

# Optimizing EMG Pattern Recognition for Prosthetic Hand Control: A Comparative Analysis of Windowing, Filtering, Features, and Classifiers Using the Ninapro Dataset

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Submitted: March 2026

## Abstract

Electromyography (EMG) enables intuitive prosthetic hand control through pattern recognition, yet optimizing signal quality, processing time, feature selection, and classifiers remains challenging. Using Ninapro DB1 (27 subjects, 52 movements), this research addresses these via four parts: general optimizations, combined analysis, individual analysis, and hybrid modeling. Key results: optimal window (500-250 ms, 69.12% accuracy, 15.76 s time), filtering (stimulus=restimulus, 75.31% accuracy), features (slope sign changes, wavelength, total power, time-frequency energy via statistical categorization/outliers/Friedman test,  $p=1.14 \times 10^{-7}$ ), classifiers (KNN/Random Forest at 90-97%). The hybrid model integrates these for 90.49% accuracy, offering a robust framework for real-time applications.

**Keywords:** Electromyography, window optimization, signal filtering, feature selection, machine learning classifiers, hybrid prosthetic model.

## 1 Introduction

The most effective approach for controlling an artificial prosthetic hand is to identify hand gestures or patterns using electromyography data. However, this technology has more potential than standard mechanically restricted movement (open-close), which frequently reduces naturalness and efficiency. Furthermore, as a real-world application for people, it is critical in human-computer interaction, virtual real-

ity, robotics, and upper/lower hand amputee assistive devices. The basic operation relies on skeletal muscle activation, which interferes with electrical information from EMG (electromyography) sensors. Recent research has shown that it is feasible to extract hand gesture patterns with greater accuracy from various muscle actions [1,2]. Using the publicly accessible NinaPro dataset, this study investigates solutions for problems with window size, filter condition, optimal features, and classifier selection [3, 4]. This Ninapro dataset comprises sEMG data from 27 healthy people doing 12 hand movement tasks and one rest position. The 1 to 8 channels contain data from eight sEMG electrodes evenly distributed around the forearm at the height of the radio humeral joint. Channels 9 and 10 provide signals from the flexor and extensor digitorum muscles. Stimulus is a pattern that repeats a movement, whereas restimulus repeats the same pattern but for a longer period. The terms repetition and rerepetition relate to the repeating of both the stimulus and the restimulus [3].

The objective is to investigate methods for enhancing classification of hand patterns using the Ninapro DB1 (Experiment 1) dataset. First, learn about the fundamental components of pattern classification systems. Then, examine each component in depth to see what difficulties are currently occurring. Finally, offer the most optimised solution for the specific challenges using conditional comparisons and statistical tests, and then suggest a better hybrid model. According to the extensive examination, the complete system includes data preparation, windowing, filter-

ing, feature extraction, and classification into appropriate class labels or hand patterns. Objectively, it is divided into four sections (A, B, C, and D), which include general findings, combined dataset analysis, individual subject analysis, and hybrid model proposals. For further examination, each section is broken into multiple experiments.

Part A: General Findings.

1. Determine the optimal window size.
2. Identify the Best Filtering Condition.

Part B: Combined Dataset Analysis (27 Subjects).

1. Find the best features using statistical analysis.
2. Analyse Outliers in Best Features.
3. Find the best classification model based on a variety of experiments.

Part C: Individual Subject Analysis.

1. Discover the best features for each subject.
2. Find the best classification model for each subject.
3. Perform a statistical analysis test on three different features.

Part D: Propose the Hybrid Model.

In the general results section (Part A), first do an experiment to establish the best window size based on the trade-off between accuracy and processing time for various classifier models. This experiment clearly demonstrates how window size influences model accuracy and processing time. Second, determine the optimal filtering conditions by computing the two parameters, mean squared error (MSE) and signal-to-noise ratio (SNR), under various stimulus, restimulus, repetition, and rerepetition circumstances. Calculate the accuracy of various classifier models with corresponding criteria at the same time.

This comparison will offer a clear direction for an optimal filter condition, utilising the trade-off between mean squared error, signal-to-noise ratio, and classifier accuracy. It utilises MSE and SNR measures

to optimise the trade-off between noise reduction and signal quality [5].

In Part B, I integrate 27 separate datasets into a single dataset and apply statistical analysis methods (mean, median, and standard deviation) to classify them as the best, moderate, and least informative features. Second, identify outliers in the best features; this helps to understand how data samples are organised in each feature. Finally, choose the optimal classifier model using several conditional tests [6, 7] (accuracy and processing time, mean accuracy and standard deviation, classification error, and confusion matrix).

In Part C, I first perform individual dataset analysis (experimenting on each of the 27 patients) to discover the most significant features in each. Second, identify the optimal classification model (KNN or Random Forest) for every subject. Finally, use the Friedman statistical non-parametric test to identify the most significant feature. This test ranks features based on their testing accuracy across multiple classifiers [8].

Part D proposes a hybrid model based on the optimisation findings from every part, as well as a feature combination-based hybrid model that incorporates features from many domains. The integration of time, frequency, and time-frequency domains strengthens machine learning models and improves their performance [9]. The MATLAB tool handled data preprocessing, feature extraction, and extraction time calculations, while the Python software handled hand pattern classification using multiple machine learning models and classification time calculations.

## 2 Theoretical Background and System Description

### 2.1 System Architecture

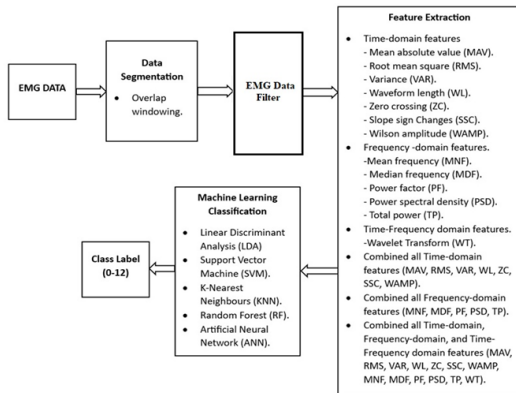


Figure 1: System architecture for pattern recognition using electromyography [10,11,12]

The system architecture processes real-time muscle activity signals to extract relevant features for prosthetic device control and pattern recognition. The data preprocessing stage extracts 10 channel EMG data from each subject, combining them in raw-wise order without changing class labels. The windowing process divides the extracted data into multiple segments, using the sliding window method to capture transitional information. The filter condition is applied to the current window, extracting features like time, frequency, and time-frequency domains. Machine learning classification models (LDA, SVM, KNN, Random Forest, and ANN) are trained to predict the accuracy of testing and training. The 13 target labels are used for predicting unseen data. Signal processing techniques like windowing, filtering, feature extraction, and pattern classification ensure the reliability and validity of surface electromyography (sEMG) data analysis, which is crucial for real-time prosthetic control, rehabilitation, and ergonomic studies.

### 2.2 EMG Data

The Ninapro dataset includes ten EMG channels in DB1, with the sEMG electrode detecting voltage changes in each channel to indicate muscle fibre contraction activity. These forms of muscle activity investigations aid in the analysis of muscle coordination, activation patterns, muscle fatigue, and motor control. These muscle signals offer the most significant contribution to pattern recognition. The flexor digitorum muscles flex the fingers, whereas the extensor digitorum muscles lengthen them. Figure 2 depicts the various EMG signals for ten channels.

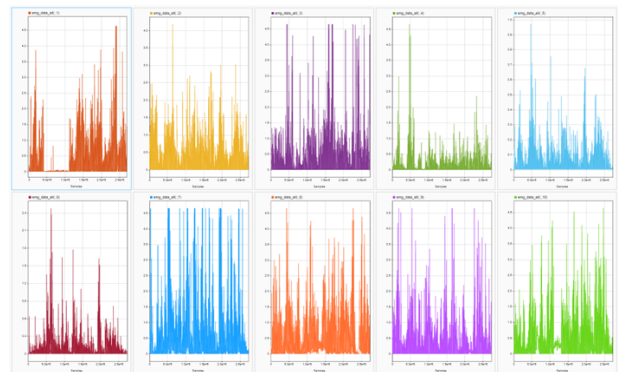


Figure 2: EMG signals for 10-channels.

#### 2.2.1 Stimulus, Repetition, Restimulus, and Rerepetition.

This study uses various techniques to analyze hand movements. Stimulus is used to ensure consistent hand movements and data collection. Repetition is the same visual prompt or action after initial exposure, examining its consistency and dependability over time. Repeating the same instructions ensures accuracy and reliability. Performance is assessed by assessing how successfully a person executes actions over time. Pattern identification aids in analyzing signal variations during the same action. Rerepetition ensures consistent results across multiple trials and monitors performance changes as the number of studies increases. Figure 3 illustrates these signal representations.

