CMDA: A TOOL FOR CONTINUOUS MONITORING DATA ANALYSIS

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ABSTRACT

Over the last few years, with the growth of time-series collecting and storing, there has been a great demand for tools and software for temporal data engineering and modeling. This paper presents a generic workflow for time series data research, including temporal data importing, preprocessing, and feature extraction. This framework is developed and built as a robust and easy-to-use Python package, called CMDA, with a modular structure that offers tools to prepare raw data for analyzing any temporal data structure for scientists and non-experts.

Keywords Time Series · Feature Extraction · Preprocessing

1 Introduction

Data generation and collection in real-time is more ubiquitous than ever due to technological progress in sensors, transmission, and data storage. This has resulted in enormous datasets, namely Big data, in various fields such as engineering, medical, physical, and social sciences. While effective modeling and simulation of such data sets can offer rich information for interpretation, prediction, monitoring, training, and decision making; their scope and complexity create significant design and implementation issues [3].

For instance, in the healthcare systems, continuous monitoring data have developed into a breakthrough for health diagnosis, subsequent treatment, and proactive health tracking [1]. The recent advancements in wearable technology and monitoring systems have offered unseen opportunities for learning mechanisms of biomedical processes from temporal data as they provide data on physiological occurrences that indicate an individual's health and comfort. However, it must be taken into account that the analysis of continuous monitoring data is highly affected by data quality issues and that even complex monitoring devices can monitor only a small subset of medical parameters involved in the process under consideration. Furthermore, the required analysis strategies for monitoring data vary significantly between monitoring a few parameters assessed on dense sampling rates (e.g., ECG, EEG, or glucose data) and monitoring several parameters over hours with low sampling frequency, as in ICU. On account of this, there is a significant demand for robust and easy-to-implement tools to facilitate continuous data wrangling and feature engineering. In order to address this, we introduce the CMDA package to provide a comprehensive monitoring data analysis workflow. CMDA, developed and built as a Python package, is a standard framework for time series data research, including temporal data importing, preprocessing, and feature extraction.

2 Structure

A general workflow for continuous data analysis consists of four steps: importing the time-series records, preprocessing (missing value imputation and denoising), feature extraction, and model creation.

Initially, continuous data are imported from various sources. Secondly, the imported data will be preprocessed based on their quality to verify that only the relevant data reaches the later steps. This can involve eliminating artifacts from the data (via filtering and outlier removal techniques) and windowing signals to a set of time bins. In the final step, before modeling, significant features relevant to a particular problem are derived from acquired filtered data.

Generally, the goal of feature extraction or feature engineering is to extract the masked characteristics of raw temporal data. In other words, it can be described as a set of features corresponding to the signal's specific form or pattern. Additionally, feature extraction is generally a procedure that reduces the dimensionality of an input signal or compresses its data and thus reduces the number of resources required to analyze it. Extracting features from a raw signal typically results in low dimensional data that can help to accurately elaborate the raw data and aid in developing a practical and interpretable input for machine learning.

CMDA offers a set of tools for the mentioned steps in a modular structure, allowing users to use modules individually or as part of another project. The process can be integrated into a pipeline, which can be executed in a parallel fashion. Figure 1 depicts the components of the feature engineering pipeline. The other noteworthy attribute of CMDA is that besides the built-in feature functions, user-defined functions can be added to the pipeline. This enables more adaptable use of the feature engineering module. A detailed description of the package architecture, including the data importing modules and feature extraction modules, as well as the list of the built-in feature functions, can be found in the next section along with tutorials in Appendix A.

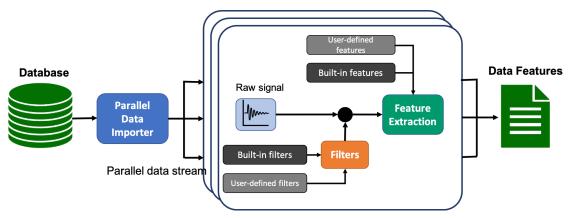


Figure 1: Illustration of the **CMDA** workflow. The pipeline consists of three components. During feature engineering, data records are imported into the pipeline consecutively or in parallel. After filtering and feature extraction, the data features are stored in a file. The features can be used later for any ML task.

