discussed in Section 5. Incorrect implementation or underestimation of the time required to complete anyone of these stages could lead to limitations in the project due to time constraints.

**Likelihood:** Possible

**Severity:** Critical

**Risk level:** High

**Original Mitigations:** A well-defined project plan has been outlined which includes milestones designed to motivate completion of the thesis within a manageable timeframe. This plan has been designed to expect delays and provide adequate time to overcome obstacles. In addition, regular correspondence with the thesis supervisor Assoc. Prof. Marcus Gallagher or one of his tutors/associates has been agreed upon to verify progress and provide guidance if required.

**Success:** The project plan was subject to changes as the aims of the thesis developed. Changes are outlined in the following section. Meetings with the thesis supervisor and/or tutors were irregular as little guidance was required. Where ever guidance or progress verification was required, the author made use of the dedicated Optimisation Thesis Project Piazza forum where correspondence was very timely.

## 5.5 Project Plan Reflection

The initial proposed project timeline for this thesis underwent significant changes over the course of the year due to adaptations in its aim and scope. These changes are summarised in Table 9.

Table 9 A month-to-month timeline comparison between the initial proposed thesis plan and the actual timeline

|  |  |  |
| --- | --- | --- |
|  | **Initial Proposal Plan** | **Actual Project Timeline** |
| **March** | Preparation of dataset | Preparation of dataset |
| **April** |
| **May** | Extract Features |
| **June** | Feature Selection using Particle Swarm Optimisation (PSO) and GA | Extract Features |
| **July** | Filter Method using GA |
| **August** | Wrapper Method using GA |
| **September** | Classification |
| **October** | Demonstration and Thesis preparation/Completion | Demonstration and Thesis preparation/Completion |
| **November** |

Key changes include:

* **Removal of PSO from scope** – this change occurred upon a suggestion for the project supervisor. This focussed the project on methods involving GA rather than broadening the scope to include a binary alteration of the PSO algorithm [25].
* **Filter to Wrapper method** – results attained through filter methods using correlation as a score metric were poor, and hence the project was adapted to focus into wrapper methods. Results and conclusions reached with the use of filter methods are still included as part of the scope of the thesis.
* **Classifiers** – initially it was proposed to use three classifiers to compare features selected using filter methods: ANN, SVM and random forest. However, due to the increased computation time requirement of implementing wrapper feature selectors, SVM was the only classifier tuned and implemented in the project. Of course, this classifier is not to be confused with the ANN implemented as part of the pre-processing AAR methods.

Overall the scope of the project was indeed scaled down from what was originally proposed. We believe that this focussing allowed for a more in-depth analysis of the GA method in this context, and provided the time required to produce significant results which may not have been possible with the broadness of the initial proposed plan.

## 5.6 Further Research

There are several avenues which could further build upon the findings presented in this thesis. Some of these are listed below:

* **Apply other optimisation algorithms** – GA was a suitable choice for this project as feature selection can be modelled as a binary string. However, many other heuristic search algorithms could be applied in this context. A thorough investigation into the performance of such algorithms could provide insight into which might be most suited to the scenario.
* **Apply this method in other contexts** – having shown GAwGBE to be an effective method of feature selection (with a focus on reduction) in this scope, there may be merit in applying this method to other such problems to understand how generally it can be effectively applied.
* **Incorporating feature reduction into GA operations** – as discussed in Section 5.2, we speculate that altering the crossover and mutation operations to encourage fewer features in subsequent generations could improve PGA. Possible alterations could include introducing some stochastic preference toward changing a 1 to a 0 over a 0 to a 1 in the next generation for either/both crossover and mutation operations.
* **Investigate other filter metrics** – this project considered correlation as a feature selector metric, however many others exist. Evaluating their effectiveness could result in a decent solution without the computation time of having to implement a wrapper.
* **Investigate and optimise performance of other classifiers** – as the project was focussed on optimisation methods for feature selection, the classification aspect was not thoroughly explored. The introduction of other classification methods could result in lower error if one is found to be better fit to the problem than the proposed SVM.
* **Reduce reliance on temporal data** – as discussed in *Results* Section 4.6, the feature set includes data which occurs up to 2.5 seconds after the initial MI event. This introduces a lag into the system, resulting in a slow BCI response. Eliminating the data in later samples from the feature set would reduce this lag but could result in increased classification error. An investigation into if the feature set can be reduced in this way, and if so by how much, could result in an improved BCI response time.
* **Including the ‘rest’ class** – this project models the problem as two-class class classification, left and right MI events. However, this model does not accurately simulate real-life BCI control as it does not consider the third, and likely most prevalent class, whereby the subject is neither imagining left- or right-hand movement but is in fact at ‘rest’. Without this class, the BCI will constantly classify between the two hands, which is not of much use for such applications as prosthesis control. Including the third class would enable the BCI system to be applied to real world scenarios.
* **Testing on live data** – implementing the BCI in a live scenario with a subject connected to an EEG sampling system would help to demonstrate if the system is indeed accurate, robust, responsive and intuitive and further outline any limitations.

Research into these areas could improve the capabilities of this technology, bringing reliable BCI control mechanism closer to reality.