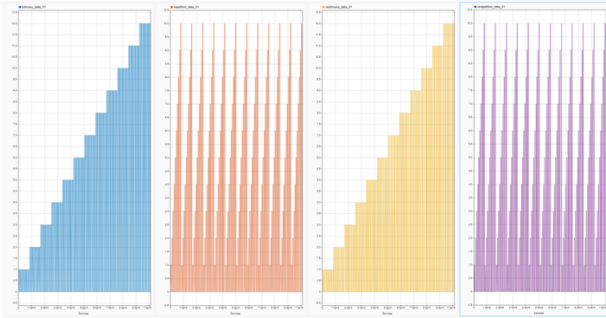


Figure 3: Signal representation of stimulus, repetition restimulus, and rerepetition.

There are noticeable points; the width of the restimulus and rerepetition signal are reduced as compared with their respective stimulus and repetition signals (see figure 4). The reason for reducing the signal width is due to multiple reasons, such as neural adaptation (the neural response of restimulus and rerepetition is reduced over time) and cognitive habituation (with repeated trials, the human cognitive system is more adept at processing restimulus and repetition actions). See figure 4. Graphical differences between stimulus and restimulus, as well as between repetition and rerepetition.

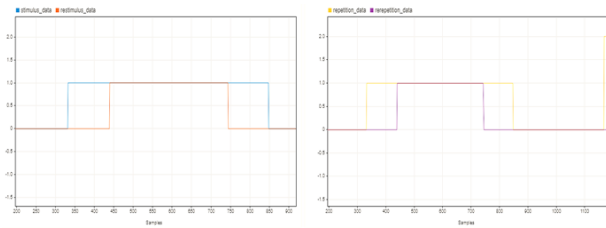


Figure 4: Graphical differences between stimulus and restimulus, as well as between repetition and rerepetition.

## 2.3 Windowing Process

The sliding window strategy is chosen in this project since the EMG data is a time series that varies constantly throughout time. The sliding window separates the EMG data into certain window lengths,

such as overlapping or non-overlapping segments (window). Each window's right features are retrieved and saved in a defined array. In this project, certain parameter names are utilised, such as winsize (actual window size), wininc (increase window size), numwin (number of windows), and curwin (current window).

The actual window sizes are 800, 500, 250, 150, 100, and 50, and increment window sizes are 400, 250, 75, 50, and 25. Figure 5 illustrate the respective window sizes, and their increment window sizes with retrieved samples. The window size specifies how many data samples are included. For example, if winsize = 500 is meant by each window, it will contain 500 samples from the signal or data. Window increment size (sliding window) is specified as how much window moves forward for the next segment. For example, if wininc = 250 is meant by each sliding or overlapping window, it will start 250 samples after the previous window's start. The number of windows (number) in each class (0 to 12) is calculated using the below equation (1). Where  $N$  is the total number of samples with the respective class. The '+1' ensures the initial window at the start of the signal; without including it, it misses the first segment window.

Finally, the representation of window size in seconds depends on the sampling rate of the EMG electrode; here, the Ninapro DB1 experiment is using 10 Otto Bock MyoBock 13E200 electrodes [13]. For example, consider sampling frequency is 1000 Hz and winsize is 500. Using equation (2), the answer is 0.5 second or 500 milliseconds.

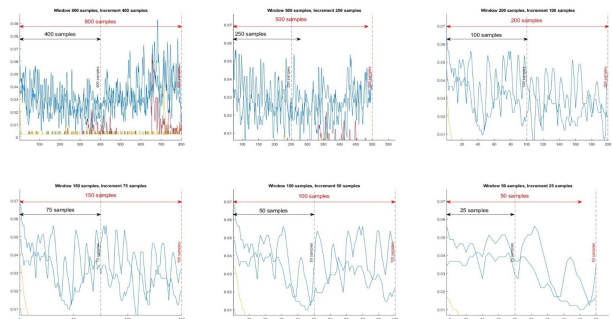


Figure 5: Window size details.

$$numwin = \left\lfloor \frac{N - winsize}{wininc} \right\rfloor + 1 \quad (1)$$

Equation (1). Calculation for number of windows

$$time = \frac{winsize}{sampling\ frequency} \quad (2)$$

Equation (2). Calculation for window size time in seconds.

## 2.4 Filtering

EMG data filtering relies on stimulus, restimulus, repetition, and rerepetition values, but their arrangement can be inconsistent. For instance, a combination of stimulus = 0 and restimulus = 3 indicates preparation state, causing annotation delays. The system is labeled rest state, but an upcoming action or pattern is already marked in the restimulus. To ensure consistency, accurate labeling, and noise reduction, the filtering should be chosen where both the stimulus and restimulus values are equal. This method is also applicable to repetition and rerepetition.

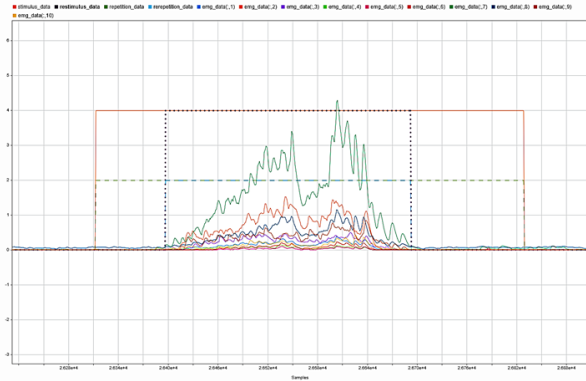


Figure 6: EMG signal extraction when stimulus and restimulus are equal.

## 2.5 Feature extraction

Various features are extracted from time, frequency, and time-frequency domains after the data filtering.

Seven time-domain features are extracted, such as mean absolute value, to provide the average of the absolute values of the EMG signal [14]. Root mean square provides an indication of the energy content and power of the EMG signal [15]. Variances indicate the power of the EMG signal [16]. Waveform length provides the cumulative length of the EMG waveform over time [14]. Slope Sign Changes The number of times the slope of the EMG signal changes sign, indicating the frequency of muscle activity changes [14]. Zero crossing is the number of times the EMG signal crosses the zero line, indicating the frequency of the signal's oscillations [14]. Willison Amplitude is the count of the number of times the amplitude difference between consecutive samples exceeds a threshold, related to muscle contraction levels [17].

In the frequency-domain feature, mean frequency (MNF) is the average frequency of the signal, providing muscle fatigue and the frequency content of the signal [18]. Median Frequency (MDF), the frequency that divides the power spectrum into two equal parts, is often used to assess muscle fatigue [18]. Peak frequency (PF), the frequency at which the power spectrum of the EMG signal is at its maximum [18]. Power spectral density (PSD), a measure of the power distribution of the signal across different frequencies, is used for analysing frequency components [19]. Total Power (TP) is the total energy of the EMG signal over a period, representing the overall power of the signal [20, 23]. Time-frequency energy (TF-Energy) features from the time-frequency domain and provide the total energy of the EMG signal over a period, representing the overall power of the signal [21, 22]. Finally, combine seven time-domain features (COMB-TD), five frequency-domain features (COMB-FD), seven time-domain and five frequency-domain features (COMB-TD & FD), and combine all time, frequency, and time-frequency domain features (COMB-TD & TFD).

## 2.6 Classification Model and Class labels

After feature extraction from various domains, these data fed to the different machine learning classification models (LDA, SVM, KNN, RF, and ANN). The

target labels of all these classification models range from 0 to 12 (as illustrated in Figure 7), and each class label is represented by a distinct hand pattern.

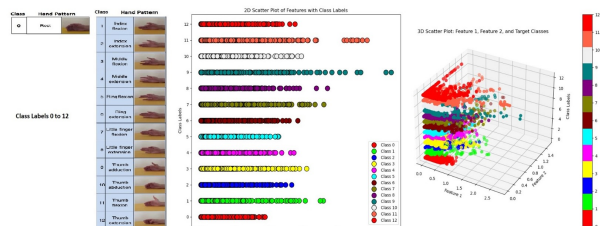


Figure 7: Classification label [3]

### 3 Literature Review

In this review of the literature, I would like to address the key findings about electromyography systems, as well as the challenges brought up by several research investigations and suggestions for advancement. The part of overall processing of EMG signals includes, a more sophisticated method of windowing, advanced filtering techniques, methods for feature extraction, and combining of multi-model data. Finally, discuss the significant machine learning model that has shown superior performance in Ninapro datasets.

Preprocessing is the first step of pattern recognition techniques for reducing the interferences of the signals, such as signal acquisition noise, electromagnetic interference, and instability of the signal due to the wrong placement of electrodes [24]. To address these filtering issues, the top filtering models in EMG signal processing are wavelet transform filtering, Kalman filtering, and Notch filtering. Wavelet Transform Filtering (WTF) is one of the best methods to avoid the EMG noise reduction. In D. Farina and R. Merletti et al. [25], address the various noise factors are affected by sEMG electrodes such as movement artifacts, electromagnetic interference, crosstalk, internal noise, and electrocardiographic (ECG) artifacts are discussed. The solution to improve the system performance mainly depend on their hardware components and algorithms. For improving design architecture, size, higher signal-to-

noise ratio these parameters sufficiently reduce the noise contents [26,27,28]. These techniques are mitigating the impact of inherent noise. In the case of movement artefacts to distort the EMG signals, it's mainly due to the displacement of electrodes and cables [29]. The solution to the concern is to use proper conductive gel to help maintain a stable interface, use wavelet transform filter techniques to reduce the motion artefact noises, and use proper electrode placement [30, 31]. Electromagnetic noise due to external devices; the solution to this problem is using adaptive filtering techniques such as FIR and IIR notch filters. These filters are effectively removing the electromagnetic interference noise [32]. Laguerre filters are the reliable technique for removing the power-line interference (PLI) [33]. The common in signal interpretation due to the size of the electrode and placement of muscle region [35]. To use of high-pass spatial filters is much capable to avoid the subcutaneous tissue thickness, muscle fatigue, and electrocardiographic artefacts issues [36,37,38]. The proposed solutions effectively address the various noise issues in EMG signals and lead to more accurate and reliable EMG signal interpretation in clinical diagnostics, rehabilitation devices, and human-machine interfaces.