3 Architecture

3.1 Data Importer Module

The data importer (cmda.import_data) is the first component in the pipeline. It offers tools for loading multiple CSV files from a server or local disk into the python coding environment (cmda.import_data.ImportCSV). It expects a CSV file where signals are represented by columns and time steps by rows.

The importer module also supports importing DAT files directly from PhysioNet databases (cmda.import_data.ImportWFDB) using WFDB package APIs [11]. PhysioNet, a funded project by the USA National Institutes of Health, is a gigantic and expanding archive of digital recordings of physiologic signals and corresponding data for the use of medical research [4].

The importer functions can save computation time by parallelizing the data loading process. Further parallelization is possible with long time series, that require windowing and segmentation. Here, the rolling window importer function (cmda.import_data.RollingWindowCSV) performs time binning and distributes each window's importing process on available nodes in parallel mode.

3.2 Preprocessing Module

The cmda.preprocessing.Filters module contains functions for applying different filters and outlier removal on a numeric signal. Users can select from a number of built-in filters, through a dynamic attribute of the Filters object, called .add that gives the user access to the functions in Table 1.

Feature function	Feature	Description
rm_outlier	Remove Outlier	Removing outliers based on a <i>low</i> and <i>high</i> threshold
rm_outliers_quantile	Remove Quantiles	Removing the values in an array based on <i>low</i> and <i>high</i>
		defined quantiles
butter_filter	Butterworth Filter	A predesigned Butterworth filter that can be used as a
		band-pass denoisng tool
interpolate	Interpolation	A <i>linear</i> or <i>cubic</i> interpolation tool that can be used for
		imputing missing values in an array

Table 1: Built-in preprocessing features

Moreover, user-defined filters can be added easily to the Filters object instance via .udf.add. This functionality makes it possible to tailor the filters to the needs of the user.

3.3 Feature Extraction Module

cmda.feature_extraction.Features is a tool for extracting multiple features from a numeric time series array. It provides several ready-to-use features of time domain, spectral domain, nonlinear, and wavelet that can be selected and added to a Features object instance via .add function.

Again, users can define and add custom functions to the Features object instance via .udf.add. The available built-in function in the current version of the package are described in the following sections.

3.3.1 Time-domain Features

Time-domain features are the signal descriptors in a time analysis span. The list of all built-in time-domain features is shown in Table 2. These functions are defined using the NumPy [5] package.

Feature function	Feature	Description
mean	Mean	Arithmetic mean of an array elements
max	Maximum	Maximum value of an array elements
min	Minimum	Minimum value of an array elements
median	Median	Median value of an array elements
std	Standard Deviation	Standard deviation of an array elements
skewness	Skewness	Skewness of an array elements
kurtosis	Kurtosis	Kurtosis of an array elements
p2p	Peak-to-Peak	Peak to peak difference (maximum - minimum values) of
		an array elements
rms	Root Means Square	Root means square of an array elements
zcr	Zero Crossing Rate	The rate at which an 1-D array changes from positive to
		zero to negative or from negative to zero to positive
mad	Mean Absolute Deviation	The median of the absolute deviations an array. Also
		called median of slopes
mns	Mean Slope	The mean differences of an array

Table 2: Built-in time-domain features

3.3.2 Frequency-domain Features

Frequency-domain features are signal characteristics in the frequency domain, which are extracted from the Fourier transform of the signal. The list of all built-in frequency-domain features is shown in Table 3. These functions are defined using the NumPy [5] and SciPy [10] packages.