# Conclusion

Despite the challenging nature of classifying MI events from EEG data, this thesis presents a viable solution for feature selection using GA within its specified aims and scope. The data was successfully filtered with an original AAR implementation before being passed to feature extraction. Features are segmented from the data across the 64 channels, into three frequency bands and over six temporal samples. Filter feature selectors using a correlation metric were found to produce poor solutions with both GFS and GA methods. Wrappers improved upon this performance, GA was found to outperform GBE in terms of classification error, which justified its use over the simpler greedy search algorithm in this context. However, due to the large number of features used in the GA solution, we developed two GA alterations to encourage the selection of fewer features: PGA and the hybrid GAwGBE. PGA did not perform particularly well despite variations in the penalty term *R*; we speculate this is because of the uninformed GA operations arbitrarily adding or removing features without knowledge of the direct influence this has on the error score. In contrast, GAwGBE provides a reasonable solution of 18.6% classification error with just 52 features (reduced from 738), resulting in a solution including just 28 of the original 64 channels. We state that this solution does satisfy the thesis aims such that it: uses GA to establish and heuristically optimal subset of features, this subset significantly reduces the number of features and has reasonably low classification error, and that the entire process could potentially be implemented as an online BCI systems in that it is automated, computationally efficient and robust. We compared the solution to another presented in a previous study by Alomari *et al*. [6] and discerned that while they’re solution yielded a very high classification accuracy (97% with SVM), variations between, and assumptions made by the two approaches were likely what led to this division. We assert that the risk management plan was followed over the course of the project with minimal variation and successfully led to a timely and hazard free thesis completion. Lastly, we state that the project plan was adapted at various points during the year to focus the scope around the implementation of GA as a wrapper in response to supervisor recommendations and experimental results.

BCI technology has the potential to provide an alternate means of control and communication across a wide range of applications. Innovations within this space may lead to significant improvements in the quality of life for a great many people. The results of this thesis, as with many related publications before it, are an indication that continued research into this field can bring this technology out of the realm of science-fiction and into reality.

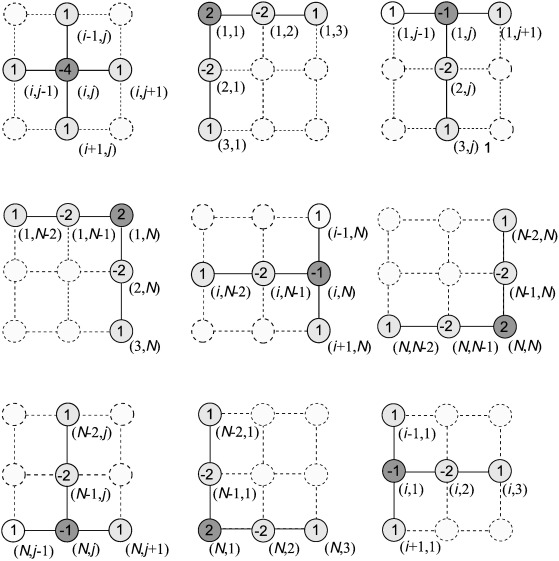
# Appendix

## A: Description of MATLAB Code

|  |  |
| --- | --- |
| Filename | Description |
| ANNFilter.m | Train and test the AAR ANN. |
| ArtefactRemoval.m | Apply preliminary filtering and AAR process to entire dataset. Uses ICAClassificationDataSetup.m |
| CORR.m | Return the correlation score of a given feature set using a precomputed correlation lookup table. |
| corrLookupGen.m | Generate the correlation lookup tables. |
| edfread.m | Reads .edf files. Source included in code. |
| featureExtraction.m | Loop through data extracting the features, uses getFeatures2.m |
| formLS.m | Constructs the surface Laplacian matrix. |
| GA\_Wrapper.m | Implements Genetic Algorithm. Despite the filename was used for both wrapper and filter methods. |
| generateData.m | Imports the data from the .edf files downloaded from the PhysioNet Database. Uses edfread.m |
| getAnnList.m | Convert the annotation list from a list of labels at time t to a list of samples with an labels. |
| getFeatures2.m | Returns a list of features for a subject. Obviously the second version. |
| goodSVMreduction.m | Performs GBE wrapper. |
| ICA.m | Performs FastICA. |
| ICAClassificationDataSetup.m | Performs preliminary actions to prepare data for AAR classification (filtering, feature extraction etc.) |
| ICATestScript.m | Used when manually labelling ICA components. |
| informedData.m | Reduces the feature set to the full 738 from a feature set including other frequency bands and channels. |
| newCORRbench.m | Performs GFS filter. |
| sampleRaw.m | Converts raw data into samples. |
| sortAnns.m | Sorts annotations into data which is usable by GA. Uses getAnnList.m |
| subArtefactRemoval.m | Reconstruct 64 channel data after artefacts are removed. |
| SVM.m | Train SVM and return classification error. |

## B: Surface Laplacian

The surface Laplacian decouples each channel from the reference electrode and instead provides relative measures between local neighbours. The diagram below explains that should the electrode of interest be the dark electrode, and have its neighbours (light electrodes) positioned as in one of the following arrangements, then the SL at that electrode is proportional to the sum of the voltage at each circle multiplied by the scalar in the circle. For instance, should our electrode have neighbours above, below, left and right of it (as is the case with most non-peripheral channels), then the SL of that electrode is given by:



## C: Pre and Post AAR Filtered EEG Data

This figure shows an example of the effects of AAR filtering on a single channel. The unfiltered data is shown in red, the filtered in blue. The filtered data is generally smaller in amplitude than the prefiltered, indicating that some artefactual component has been removed. Particularly large spikes (high frequency) have been significantly reduced. It is difficult to conclude what type, or types of artefacts are present as they are most recognisable as components compiled through the combination of multiple channels (through ICA). Despite this it is worth presenting this data as evidence that the AAR algorithm is indeed processing the data.

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|  |  |
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1. Tuning of SVM classifier is described in detail in *Methods* section 3.5.3. [↑](#footnote-ref-1)