The most common method used in electromyography field or time-series data is the sliding window method, and it's powerful for analysing the dynamic muscle activity. The key factors in windowing are the selection of the actual window and sliding window sizes. Window size is a trade-off between many factors in electromyography systems, especially accuracy and processing time. A larger window size can capture the general pattern of the signal but might lead to underfitting by missing short-term significant events. Smaller window sizes are providing higher temporal resolution and able to capture the small changes, so it probably leads to overfitting [39, 40]. The characteristics of the EMG signal are dependent on the muscle activity and their movement. Therefore, the fixed windowing method is not applicable for all types of movements. The solution to the issue is to use the adaptive windowing technology so it automatically changes the window size as per the characteristics of the EMG signals [41]. However, the cross-validation method is a considerable option to

analyse the consistency of the window process [42]. One of the real-time limitations is that smaller window sizes capture a high amount of data, and the cost of computation is high. The suitable solution of the computational overhead to implement the dimensionality reduction techniques such as linear discriminant analysis (LDA) without loss of information [43]. The best trade-off between actual window size and incremental window size is fifty percent, and it balances the computation time and quality of the signals [44].

The time-domain, frequency-domain, and time-frequency domain features are commonly extracted from the Ninapro dataset. The time-domain features contain the characteristics of statistical and waveform-based information of the EMG signal. In the frequency domain, using Fourier transform, the features are derived by transforming the signal into the frequency domain. The time-frequency domain features are a combination of both and provide more detailed information about the non-stationary EMG signals. In-time-domain features are widely used due to their simplicity and low computational power. In the context of the latest research on pattern recognition systems, deep learning methods are used for getting higher accuracy and robustness. However, convolutional neural networks (CNNs) and recurrent neural networks (RNNs) provide higher accuracy as compared to traditional features [45, 46]. The main challenges are noticed variability of EMG signals, data collection, and normalization. The variability of electromyography (EMG) signals is measured by the accuracy of the model being changed due to the changes of individuals, which affect the consistency. Secondly, for ensuring the normalisation technique is consistent for feature extraction with an extensive amount of data. The solution to these challenges is to standardise the EMG signals across different users and sessions, such as z-score normalisation or min-max scaling to reduce variability. In the context of an EMG signal, the wavelet transform (WT) feature is one of the powerful features for analysing non-stationary EMG signals. It contains both time and frequency domain information, and this combined information nature will provide higher accuracy in particular machine learning models [47]. The main challenges are the complexity of the feature extraction

and ensuring that feature is computationally efficient. Finally, the entropy-based feature extraction method plays critical roles if the electromyography signals are of an irregular nature. Phinyomark et al. introduced how the integration of entropy-based both time and frequency domain features significantly increases the accuracy and stability of pattern recognition systems [48]. The main challenges are to find the best features and avoid the redundant or irrelevant features because it reduces the system performance. The solution for this issue is to implement the feature selection method such as recursive feature elimination and LASSO (Least Absolute Shrinkage and Selection Operator). Moreover, the hybrid method or integration of features from different domains provides a comprehensive solution to the above issues [49].

LDA is a linear classifier, and it works better in a linear dataset, and that best separates the classes (hand pattern) in the dataset. If the data is a non-linear combination, LDA is less efficient in terms of accuracy prediction and least computation time with a low memory requirement [1]. As discussed in various research papers, the solution to solving the non-linear data is to use higher complex algorithms such as support vector machines (SVMs) and neural networks (CNNs and RNNs) that can handle the non-linear relationships in the data [50, 51]. The use of kernel functions for the SVM algorithm is more efficient in non-linear complex data due to the hyper-plane mechanism [52]. The most common challenges for the data preprocessing part are features selection, nonlinearity in data, training time complexity, and class imbalance. The solution for this issue to choose relevant features with help of feature engineering method and reducing the dimension of the data. Due to these techniques to avoid the overfitting and improve the computational efficiency. However, the method of regularisation and resampling techniques is better to reduce the complexity of the data, training time, and provide better class imbalance issues.

k-Nearest Neighbours (k-NN) is more efficient in the case where the data distribution is unknown or of a nonlinear nature. Due to the simplicity, the computation time is comparatively low and performs well in moderate-sized datasets but might be overfitting [53]. Overall, as per the research paper analysis, the com-

mon challenges are computational time, scalability, distance metric choice, and choice of k value. The issues of computational complexity and scalability can solve nearest neighbours (ANN) algorithms (e.g., KD trees, ball trees). For improving scalability, the clustering method is highly efficient. Random Forest is a method of ensemble learning, and it includes multiple decision trees during training and provides the result based on the mode of individual trees. As per the research paper analysis, it outperformed in complex data and indicates the accuracy range of 85 to 95 percent. The noticeable point is that the number of trees is higher, and the computation cost is also high. These issues can be solved by using a threshold value that is determined due to the experiment between the quantity of characteristics and the accuracy of the model [54].

The artificial neural network (ANN) model is more powerful as compared with the above classifiers, but the common factor is the higher computation time. As a result, the complex algorithms are not suitable for real-time applications due to the key factor processing time. The highlighted characteristic is automatically learning the data pattern with multiple epochs and providing better classification accuracy. One of the drawbacks for ANNs is that it requires a large amount of data, so obviously it led to higher computation power. The challenges are overfitting, high dimensionality, and hyperparameter tuning. The implied solutions for finding the overfit are to experiment with the performance accuracy in training and testing datasets. It helps to mitigate the overfitting by validating the model's performance on unseen data. The high-dimensionality issue was solved by applying a diffusion map for dimensionality reduction in fuzzy classification. Finally, the hyperparameter tuning issue was solved by experimenting with different hidden layers and activation functions to find the optimal configuration [55].

## 4 Simulation Details

This project simulation is divided into two parts: the first (Part 1) uses MATLAB for feature extraction and extraction time calculation, and the second is

a machine learning classification model created in Python using a Jupyter Notebook. The reason for separating is that the MATLAB tool offers superior capabilities for mathematical calculation in the case of feature extraction, however, Python provides a diverse set of machine learning libraries to do numerous tasks in comparison to the MATLAB tool. For a better method, utilise multiple tools depending on their performance in different contexts, and then combine both files to use batch files (Windows) or shell scripts (Linux) via a sequence of instructions run by the command prompt. It enables you to automate many processes by executing several commands in order.

The simulation operates in two separate scenarios. The first step is to integrate the 27 individuals from the Ninapro DB1 dataset into a single data set. Second, work with individual participants from the Ninapro DB1 dataset. This type of experiment was used to study and compare the assessment, processing time, and feature selection in both domains. However, both techniques might serve to create a more sophisticated hybrid model. The final hybrid model is trained using all the participants' data; thus, it has various properties and a wider prediction range than individual trained models. Figure 8 shows the flowchart for the total simulation process.

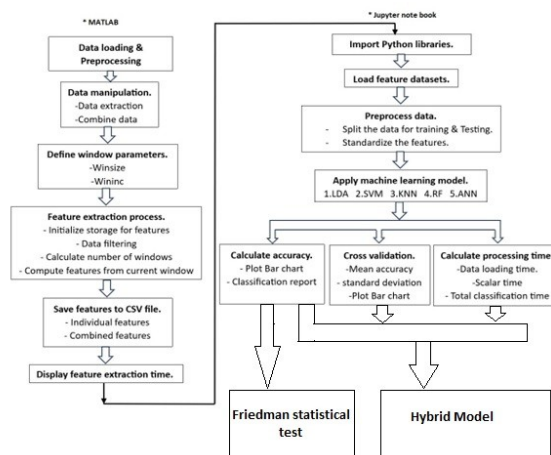


Figure 8: Flowchart for overall simulation.

## 4.1 Part:1 (MATLAB)

### 4.1.1 Data Loading and Preprocessing

The data loading and preprocessing phase begins by recording the overall execution time using the command `startTime = datetime('now');`. The `clc` command is used to clear the MATLAB command window. Using the `load` function, data for 27 participants are loaded and stored in separate variables (e.g., `data_01`, `data_02`, etc.). After loading the datasets, the script prints the dimensions of various data matrices, including EMG data, stimulus, restimulus, repetition, and rerepetition. This structured representation ensures that the data is correctly loaded and provides a quick diagnostic overview. See Figure 9 for the data loading and preprocessing steps.

```
Subject 01:
EMG data size: [101014 10]
Stimulus data size: [101014 1]
Restimulus data size: [101014 1]
Repetition data size: [101014 1]
Rerepetition data size: [101014 1]
Subject 02:
EMG data size: [100686 10]
Stimulus data size: [100686 1]
Restimulus data size: [100686 1]
Repetition data size: [100686 1]
Rerepetition data size: [100686 1]
```

Figure 9: Data loading and preprocessing.