Feature function	Feature	Description
periodogram	Periodogram	The power spectrum density of an array [8]
welch	Welch Periodogram	The power spectrum density of an array
		computed using welch method.
mnf	Mean frequency	The center of the distribution of power
		across frequencies. It is calculated as the
		sum of product of the power spectrum and
		the frequency divided by the total sum of
		the power spectrum [8].
mdf	Median frequency	The frequency at which the power spectrum
		is divided into two parts with equal powers
	Mariana Garage	[8].
vcf	Variance of central fre-	The variance of power spectrum amplitudes
,	quency	from its mean frequency [8].
peaks	Peak Frequency	The frequency at which the maximum of
	Danier and atmospheric	power spectrum occurs [8].
psr	Power spectrum ratio	The ratio between the energy around the maximum value of the power spectrum and
		the whole energy of the power spectrum
		[8].
band_power	Band Power	The power spectral density in a specified
		frequency band.
band_std	Band Standard Deviation	The power spectral standard deviation in a
		specified frequency band.
band_mnf	Band Mean Frequency	The mean frequency of a specified fre-
		quency band.
band_mdf	Band Median Frequency	The median frequency of a specified fre-
		quency band.

Table 3: Built-in frequency-domain features

3.3.3 Entropy Features

Shannon entropy and its derivations are methods for estimating the complexity of dynamic systems. Table 4 shows the available entropy built-in functions in the CMDA package.

Feature function	Feature	Description
entropy	Shannon Entropy	A technique for estimating the irregularity in temporal
		data [9]
sample_entropy	Sample Entropy	A technique for estimating the complexity of dynamic
		physiological systems based on the existence of patterns
		[9]
perm_entropy	Permutation Entropy	A derivation of Shannon entropy, used to assess the com-
		plexity of time series data, proposed by Bandt[2]. It quan-
		tifies the complexity of a system based on similar ordinal
		patterns [2].
spectral_entropy	Spectral Entropy	A measure of the uniformity of a power spectrum fre-
		quency distribution [7].

Table 4: Built-in entropy features

3.3.4 Wavelet Features

Wavelet features stem from the wavelet transform method which transforms from joint time to frequency. It was developed to deal with non-stationary signals and can be used for discrete and continuous data. As the continuous method is resource intensive the Discrete Wavelet Transform (DWT) is extensively used. A problem lies within the downsapling resolution of DWT. With higher decomposition levels the time invariance property is no longer valid. The Stationary wavelet transform (SWT) [6] is a wavelet transform algorithm designed to overcome the downsample

problem of DWT and preserves the time information. In the CMDA package, we offer a built-in swt_features function that can return a set of time-domain and frequency-domain features of the decomposition levels coefficients. The parameters in Table 5 can be given to the function.

Feature Function	Feature
mnf	Mean Frequency
psr	Power Spectrum Ratio
peak	Peak Frequency
mds	Median Absolute Deviation
mns	Mean Slope
see	Shannon Entropy
perm-ent	Permutation Entropy
nse	Normalized Energy

Table 5: Built-in SWT features. The description of each parameter can be found in Tables 2, 3, and 4.

3.4 Feature Engineering Pipeline

The cmda.pipeline.Pipeline is a tool that combines and automates importing, pre-processing and feature extraction of multiple records (files). It recieves an importer object instance (cmda.import_data), a feature object instance cmda.preprocessing.Features, and an optional filter object instance (cmda.preprocessing.Filters), which are combined into a pipeline. The user can run the process consecutively or in parallel by accessing its .run function.

4 Discussions

CMDA Python package is developed with the purpose of creating a monitoring data analysis workflow that includes a wide range of functional elements between from reading multiple data to denoising and feature extraction. It is highly flexible and through its modularity and support of user-defined features, it can be integrated in various applications. The package provides a fast and straightforward workflow with minimal coding requirements, making it feasible for researchers to prepare ready-to-analyze, extensive monitoring data sets that are suitable for sophisticated modeling strategies. In the future development of CMDA, we aim to add more built-in features and pre-processing functions to cover broader applications. In addition, embedding unsupervised feature engineering methods such as deep autoencoders and signal-to-image transformation is another objective of the package expansion. Until then, we hope CMDA benefits healthcare data expertise by providing a standard setting for monitoring data analysis.

