## Technical Roadmap - Part I

This document describes the final tasks completing the technical part of your term project. Part II details the format and logical organization of the final project report. Building on your skills and experience from the past three group assignments, the tasks listed here expand on the problem scope for unsupervised intrusion detection using time series analysis and forecasting applied to stream data from a supervisory control system. There are several goals: enriching the *feature engineering phase*, allowing for models with arbitrary many states, and advancing the *anomaly detection phase* by analyzing several datasets with injected anomalies. The data analysis, the design and selection of models, and the presentation of the experimental results form the technical basis of five main aspects to be addressed in your project report (as detailed in Part II).

All groups will work with the same dataset for model training, testing and anomaly detection. This way, the results of your experiments will be comparable to the results of other groups and naturally allow for ranking model performance across all groups. The dataset is going to be uploaded to the TERM PROJECT folder listed under 'Assignments' on the course home page.

## Please complete the following four tasks:

1. Feature Scaling is an essential preprocessing technique of input data for machine learning algorithms and beyond. Two widely used feature scaling techniques are *normalization* and *standardization*. For each of the following questions explain clearly and concisely: (I) Why scaling features of a dataset is necessary? — (II) What does normalization and standardization do to the data and the noise? — (III) Which feature scaling method do you choose for HMM training for anomaly detection purposes and why? Your answers to all three questions should not exceed one page in total but provide technical descriptions including the use of mathematical notation. (Please check the information available online.)

Note: You need to apply your scaling method of choice to the dataset provided across all the features prior to engaging in the Feature Engineering task.

2. Feature Engineering. Choose a subset of the response variables for training of multivariate HMMs on normal electricity consumption data. For deciding on the subset of variables that are most suitable for training your models, you need to perform a Principal Component Analysis (PCA)<sup>1</sup>, a superior alternative to using a correlation matrix. Provide a proper rational for your final choice of response variables based on your PCA results. This method is explained in more detail on the last page.

<sup>&</sup>lt;sup>1</sup> Principal component analysis is a statistical procedure that uses an orthogonal transformation to convert a set of observations of possibly correlated variables into a set of values of linearly uncorrelated variables called *principal components*.

3. HMM Training and Testing. Partition your scaled data into train and test: use the first three years for training and reserve the fourth year for testing. Choose a weekday or a weekend day and a time window between 2 to 6 hours on that day. For this time window, train various multivariate HMM models on the train data with different numbers of states. For models with at least 4 and not more than 20 states evaluate and compare the results of log-likelihood and BIC to select the 'best performing' model(s) with an overall good fit on the train data. Note that you do not need to train a model for each and every number of states across the given range. Use the test dataset to determine the best performing HMM model, i.e. one that is neither overfitted nor underfitted.

Note that you need to compare normalized log-likelihood of the train data and the test data. *Hint:* For calculating the log-likelihood of the test data, look at the fit-section on Page 15 and the forwardbackward-section on Page 21.

https://cran.r-project.org/web/packages/depmixS4/depmixS4.pdf

4. Anomaly Detection. In anomaly detection using HMMs, calculating the log-likelihood of new data streams is a foundational method for assessing how significantly a data sequence deviates from the model's learned behaviour, thereby indicating its level of anomaly. This process necessitates establishing a threshold to distinguish normal from anomalous observations. Such a threshold should be derived from the distribution of log-likelihoods for a validation set comprising known normal data, or determined through cross-validation techniques. This approach ensures that the threshold is empirically grounded, enhancing the model's accuracy in anomaly identification.

Finally, partition your test dataset into 10 roughly equal-sized subsets representing consecutive weeks. Compute the log-likelihood for each of the subsets using your "best" performing model in order to determine the maximum deviation from the train log-likelihood value. The result obtained is a threshold for the acceptable deviation of any unseen observations to be considered normal behaviour.

## PRINCIPAL COMPONENT ANALYSIS

Principal Component Analysis is a useful technique for analyzing datasets with many variables. It is basically a type of linear transformation which takes a dataset with many variables (i.e., number of responses and number of samples), and simplifies it by turning it into a smaller number of variables, called *principal components*.

This technique also allows you to visualize how the data is spread out in a dataset. The underlying mathematics is somewhat complex though, so we won't go into too much detail, but PCA gives us a number (percentage) for each variable which indicates how much variance there is in the data for that variable.

To have a better understanding, let's assume you have a year worth of multivariate data which has 7 responses. Further assume you choose a time window from <start time> to <end time> on a <weekday>. Therefore, you would have 52 samples for each of these 7 responses. After applying PCA on this data, you obtain 7 principal components. Each of these PCs is represented by a number which explains a percentage of the total variation in the dataset. If PC1 is 65%, it means it has 65% of the total variance; in other words, nearly two-thirds of the information in the dataset can be encapsulated by using this one principal component.

In this part, you should (I) compute the principal components of the original dataset; (II) plot the results (PCs); and then (III) interpret the results. In order to compute the principal components, we recommend to use the stats package (the important commands you may need are prcomp() and summary()). To plot the result we recommend to use the ggbiplot package (it is based on the ggplot package).

Please read about **Principal Component Analysis** to gain a better understanding of this concept and also refer to the documentation of the packages you use. There are also three short videos on YouTube explaining PCA in a generally understandable way. You can find these YouTube videos at the following URLs:

https://youtu.be/HMOI\_lkzW08 https://youtu.be/FgakZw6K1QQ https://youtu.be/0Jp4gsfOLMs

## **SUBMISSION**

Please submit a PDF copy of your term project report: presenting, explaining and illustrating your experimental analyses and the evaluation and interpretation of your quantitative and qualitative results, together with the <u>R code</u> of your technical solutions, through the course home page by end of Sunday, 24 NOVEMBER 2024.