### 4.1.2 Data Manipulation

The data manipulation section first performs specific variable extraction (EMG, stimulus, restimulus, repetition, and rerepetition) from the loaded data for each participant using the `eval` function. This function evaluates the string for the 27 participants' data and assigns respective variable names; for example, `data_01` is mapped to `emg_data_01`, `stimulus_data_01`, and so on. The `sprintf` command is used to dynamically create these variable names for all subjects. Secondly, individual variables from all participants are combined into a single dataset using concatenation (`[]`), resulting in variables such as `emg_data_all` and `stimulus_data_all`.

Finally, the script displays the sizes of both the individual and combined datasets.

```
Combined data size of 1 to 27 Participants: [2731393 14]
EMG data_all size: [2731393 10]
Stimulus_data_all size: [2731393 1]
Restimulus_data_all size: [2731393 1]
Repetition_data_all size: [2731393 1]
Rerepetition_data_all size: [2731393 1]
```

Figure 10: Data Manipulation process.

### 4.1.3 Define window parameters.

Windowing is a crucial step in electromyography (EMG) for analyzing non-stationary signals. It divides the continuous signal into small segments and analyzes them independently, allowing for the extraction of features over time. Window size (`winsize`) refers to the number of data points within each segment, while window increment (`wininc`) is the number of samples moved forward for the next segment. This overlapping method ensures no significant information is lost between segments and improves feature extraction smoothness. The selection of `winsize` and `wininc` is a trade-off between temporal resolution and computational efficiency. See figure 11: Window parameter initialisation.

```
% Define window size and increment
winsize = 500; % Window size in samples (800, 500, 200, 150, 100, 50)
wininc = 250; % Window increment in samples (400, 250, 100, 75, 50, 25)
```

Figure 11: Window parameter Initialisation.

### 4.1.4 Feature extraction process

1.storage for features

Initialise the array storage for each feature across all classes. These include time-domain, frequency-domain, and time-frequency domain. Similarly, initialise the timing variables to record the time taken for extracting each feature type across all classes. See figure 12 for feature storage initialisation.

```

% Initialize storage for each feature across all classes
all_mav_features = [];
all_rms_features = [];
all_var_features = [];
all_wl_features = [];
all_zc_features = [];
all_ssc_features = [];
all_wamp_features = [];

all_mnf_features = [];
all_mdf_features = [];
all_pf_features = [];
all_psd_features = [];
all_tp_features = [];

all_tf_energy_features = [];

```

Figure 12: feature storage initialisations

2.Data filtering and display number of windows in each class

The EMG data is filtered based on the condition that both the stimulus and restimulus values match the current class label, and the value of repetition matches the value of rerepetition. Finally, this process loops through different classes, filters the data by stimulus and repetition labels, calculates the number of windows in each calss, and allocates storage for various features. See Figure 13 for data filtering and number of windows.

```

Number of windows for class 0: 4020
Number of windows for class 1: 387
Number of windows for class 2: 388
Number of windows for class 3: 410
Number of windows for class 4: 371
Number of windows for class 5: 371
Number of windows for class 6: 391
Number of windows for class 7: 384
Number of windows for class 8: 398
Number of windows for class 9: 365
Number of windows for class 10: 343
Number of windows for class 11: 342
Number of windows for class 12: 380

```

Figure 13: data filtering and number of windows.

3.Compute features from current window and extraction time calculation.

The feature extraction and their computing time are recorded for each window to evaluate the efficiency and performance of the process. The extracted features have different aspects of the signal, including amplitude, frequency content, and energy distribution. These features are stored in pre-allocated matrices for each feature type. After extracting these features for each class, append the class label to the last column for each feature matrix. See figure 14 for feature extraction and their computation time.

```

Total processing time for MAV features across all classes: 1.18 seconds
Total processing time for RMS features across all classes: 0.59 seconds
Total processing time for VAR features across all classes: 0.90 seconds
Total processing time for WL features across all classes: 0.61 seconds
Total processing time for ZC features across all classes: 1.30 seconds
Total processing time for SSC features across all classes: 1.26 seconds
Total processing time for WAMP features across all classes: 0.86 seconds
Total processing time for MNF features across all classes: 6.34 seconds
Total processing time for MDF features across all classes: 4.61 seconds
Total processing time for PF features across all classes: 0.54 seconds
Total processing time for PSD features across all classes: 3.83 seconds
Total processing time for TP features across all classes: 0.13 seconds
Total processing time for TF Energy features across all classes: 74.36 seconds

```

Figure 14: feature extraction and their computation time

#### 4.1.5 Save features to csv file.

The process is storing features from EMG data to CSV files, then integrating time-domain, frequency-domain, and time-frequency domain features into a single dataset with respective names (for example, combined time and frequency domain feature). Each channel has a dynamic header name, which includes (channel names and target label) Feature 1, Feature 2, and class. Figure 15 illustrates feature saving as csv files.

```

All feature sets saved successfully as CSV files.
Combined feature sets saved successfully as CSV files.
Total execution time: 00:01:57

```

Figure 15: features saving as csv files.

## 4.2 Part: 2 (Python-Jupyter Notebook)

### 4.2.1 Development for classification model.

Part 2 mainly focuses on machine learning classification-based models, evaluating metrics such as accuracy, mean accuracy, standard deviation, and classification time. The primary libraries utilized are scikit-learn, NumPy, Pandas, Matplotlib, and Seaborn.

Scikit-learn is a widely used machine learning library for data preprocessing, model training, and evaluation. It includes classes such as `sklearn.discriminant_analysis.LinearDiscriminantAnalysis` (LDA), `sklearn.svm.SVC` (SVM Classification), `sklearn.neighbors.KNeighborsClassifier` (KNN Classification), `sklearn.ensemble.RandomForestClassifier` (Random Forest Classification), and `sklearn.neural_network.MLPClassifier` (ANN Classification). Additionally, the NumPy library is used across all models for numerical operations and array handling, while the Pandas library is employed for data manipulation and preparation. For data visualization and plotting, Matplotlib and Seaborn are used. In the case of accuracy, calculation is evaluated by various machine learning models such as linear discriminant analysis (LDA), support vector machines (SVM), K-nearest neighbours (KNN), random forests (RF), and artificial neural networks (ANN). Each classifier follows a consistent workflow: first load the extracted feature dataset and split it into features (channels or columns in a dataset) and target labels (class labels). Then, after data split into training and testing sets, features were scaled using “StandardScaler” to standardise the input. Each classifier is trained on scaled training data, and predictions are made on the test set. The performance of the model is based on their testing and training accuracies. The results are visualised through bar plots with the help of the “matplotlib” library. Additionally, a detailed classification report, including precision, recall, and F1-score, is generated for better comparison between the classifier models. See figure 16 results for accuracy and classification

report.

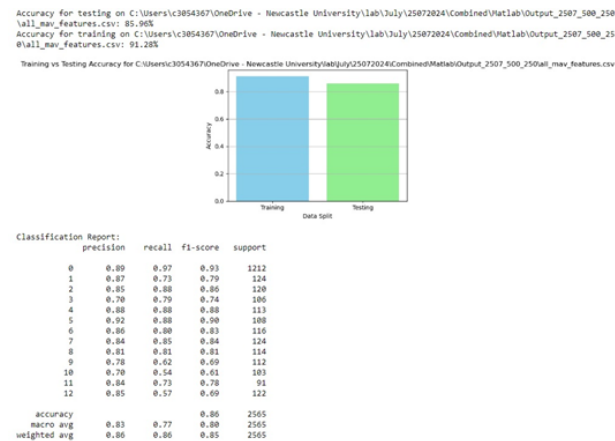


Figure 16: results for accuracy and classification report.

Secondly, for calculating mean accuracy and standard deviation for each model with the help of cross-validation techniques. This technique ensures the stability or robust assessment of model performance. For each classifier (LDA, SVM, KNN, RF, and ANN), the dataset is divided into multiple folds, and the model is trained and tested on different subsets of the data. Calculate the accuracy scores in each fold and compute the mean accuracy; it indicates the average performance of the model across different subsets of the data. The standard deviation is also calculated to assess the variability or consistency of the model’s performance. The library “random.state” ensures that the initialisation of the model parameters is consistent across multiple runs, which helps in stabilising results. Overall, this approach provides information on how each model performs on average and how stable the performance is across different data splits. See figure 17 result for mean accuracy and standard deviation.

```

Results for MAV_Features:
Mean Test Accuracy: 60.36%
Standard Deviation of Test Accuracy: 1.49%

Results for RMS_Features:
Mean Test Accuracy: 61.19%
Standard Deviation of Test Accuracy: 1.82%

Results for VAR_Features:
Mean Test Accuracy: 60.75%
Standard Deviation of Test Accuracy: 1.44%

Results for WL_Features:
Mean Test Accuracy: 59.36%
Standard Deviation of Test Accuracy: 1.73%

```

Figure 17: Result for mean accuracy and standard deviation.