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A Examples

A.1 Feature Object

cmda.feature_extraction.Features is a tool for extracting multiple features at the same time from a numeric signal x. One can select from a wide variety of built-in features, as well as adding user-defined features. This functionality makes it possible to tailor the features to the need of the user. The following subsections show examples of time domain, frequency domain, and user-defined feature extraction.

A.1.1 Data Import

We start by using sample data contains an ECG signal and ABP (Arterial Blood Pressure) signal with a length of 10 seconds and 125 Hz sampling rate. The toy data can be accessed from cmda.data.ecg_abp_sample.

```
import numpy as np
import matplotlib.pyplot as plt

from cmda.data import ecg_apb_sample

data = ecg_apb_sample()

ecg = data["ECG"]
abp = data["ABP"]

fs = 125
t = np.linspace(1, 1 / fs, len(ecg))
fig, (ax1, ax2) = plt.subplots(2, sharex=True)
ax1.plot(t, ecg)
ax1.set_ylabel("ECG [mV]")
ax2.plot(t, abp, c="red")
ax2.set_ylabel("ABP [mmHg]")
ax2.set_xlabel("Time [seconds]")
plt.show()
```

Listing 1: Importing a sample data

The output graph can be seen in Figure 2.

A.1.2 Time-domain feature extraction

In this example, we extract some built-in time-domain features of the ABP signal. To start, we call the cmda.feature_extraction. Features object and add the desired features using the add function.

After adding the features, we apply the transform function to the signal which calculates the features. The transform function returns the extracted features as a python dictionary.

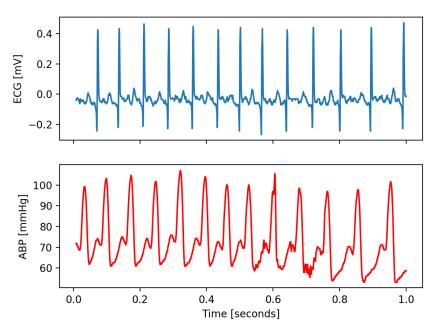


Figure 2: Imported ECG and ABP sample signals

```
from cmda.feature_extraction import Features
2 from cmda.data import ecg_apb_sample
 data = ecg_apb_sample()
 x = data["ABP"]
  feature = Features()
9 # add built-in time-domain features
10 feature.add.mean()
feature.add.max()
12 feature.add.min()
13 feature.add.median()
14 feature.add.skewness()
15 feature.add.kurtosis()
16 feature.add.std()
17 feature.add.p2p()
18 feature.add.zcr(center=True)
20 feature.transform(x=x, fs=125)
```

Listing 2: Time-domain feature extraction

Output:

```
1 {'mean': 72.385,
2   'max': 107.03125,
3   'min': 53.125,
4   'median': 69.53125,
5   'skewness': 0.9154627781784881,
6   'kurtosis': -0.15299373030467667,
7   'std': 13.268216310510619,
8   'p2p': 53.90625,
9   'zcr': 42}
```

Listing 3: Time-domain feature extraction output

A.1.3 Frequency-domain feature extraction

In this part, we extract some built-in frequency-domain features of the ECG signal. Similar to the previous example, we call the cmda.feature_extraction.Features object, and append the desired features using the add function. Using the transform function returns the extracted features as a python dictionary.

```
from cmda.feature_extraction import Features
from cmda.data import ecg_apb_sample

data = ecg_apb_sample()
x = data["ECG"]

feature = Features()

# add frequency-domain built-in features
feature.add.mnf(spectrum="ps")
feature.add.mdf(spectrum="ps")
feature.add.stdf(spectrum="ps")
feature.add.stdf(spectrum="ps")
feature.add.psr(spectrum="welch", int_limit_ratio=0.01)
feature.add.psr(spectrum="welch", n_peaks=1, height=True, width=True)
feature.add.band_power(spectrum="ps", low=1, high=7)
feature.add.band_mnf(spectrum="ps", low=1, high=7)

feature.add.band_mnf(spectrum="ps", low=1, high=7)

feature.transform(x=x, fs=125)
```