For calculating the classification time for each feature set with respective models based on required time for both training and prediction phases. This is achieved by calculating the start and end times of the classification processes for each feature set and model combination. This result provides a clear understanding of the computational efficiency of each feature with respective models and valuable insights into their practical performance and suitability for real-time applications. However, performing statistical analysis on each feature using mean, median, and standard deviation. It's helping to understand data distribution. Standard deviation indicates the amount of variation or dispersion from the mean. Secondly, calculate the outliers of those features. The outlier indicates the sum of data distribution, central tendency, variability, and presence of noise. See figure 18. for statistical analysis tests.

```

Summary statistics for C:\Users\c3854367\OneDrive - Newcastle University\lab\July\25872824\Combined\Matlab\Output_2587_500_258
\all_ssc_features.csv:

```

	mean	median	std	range
Feature1	175.376659	177.0	54.655574	385.0
Feature2	138.166667	157.0	86.713669	393.0
Feature3	148.563275	157.0	68.550909	368.0
Feature4	117.756691	169.0	105.348250	402.0
Feature5	134.362222	113.0	89.456761	376.0
Feature6	147.623263	167.0	103.620858	387.0
Feature7	174.607251	179.0	56.614676	388.0
Feature8	191.514269	184.0	38.743244	346.0
Feature9	132.773673	144.0	95.277312	399.0
Feature10	177.297895	184.0	62.752823	354.0

Figure 18: Statistical analysis test

Finally, calculate the confusion matrix for the best model and provide a detailed breakdown of the model's performance by showing the number of true positives, true negatives, false positives, and false

negatives. Additionally, in the individual data set, perform the accuracy comparison and find the best features and subjects out of 27. At last, perform the Friedman non-parametric test in the best features (wavelength, total power and time-frequency energy) of individual subjects. See figure 19. for Friedman test statistic and P-value

```

Processing wl_features for S27 from file: C:\Users\c3854367\OneDrive - Newcastle University\lab\Individual\Output_files\S27_wl_
features.csv
Accuracy for testing on C:\Users\c3854367\OneDrive - Newcastle University\lab\Individual\Output_files\S27_wl_features.csv: 81.7
1%
Processing tp_features for S27 from file: C:\Users\c3854367\OneDrive - Newcastle University\lab\Individual\Output_files\S27_tp_
features.csv
Accuracy for testing on C:\Users\c3854367\OneDrive - Newcastle University\lab\Individual\Output_files\S27_tp_features.csv: 87.4
0%
Processing tf_energy_features for S27 from file: C:\Users\c3854367\OneDrive - Newcastle University\lab\Individual\Output_files
\S27_tf_energy_features.csv
Accuracy for testing on C:\Users\c3854367\OneDrive - Newcastle University\lab\Individual\Output_files\S27_tf_energy_features.cs
v: 83.33%
Friedman test statistic: 31.988582524271846, p-value: 1.1361387919667559e-07
There is a significant difference between the feature types.
Results saved to C:\Users\c3854367\OneDrive - Newcastle University\lab\Individual\Output_csv_file\combined_results.csv

```

Figure 19: Friedman test statistic and P-value.

## 5 Results and Discussion

### 5.1 Part A. General Findings

#### 5.1.1 Determine the optimal window size.

Finally, considering trade-off aspects, the window size 500-250 (winsize = 500, wininc = 250) is chosen over 800-400 (winsize = 800, wininc = 400). Table 01 compares the average accuracy and processing time for various windows. Figure 20 depicts a graphic representation of window size, accuracy, and processing time. When the window size is raised, the accuracy of all models improves, and processing times drop systematically. Shorter window sizes need more time than larger ones. The requirements include large window sizes (800-400) capable of catching long-term trends and overall signal patterns, but they may result in underfitting by missing short-term important events or temporal resolution of the signal. Smaller window sizes, on the other hand, provide increased temporal resolution while still recording rapid signal changes. However, the resulting obtained features more sensitive to noise and volatility over time. Choosing the appropriate window size involves balancing the model's accuracy against its processing time. Finally, I chose the window size 500-250, which had an average total accuracy of 69.12 percent and an average total processing time of 15.76 seconds.

It delivers reasonably greater precision while reducing processing power and short-term fluctuations or noise. The study’s goal is to strike a compromise between window size and system processing time to improve classification accuracy while minimising latency. Iterating over subjects and extracting sequential features are among the challenges. For better performance, a parallel windowing solution is advised.

Table 1: Comparison between average accuracy and processing time for different windows

Window size	Parameter	LDA	SVM	KNN	RF	ANN	Avg. Total
800-400	AVG. Accuracy	60.51	63.19	73.69	77.90	72.59	69.58
	AVG. Processing time	12.28	12.72	12.32	12.94	28.54	15.76
500-250	AVG. Accuracy	57.80	60.88	75.50	78.46	72.94	69.12
	AVG. Processing time	14.72	16.99	14.85	18.26	36.49	20.26
200-100	AVG. Accuracy	56.52	58.48	71.94	75.38	70.00	66.46
	AVG. Processing time	18.20	35.17	18.64	27.83	71.15	34.20
150-75	AVG. Accuracy	55.10	56.69	67.14	73.39	68.04	64.07
	AVG. Processing time	20.30	53.60	20.59	33.02	77.39	40.98
100-50	AVG. Accuracy	53.90	55.08	68.32	72.25	66.18	63.15
	AVG. Processing time	20.81	109.55	21.53	169.36	113.05	86.86
50-25	AVG. Accuracy	52.77	53.64	66.61	71.24	64.30	61.71
	AVG. Processing time	31.93	589.42	34.42	71.93	188.09	183.16

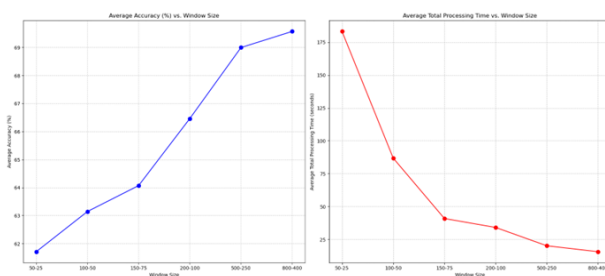


Figure 20: Graphical representation of window size and accuracy, and window size and processing time

### 5.1.2 Identify the best filtering condition.

Model 4 condition improves SNR moderately after filtering (1.027378 to 1.267982) and has less impact on MSE compared to models 1 and 2 condition. The moderate improvement in SNR after filtering indicates better quality of features used for classification and maintains a relatively low MSE, suggesting good accuracy in predictions.

In terms of accuracy, the filtering condition of model 1 performs very poorly (average accuracy =

62.48%) compared with other models (2, 3, and 4). In Model 2 and Model 3, there is no significant difference in testing, training, or F1score. Random Forest and KNN generally outperform the other classifiers in terms of accuracy and F1 scores across all models. Finally, model 4 shows the strongest results overall in terms of testing (75.31%), training (79.85%), and F1-score (0.73). Table 02 and Figure 21 provide a clear comparison incorporating accuracy, F1-score, mean squared error (MSE), and signal-to-noise ratio (SNR) for each model condition.

Table 2: Comparison for different filter conditions

Filter Conditions	Average MSE (Original)	Average MSE (Filtered)	Average SNR (Original)	Average SNR (Filtered)	Classifiers	Testing Accuracy	Training Accuracy	Weighted Avg. F1-Score
Model 1 (Stimulus filtered by Repetition condition 1 & 2)	0.123535	0.200242	1.027378 dB	1.413006 dB	LDA	35.08%	36.54%	0.340
					KNN	85.25%	90.32%	0.850
					SVM	40.25%	40.89%	0.390
					RF	89.33%	100.00%	0.890
					<b>Average</b>	<b>62.48%</b>	<b>66.94%</b>	<b>0.618</b>
Model 2 (Stimulus filtered by Repetition condition 5,6,7,8,9,10)	0.123535	0.194971	1.027378 dB	1.361528 dB	LDA	53.44%	53.61%	0.480
					KNN	86.68%	92.13%	0.860
					SVM	56.13%	56.72%	0.510
					RF	91.11%	100.00%	0.910
					<b>Average</b>	<b>71.84%</b>	<b>75.62%</b>	<b>0.690</b>
Model 3 (Stimulus filtered by all Repetition condition (1 to 10 and including 0))	0.123535	0.123535	1.027378 dB	1.027378 dB	LDA	53.44%	53.61%	0.480
					KNN	86.68%	92.13%	0.870
					SVM	56.13%	56.72%	0.510
					RF	91.11%	100.00%	0.910
					<b>Average</b>	<b>71.84%</b>	<b>75.62%</b>	<b>0.693</b>
Model 4 (Stimulus = Restimulus filtered by Repetition value = Rerepetition)	0.123535	0.168031	1.027378 dB	1.267682 dB	LDA	60.70%	62.07%	0.560
					KNN	88.15%	92.25%	0.880
					SVM	62.96%	65.06%	0.590
					RF	89.43%	100.00%	0.890
					<b>Average</b>	<b>75.31%</b>	<b>79.85%</b>	<b>0.730</b>

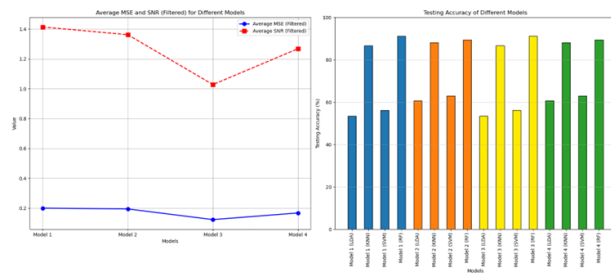


Figure 21: Graphical representation of different filter condition.

## 5.2 Part B. Combined Dataset Analysis (27 Subjects).

### 5.2.1 Find the best features using statistical analysis.

The root mean square feature (RMS feature) has slightly higher mean and median values (0.20 and 0.10) compared to MAV features (0.17 and 0.08). It indicated RMS features more informative data than MAV features, but higher variability could be challenging. Wavelength (WL Feature) features have high mean (4.71) and median (2.93) values along with a significant standard deviation (5.35), so this feature can capture the complex patterns and distinctions between classes. Slope Sign Change feature (SSC feature) and finally, in time domain sessions, slope sign change, wavelength, and root mean square features provide better accuracy based on their ranges of values, such as mean, median, and standard deviation.

In the frequency domain, the best feature is total power (TP feature) due to the higher mean (.83), median (0.46), and standard deviation (1.07) compared with frequency domain features. Relatively small values mean median and standard deviation indicate that feature capture subtle frequency characteristics. In the time-frequency domain, the time-frequency energy features (TF Energy Features) provide extremely high values for both mean (389.57) and standard deviation (485.15). It indicates that this feature can capture more information but may also be very noisy. Evaluation based mean, median, and standard deviation, each feature is categorised into best, moderate, and less informative features. See table 04 for categorised statistical features.

Table 3: Statistical comparison of features

Dataset	Mean (Average of 10 channels)	Median (Average of 10 channels)	Standard Deviation (Average of 10 channels)
MAV Feature	0.17	0.08	0.24
RMS Feature	0.20	0.10	0.27
VAR Feature	0.04	0.00	0.13
WL Feature	4.71	2.93	5.35
ZC Feature	0.01	0.00	0.11
SSC Feature	153.77	157.10	76.19
WAMP Feature	4.38	0.20	13.91
MNF Feature	0.09	0.08	0.04
MDF Feature	0.05	0.03	0.05
PF Feature	0.0004	0.00	0.0010
PSD Feature	0.0017	0.0009	0.0021
TP Feature	0.83	0.46	1.07
TF_Energy_Feature	389.57	223.09	485.15

Table 4: Categorised statistical features

Category	Features
Best informative Features	SSC, WL, TP, TF_Energy
Moderate informative Features	WAMP, RMS, MNF, MDF, PSD
Less informative features	MAV, VAR, ZC, PF

### 5.2.2 Analyse Outliers in best features.

The below box plot (figure. 22) indicates the graphical representation of data distribution, central tendency, variability, and presence of outliers. It's helpful to identify the overall range data distribution in each channel and potential anomalies. The highest number of outliers are presented in total power feature (TP feature) and time-frequency energy feature (TF-Energy feature), indicating high variability. Wavelength feature (WL features), generally a high number of outliers are presented across all channels. In the slope sign change feature (SSC feature), some channels (4, 5, and 6) having no outlier are presented, which might imply more stability in those features. See table 05. Outlier in each channel for different features.

Table 5: Outlier in each channel for different features

Channels	SSC Features	WL Features	TP Features	TF Energy Features
Outlier channel 1	676	295	335	318
Outlier channel 2	234	286	325	247
Outlier channel 3	252	357	461	445
Outlier channel 4	0	548	614	552
Outlier channel 5	0	594	730	640
Outlier channel 6	0	611	750	661
Outlier channel 7	1,144	545	683	683
Outlier channel 8	375	306	479	416
Outlier channel 9	1	616	638	587
Outlier channel 10	967	497	532	447

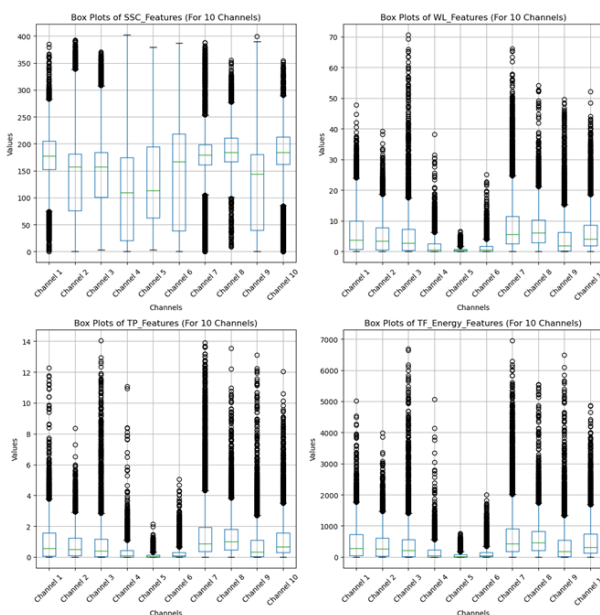


Figure 22: Graphical representation of data distribution

### 5.2.3 Find the best classification model based on a variety of experiments.

1. Finding the best ML model and features based on accuracy and processing time.

In terms of accuracy comparison, the Random-Forest model has the highest average accuracy (78.46%) across all models, but cost of computation is higher. It performs well in combined all time-domain

features (91.7%) and combined all time, frequency, time-frequency domain features (90.21%). Secondly, the KNN model is the second highest performer in average accuracy (75.50%). It also shows strong performance with combined all time-domain features (91.07%) and combined all time, frequency, time-frequency domain features (86.59%). Thirdly, artificial neural networks (ANN) perform quite well, with an average accuracy of 72.94%, and they also perform well in combined all-time domain features (90.49%) and combined all-time, frequency, time-frequency domain features (88.42%). Support Vector Machine (SVM) has an average accuracy of 60.88%, and it performs well in combined all-time, frequency, and time-frequency domain features (80.43%). Finally, Linear Discriminant Analysis (LDA) has the lowest average accuracy (57.80%) and moderate performance in combined all-time, frequency, and time-frequency domain features (70.06%).

In terms of processing time, LDA has the best model, and the average processing time is 14.72 seconds. KNN and SVM have nearly similar processing times of 14.85 seconds and 16.99 seconds, respectively. Random forest has slightly slower average processing time of 18.26 seconds. Finally, ANN is the slowest model, with an average processing time of 36.49 seconds. See table 6.

Table 6: Comparison of classifiers based on accuracy and processing time

Models	Window Size: 500-250									
	LDA		SVM		KNN		Random Forest		ANN	
Features	Total Processing time(seconds)	Accuracy	Total Processing time(seconds)	Accuracy	Total Processing time(seconds)	Accuracy	Total Processing time(seconds)	Accuracy	Total Processing time(seconds)	Accuracy
MAV	1.1	59.84	3.04	61.72	1.29	85.96	4.38	89.12	14.83	81.52
RMS	0.63	60.7	2.39	62.96	0.85	88.13	4.01	89.45	14.49	83.12
VAR	0.87	59.86	2.57	63	1.1	76.49	4.16	89.86	20.42	72.44
WL	0.66	58.79	2.79	61.25	0.89	87.64	3.97	88.3	7.4	82.61
ZC	1.82	47.25	1.47	47.25	1.51	47.25	1.68	47.29	44.79	47.25
SSC	1.23	44.6	3.98	47.25	1.46	72.24	3.27	70.06	14.17	63.59
WAMP	0.91	54.97	3.22	55.91	1.14	60.66	1.86	68.73	43.78	63.82
MEV	5.61	53.02	7.88	55.65	5.84	60.7	9.48	62.18	26.06	58.13
MDF	4.62	54.42	6.38	55.83	4.85	64.13	6.37	67.25	26.55	60.19
PF	0.62	48.73	3.37	49.86	0.86	48.62	1.14	49.4	12.44	49.98
PSD	3.62	60.7	5.7	63.43	3.85	67.02	7.18	69.24	25.46	62.53
TP	0.19	60.7	2.08	63.43	0.41	67.02	3.76	69.24	23.43	62.07
TP_Energy	85.58	59.88	87.59	62.03	85.8	85.42	89.18	86.51	108.61	79.53
Combined all Time-domain features (MAV, RMS, VAR, WL, ZC, SSC, WAMP)	6.77	67.49	9.17	76.3	6.53	91.07	12.08	91.7	19.46	90.49
Combined all Frequency-domain features (MNF, MDF, PF, PSD, TP)	14.71	63.66	18.1	69.82	14.57	79.06	21.47	86.82	26.83	81.29
Combined all Time, Frequency, Time-Frequency domain features (MAV, RMS, VAR, WL, ZC, SSC, WAMP, MNF, MDF, PF, PSD, TP, WF)	107	70.06	110.16	80.43	106.66	86.59	116.14	90.21	115.13	88.42
Average	14.72	57.80	16.99	60.88	14.85	75.50	18.26	78.46	36.49	72.94