Listing 4: Frequency-domain feature extraction

Output:

```
1 {'mnf': 9.857968968908676,
2 'mdf': 9.0,
3 'stdf': 6.657686374206612,
4 'psr_0.01': 0.1086823980353686,
5 'peak_freq_1': 3.90625,
6 'peak_height_1': 0.0009058970155242925,
7 'peak_width_1': 4.386465062242469,
8 'power_[1,7]Hz': 0.3921784430303598,
9 'mnf_[1,7]Hz': 4.277888519896743}
```

Listing 5: Frequency-domain feature extraction output

A.1.4 User-defined feature extraction

In this section, we derive some features, including a user-defined feature from the ABP signal. Similar to the previous example, we call the <code>cmda.feature_extraction.Features</code> object, Define the feature and add them to the feature object, using <code>add_udf</code> function. The <code>transform</code> function returns the extracted features as a python dictionary.

```
import numpy as np
2 from cmda.feature_extraction import Features
3 from cmda.data import ecg_apb_sample
5 data = ecg_apb_sample()
6 x = data["ABP"]
8 # Define the feature
 # Please take note, that the output must be a dictionary
10 # ,where the keys are the extracted features
 def quantile(x,low = 0.1,high =0.9):
11
      q_low = np.quantile(x,q=low)
12
      q_high = np.quantile(x,q=high)
14
      res = {"q_low":q_low, "q_high":q_high}
15
      return res
16
19 # Call the Feature object
```

```
feature = Features()

feature.add built-in features
feature.add.mean()
feature.add.max()
feature.add.zcr(center=True)
feature.add.mnf(spectrum="ps")

# add user-defined feature
feature.udf.add(quantile)

feature.transform(x=x, fs=125)
```

Listing 6: User-defined feature extraction

Output:

```
1 {'mean': 72.385,
2  'max': 107.03125,
3  'zcr': 42,
4  'mnf': 2.0056974184541025,
5  'q_low': -13.79125,
6  'q_high': 23.70875}
```

Listing 7: User-defined feature extraction output

A.2 Multivariate Feature Extraction

In this example, we show how to extract features from a multivariate signal. In the first example, we extract the same features from multiple variables variables, and in the second part, we extract different features from multiple variables. How a feature object is created can be seen in Section A.1. We use the same data as used in the Feature Object example (A.1). The data contain two signals (ECG, ABP) with a sampling frequency = 500.

A.2.1 Feature extraction with one feature object

To extract features from a data set, we use cmda.feature_extraction.extract_features function.

```
from cmda.feature_extraction import Features
from cmda.feature_extraction import extract_features
from cmda.data import ecg_apb_sample

# load the data
data = ecg_apb_sample()

# Create the feeature object
features = Features()

# Add built-in features
features.add.std()
features.add.std()
features.add.mnf()
features.add.stdf()

# extract eatures
res = extract_features(data=data, feature_obj=features)

print(res)
```

Listing 8: Multivariate feature extraction using one feature object

Output:

```
1 {'ECG_std': 0.0912508348533301,
2 'ECG_mnf': 0.0788637517512694,
3 'ECG_stdf': 0.053261490993652905,
4 'ABP_std': 13.268216310510619,
```

```
'ABP_mnf': 0.01604557934763282,
'ABP_stdf': 0.012480758701313835}
```

A.2.2 Feature extraction with multiple feature objects

In various real applications, extracting similar features from different signals may lead to wrong data compression and unnecessary information. To avoid that, we show how to create and apply signal-specific feature objects.

```
from cmda.feature_extraction import Features
2 from cmda.feature_extraction import extract_features
3 from cmda.data import ecg_apb_sample
5 # load the data
6 data = ecg_apb_sample()
8 # Create the feature object for ECG signal
9 ecg_features = Features()
# Add built-in features
12 ecg_features.add.mnf()
13 ecg_features.add.stdf()
15
16 # Create the feature object for ECG signal
17 abp_features = Features()
19 # Add built-in features
20 abp_features.add.mean()
abp_features.add.std()
22 abp_features.add.max()
23 abp_features.add.min()
24
25 # Create the feature object as a dict
26 features = {'ECG':ecg_features, 'ABP':abp_features}
28 # extract eatures
29 res = extract_features(data=data, feature_obj=features)
31 print (res)
```