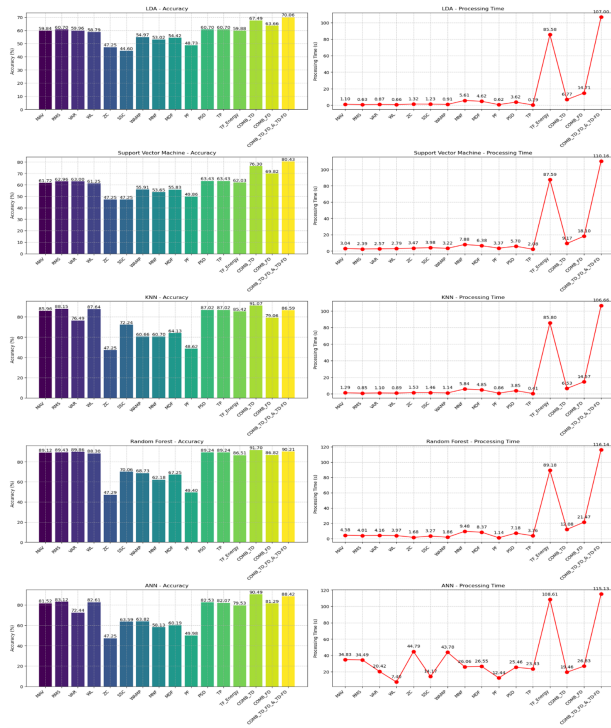


Figure 23: Feature accuracy and processing time for different classifiers.

2. Find the best model based on mean accuracy and standard deviation.

This session focuses on determining optimal model and feature selection based on mean accuracy and standard deviation, with higher mean accuracy indicating better prediction performance and lower standard deviation indicating consistency. In our scenario, KNN and Random Forest achieve the highest mean accuracy consistently for most of the features. Particularly, the combined time-domain features of KNN reach 91.70% and Random Forest achieve 90.74%. The LDA and ANN generally perform lower than KNN and Random Forest, and their mean accuracy ranges in between 60 and 70 percent. SVM provides moderate performance, and the highest mean accuracy reached 81.74% at the combined time, frequency, and time-frequency domain feature. The noticeable point is that some features, such as MAV, RMS, PSD, TP, and TF-Energy features, are

consistently achieving more than 80 percent mean accuracy in KNN and Random Forest models. In the case of standard deviation, KNN and Random Forest exhibit lower standard deviations, indicating more consistent performance. The range of standard deviation of KNN is mostly under 1.5, with the lowest at 0.54. Overall KNN and Random-Forest are performed better than other classifiers.

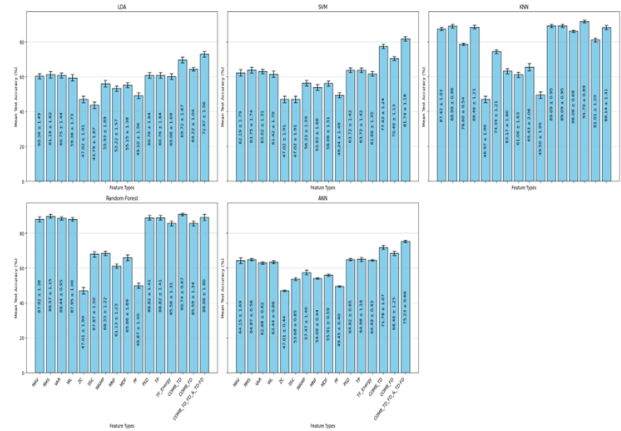


Figure 24: Mean accuracy and standard deviation for different classifiers.

3. Find the best model based on mean classification error.

The mean classification error in different models, the KNN and Random Forest, has the lowest error rate (23.19% and 22.35%), and it indicates the best-performing models. See figure 24 for classification error comparison for different classifiers. Finally, see Table.07 for the best classifier models and their best features.

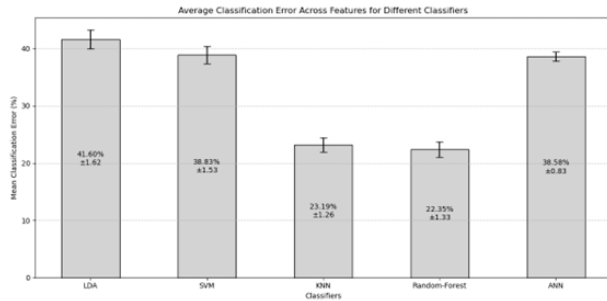


Figure 25: Classification error comparison for different classifiers



Figure 26: Confusion matrix for best classifier models.

Table 7: Best classification models and their best features

Top Models and Their Best Features					
KNN	Mean Accuracy (%)	Std Deviation	Random-Forest	Mean Accuracy (%)	Std Deviation
Combined Time-Domain Feature	91.70	0.89	Combined Time-Domain Feature	90.74	0.67
Power Spectral Density Feature (PSD)	89.09	0.95	Root Mean Square Feature (RMS)	89.57	1.15
Total Power Feature (TP)	89.09	0.95	Combined Time, frequency, and Time-Frequency Domain	88.98	1.80
Wavelength Feature (WL)	88.82	1.41	Total Power Feature (TP)	88.82	1.41
Mean Absolute Value (MAV)	87.42	1.03	TF Energy Feature	88.82	1.31

4. Confusion matrix for KNN model with their best features.

The confusion matrix helps to identify which classes perform well or not in their classifier model. Random Forest indicates good performance in class 0 (6019) and weakest true positive in class 10 (317). Classes 2 and 7 have also provided better true positives for effective classification. On the other hand, classes 10 and 11 need more attention to improvement. The KNN classifier model performed significantly well in class 0, showing a high number of true positives (5934). The weakest performance was class 10, with the fewest true positives (277). Finally, Random Forest is slightly better than KNN, especially in classes with lower true positives. See Figure 25 confusion matrix for best classifier models.

### 5.3 Part. C Individual Subject Analysis (S1\_A1\_E1 to S27\_A1\_E1).

#### 5.3.1 Discover the best features for each subject.

In this part, deal with individual subjects (1 to 27) and find the accuracy in all time- domains and frequency- domains. The combined both time and frequency domain features into a single data set to find the accuracy in KNN and Random Forest models. See table 08 for the best features in individual subjects. Based on KNN and Random Forest mean testing accuracy comparison, the Random Forest classifier has a higher than k-NN (Random Forest: 93.18% and KNN: 90.20%). See figure 26 for mean accuracy for combined time and frequency domain features in individual subjects.

Table 8: Best features in individual subjects

Subjects	Features name	Random forest Testing Accuracy	KNN Testing Accuracy
S1	Combined_Time_and_Frequency_Domain_Feature	95.91	94.55
S2	Combined_Time_and_Frequency_Domain_Feature	95.69	93.97
S3	Combined_Time_and_Frequency_Domain_Feature	95.95	93.69
S4	Combined_Time_and_Frequency_Domain_Feature	92.47	89.96
S5	Combined_Time_and_Frequency_Domain_Feature	92.86	84.92
S6	Combined_Time_and_Frequency_Domain_Feature	93.64	93.22
S7	Combined_Time_and_Frequency_Domain_Feature	95.30	91.45
S8	Combined_Time_and_Frequency_Domain_Feature	96.83	95.93
S9	Combined_Time_and_Frequency_Domain_Feature	89.18	83.98
S10	Combined_Time_and_Frequency_Domain_Feature	94.89	89.79
S11	Combined_Time_and_Frequency_Domain_Feature	88.48	84.77
S12	Combined_Time_and_Frequency_Domain_Feature	92.83	87.34
S13	Combined_Time_and_Frequency_Domain_Feature	95.77	93.90
S14	Combined_Time_and_Frequency_Domain_Feature	92.31	88.89
S15	Combined_Time_and_Frequency_Domain_Feature	96.14	92.27
S16	Combined_Time_and_Frequency_Domain_Feature	93.27	89.69
S17	Combined_Time_and_Frequency_Domain_Feature	94.55	92.73
S18	Combined_Time_and_Frequency_Domain_Feature	89.47	80.70
S19	Combined_Time_and_Frequency_Domain_Feature	93.99	92.70
S20	Combined_Time_and_Frequency_Domain_Feature	93.30	93.30
S21	Combined_Time_and_Frequency_Domain_Feature	97.10	93.78
S22	Combined_Time_and_Frequency_Domain_Feature	89.52	88.31
S23	Combined_Time_and_Frequency_Domain_Feature	90.64	87.66
S24	Combined_Time_and_Frequency_Domain_Feature	96.09	94.78
S25	Combined_Time_and_Frequency_Domain_Feature	92.08	90.42
S26	Combined_Time_and_Frequency_Domain_Feature	92.14	89.08
S27	Combined_Time_and_Frequency_Domain_Feature	85.37	83.74

### 5.3.2 Find the best classification model for individual subject.

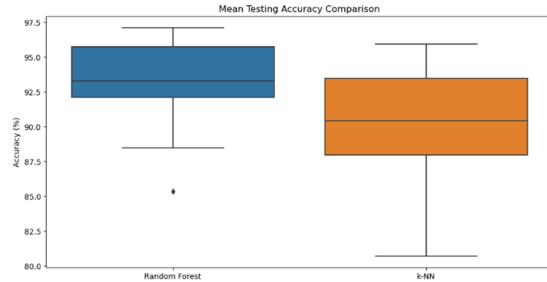


Figure 28: Mean testing accuracy for Random-forest and KNN

During all subjects, Random Forest performed slightly better than the KNN model, and the difference in accuracy ranged from 0.90 percent to 5.10 percent. The top performing subject is 21, and it achieves 97.10 percent in the random forest classifier and 95.93 percent in subject 8. In subjects 8, 15, and 24, they also indicate better accuracy just above the 96 percent in the Random Forest classifier, and similarly, subjects 24 and 1 achieve 94.78 and 94.55 percent, respectively. The top subjects based on accuracy in both classifiers are 1, 2, 3, 8, 13, 21, and 24. The lowest accuracy (85.37%) achieved in subject 27 at Random Forest and KNN is subject 18 (80.70%). See table 08 for best features in individual subjects. The high-performance subjects (S21, S8, S24, S15, and S3) are exceedingly above 95 percent. The highest accuracy subject remains the same as 27 (97.10%). Several subjects, including S13, S1, S2, and S17, are performed in moderate as compared to the highest accuracy subject (S27).

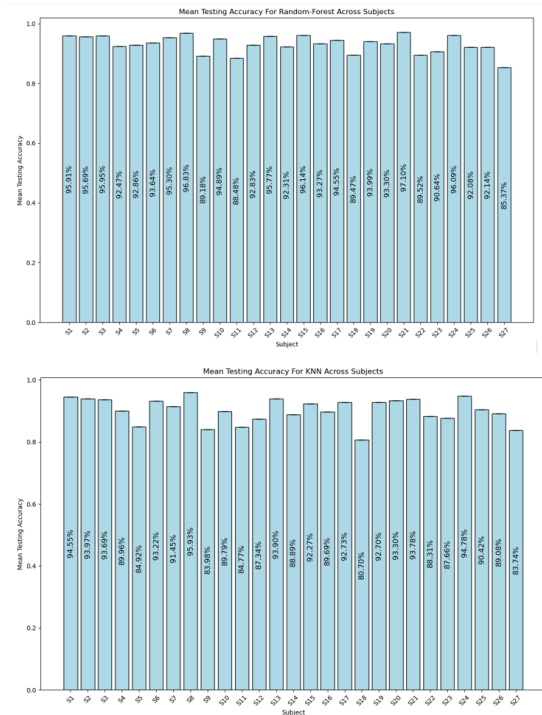


Figure 27: Mean accuracy for combined time and frequency domain feature in individual subjects.

### 5.3.3 Perform a statistical analysis test on three different features.

This statistical non-parametric Friedman test [56] was used to detect differences in treatments across multiple test attempts. For this experiment, I chose three different features (wavelength (WL), total power (TP), and time-frequency energy (TF en-

ergy)). See table 09 for testing accuracy and ranking for three different features. Then, perform the Friedman test to statistically compare the accuracy results across the three features. The results indicated a Friedman test statistic of 31.98 and a p-value of  $1.14 \times 10^{-7}$ , demonstrating a significant difference in performance between the features. A lower mean rank indicates better performance; in the Friedman test and ranking analysis, the tf-energy feature (time-frequency domain) has the best average rank (1.22), meaning it performed the best as compared to the other features across all subjects. Overall, the lowest p-value indicates that the observed differences in performance are statistically significant and not due to random chance. This significant result provides tf-energy features, which is the best performance feature.

Table 9: Testing accuracy and ranking for three different features.

Subjects	Testing accuracy			Ranking		
	wl_features	tp_features	tf_energy_features	wl_features	tp_features	tf_energy_features
S1	95.45	93.18	92.73	3	2	1
S2	92.24	95.26	91.38	2	3	1
S3	95.95	95.95	93.69	2.5	2.5	1
S4	90.80	93.31	90.38	2	3	1
S5	88.10	93.25	84.52	2	3	1
S6	89.41	94.92	87.71	2	3	1
S7	93.16	94.44	90.17	2	3	1
S8	95.93	96.38	94.57	2	3	1
S9	87.88	88.31	83.55	2	3	1
S10	94.89	94.47	92.77	3	2	1
S11	87.24	89.71	83.95	2	3	1
S12	85.23	90.72	83.54	2	3	1
S13	92.49	96.71	92.49	1.5	3	1.5
S14	85.47	90.60	81.62	2	3	1
S15	93.56	95.71	90.99	2	3	1
S16	91.48	91.93	91.48	1.5	3	1.5
S17	95.00	95.45	95.00	1.5	3	1.5
S18	86.84	88.60	79.82	2	3	1
S19	90.13	93.56	87.98	2	3	1
S20	93.75	91.52	90.18	3	2	1
S21	95.44	95.02	95.44	2.5	1	2.5
S22	87.90	89.92	85.08	2	3	1
S23	86.81	91.06	86.38	2	3	1
S24	93.04	95.22	90.43	2	3	1
S25	91.25	93.75	90.00	2	3	1
S26	89.96	88.65	90.83	2	1	3
S27	81.71	87.40	83.33	1	3	2
	<b>Mean Ranking</b>			<b>2.06</b>	<b>2.72</b>	<b>1.22</b>

## 5.4 Part. D Propose the hybrid model.

This hybrid approach is experimented with in combining 27 subjects into a single dataset and extracting the best features such as wavelength, total power, and time-frequency energy. Then, after these features

are extracted, and combine into a single feature and named as hybrid features. It contains multiple information from various domains. Finally, calculate the testing and training accuracy using KNN and Random Forest. See table 10 for an accuracy comparison for hybrid models.

Table 10: Accuracy comparison for hybrid models.

Metric	KNN	Random-Forest
<b>Testing Accuracy</b>	88.89%	90.49%
<b>Training Accuracy</b>	93.33%	100.00%
<b>Macro Average Precision</b>	0.85	0.9
<b>Macro Average Recall</b>	0.82	0.84
<b>Macro Average F1-Score</b>	0.83	0.86
<b>Weighted Average Precision</b>	0.89	0.91
<b>Weighted Average Recall</b>	0.89	0.9
<b>Weighted Average F1-Score</b>	0.89	0.9

Random Forest outperformed KNN in accuracy, precision, recall, and F1-score metrics, but has perfect training accuracy (100%), potentially indicating overfitting. KNN computation time is more efficient in real-time scenarios. A combined hybrid approach with 27 subjects helps learn more robust patterns but increases training data size. This approach may capture variations and simplify the model development process.

## 6 Conclusions

This research has optimized key components of EMG pattern recognition for prosthetic hand control using the Ninapro DB1 dataset. Key findings include an optimal window size of 500-250 ms, providing a balance between 69.12% accuracy and 15.76 s processing time; Model 4 filtering condition (stimulus=restimulus), achieving 75.31% accuracy with improved SNR and MSE; statistically selected features such as slope sign changes, wavelength, total power, and time-frequency energy, with TF-energy outperforming others (Friedman rank 1.22,  $p=1.14 \times 10^{-7}$ ); and classifiers KNN and Random Forest, with RF yielding up to 97.10% in individual analysis. The proposed hybrid model integrates these elements to

achieve 90.49% accuracy, demonstrating robustness across combined and individual datasets.

Limitations include reliance on healthy subjects, potentially limiting generalizability to amputees, and absence of deep learning comparisons, which may offer higher accuracies in complex scenarios. Future work could involve clinical trials with amputee participants, integration with real-time hardware for edge computing, and hybridizing with DL models to further enhance performance. These advancements contribute to more efficient and natural prosthetic control systems.

## 7 Acknowledgement

I express my heartfelt gratitude to the following people and institutions for their invaluable contributions to this work:

Dr. Simon Stuttaford provided excellent assistance and insightful criticism throughout the project's exploration.

Dr. Rishad Shafik and Dr. Kabita Adhikari for their valuable insights into machine learning fundamentals and inspiration for choosing machine learning based project.

Dr. Mathew Dyson for his significant suggestions and advice in improving the entire project.

Dr. Andrew Smith for assisting with the technical documentation template format and providing helpful comments.

With gratitude to the management team at Newcastle University for providing a top-notch computer lab in the Stephenson Building and granting permission to operate after hours.

Finally, my friends and family support and encourage me to accomplish my dissertation work.

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## 9 Appendix

Data and Code Availability The source code for the EMG pattern recognition analysis, including the feature extraction and classification models, is available at the following GitHub repository:

<https://github.com/Arun101DotCom/EMG-Pattern-Recognition-Analysis>