Listing 9: Multivariate feature extraction using one feature object

Output:

```
1 {'ECG_mnf': 0.0788637517512694,
2 'ECG_stdf': 0.053261490993652905,
3 'ABP_mean': 72.385,
4 'ABP_std': 13.268216310510619,
5 'ABP_max': 107.03125,
6 'ABP_min': 53.125}
```

A.3 Feature Extraction Pipeline

This example demonstrates the usage of cmda.pipeline.Pipeline to extract features from a PhysioNet WFDB. We extract features from the ECGRDVQ database. The database contains 5232 extracted 10-second standard 12 lead ECG segments. The meta-data of the database (ecgrdvq_metadata) is available in cmda.data.wfdb in form of a dictionary, containing record names under record_name and their corresponding public directories public_dir. The pipeline imports the data, applies filters to the records, and extracts features from records. To import the data from PhysioNet WFDB, we use the function cmda.read_data.ReadWFDB. We create the filtering object by cmda.filter.Filters and create the feature object by cmda.feature_extraction.Features. A detailed example on creating feature objects can be found in A.1.

```
from cmda.data.wfdb import ecgrdvq_metadata
from cmda.read_data import ReadWFDB
```

```
3 from cmda.preprocessing import Filters
4 from cmda.feature_extraction import Features
5 from cmda.pipeline import Pipeline
7 # Load the meta-data
8 metadata = ecgrdvq_metadata()
10 # get the record names and their corresponding public directories
record_names = metadata['record_name']
public_dir = metadata['public_dir']
^{14} # Build the importer object using ReadWFDB
15 # Set the channels to ["II",'V1'] to import these channels exclusively.
importer = ReadWFDB(record_names=record_names, public_dir=public_dir, channels=['
     II','V1'])
17
18 # Create the filter object and add a buuterworth low-pass filter
19 filter_obj = Filters()
20 filter_obj.add.butter_filter(cutoff=60, btype="lowpass")
22 # Create the feature object
23 feature_obj = Features()
24 feature_obj.add.mnf()
25 feature_obj.add.mdf()
26 feature_obj.add.psr()
27 feature_obj.add.stdf()
28 feature_obj.add.peaks(n_peaks=1,spectrum='welch',height=False,width=False,nperseg
     =512)
29 feature_obj.add.band_power(low=0.6, high=2,spectrum='welch')
30 feature_obj.add.band_power(low=2, high=4,spectrum='welch')
feature_obj.add.band_power(low=4, high=6,spectrum='welch')
feature_obj.add.band_power(low=6, high=10, spectrum='welch')
feature_obj.add.band_power(low=10, high=15, spectrum='welch')
feature_obj.add.band_power(low=15, high=30, spectrum='welch')
36 # Build the pipeline
37 ecgrdvq_pipeline = Pipeline(importer=importer,features=feature_obj, filters=
     filter_obj)
39 # Run the pipeline
40 # Set the dataframe_output to True, to get the extracted as a Pandas dataframe
41 # n_jobs set the number of cpu cores for parallel feature extraction
42 res = ecgrdvq_pipeline.run(n_jobs=4, dataframe_output = True)
```

Listing 10: Multivariate feature extraction using one feature object

Output:

```
Running the pipeline on 5232 instances...

100% [||||||||||| 5232/5232 [00:05<00:00, 2.23it/s]

4 finished!
```

The required time for parallel running the above pipeline was around 2640 seconds for a PC with a 3,5 GHz Dual-Core Intel Core i7 processor and 16 GB 2133 MHz LPDDR3 memory. The process included retrieving the data from PhysioNet data servers.