Clustering techniques selection for a hybrid regression model: a case study based on a solar thermal system

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Clustering techniques selection for a hybrid regression model: a case study based on a solar thermal system

This work addresses the performance comparison between four clustering techniques with the objective of achieving strong hybrid models in supervised learning tasks. A real dataset from a bio-climatic house named Sotavento placed on experimental wind farm and located in Xermade (Lugo) in Galicia (Spain) has been collected. Authors have chosen the thermal solar generation system in order to study how works applying several cluster methods followed by a regression technique to predict the output temperature of the system.}

With the objective of defining the quality of each clustering method two possible solutions have been implemented. The first one is based on three unsupervised learning metrics (Silhouette, Calinski-Harabasz and Davies-Bouldin) while the second one, employs the most common error measurements for a regression algorithm such as Multi Layer Perceptron.

Keywords: clustering; regression; hybrid model; learning metrics; spectral clustering; Gaussian mixture clustering; agglomerative clustering; k-means

Subject classification codes: include these here if the journal requires them

1. Introduction

In general terms there are a lot of different hot topics, and of course for the most of possible applications, and regardless of the field of the final use. Representative cases of them are: ecological, zero impact, environment safety, sustainability, and so on [5, 16]. Usually, these topic examples go in opposition with other issues like benefits, comfort, luxury, etc. [22, 21]. Furthermore, it is a challenge the compromise between the two trends; for instance, people like comfort homes, and therefore, it is desirable this achievement comes from renewable energies.

In relation to energy needs, renewable energies play a key role in contributing to a reduction in environmental impact and emissions [23]. Nevertheless, the impact of the power-plant implementation itself based on renewable sources has to be taken into account, there is not usually any zero impact [30].

Because it is not possible to achieve the null impact, even with the alternatives and use of renewable energies, there is a legal obligation to optimize and plan installations with maximum efficiency [36]. Moreover, the facilities performance must be measured in accordance with the right ratios and criteria with the aim of ensuring the desired minimum impact [18].

For an optimal performance of the renewable energy systems, due to some different reasons, commonly it is necessary to make predictions of the used variables for the facility right management [20]. There are many techniques to make predictions, from the traditional ones to the most advanced through the middle ones between both [4]. When a specific system to be modelled has a performance with a very non-linear component for instance, the modelling based on hybrid systems frequently gives very satisfactory results [6, 31, 11, 9, 25, 10].

When hybrid systems are used for modelling tasks, during the clustering stage frequently is used K-means method as a standard [34]. However, there are many clustering techniques with a satisfactory performance and, in a lot of cases, with a better performance versus K-means technique [34].

The present research accomplishes a performance study of two clustering techniques, Gaussian Mixture and Spectral Clustering. For comparing their be- haviour, two approaches have been implemented. Firstly a set of error non- supervised measurements and following an MLP (Multi Layer Perceptron) regressor for establishing the quality when a hybrid model is developed. The work has been accomplished over a real system based on a solar thermal panel, installed in a bioclimatic house. The rest of the document is structured as follows. Section 2 describes briefly the case of study. After section 2, the model approach used to compare the clustering measurement is shown. After that, the techniques applied to achieve the classification are explained. Section 5 details the experiments and achieved results and finally, the conclusions and future works are exposed in Section 6.

2. Case study

The case study of this research is part of the installation of the Sotavento Galicia Foundation bioclimatic house. This Foundation was created with the aim of studying both new renewable energies and their use in building, and for this last point, they built the bioclimatic house.

Sotavento bioclimatic house

The real house is shown in figure 1, and it was built with the aim of reducing the amount of energy consumed inside. It is located in Xermade council, in Lugo, in the Sotavento Experimental Wind Farm, that is a place where the Foundation has its own wind farm to study different types of wind turbines.



Figure 1. Sotavento bioclimatic house

This research is focused on the solar thermal energy collectors, that are only a part of the whole thermal system of the house. Figure 2 shows the thermal energy

schematic of the house, which includes solar (1), biomass (2) and geothermal (3) as primary energies. The schematic is divided into three parts: generation, accumulation and consumption. The thermal energy consumption of the house is the Domestic Hot Water, DHW, (7) and the Heating system (6). The accumulation part has two different water store deposit, one is the solar accumulator (4), and the other is the DHW and Heating accumulator (5); this part also include the preheating for the DHW (8).

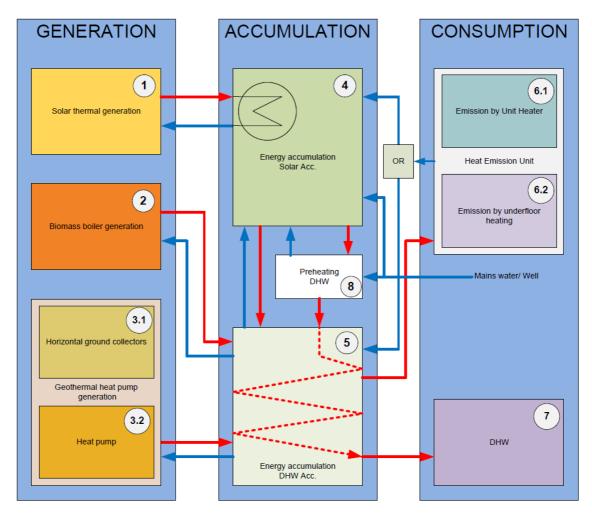


Figure 2. Thermal energy installation schematic

As the bioclimatic house is made for study not only thermal energies, the house includes also electrical energies like wind, photovoltaic and grid connection. As this type of energy is not part of the research, it is not described in this paper.

The thermal solar system is presented in figure 3, that shows the schematic of this part of the installation. This research uses only the temperature sensors S1, S2, S3 and S4, and also the flow-meter (red arrow in the figure). The solar collector (with a total surface of 20 m2) is made with eight panels, distributed in two strings of four panels. The top and the bottom string have input and output temperature sensors (S1, S2, S3 and S4); the rest of the schematic is the same for both strings. Figure 3 also includes the solar accumulator, with a capacity of 1000 L, and the necessary valves and pumps to ensure that the system could work properly.

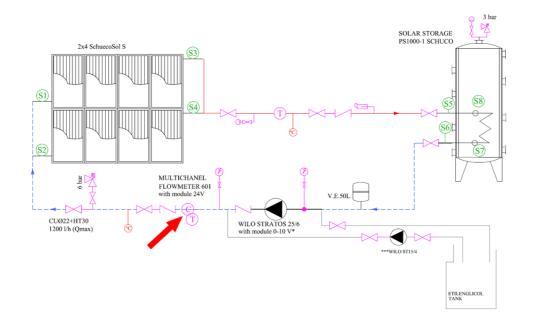


Figure 3. Solar thermal energy layout

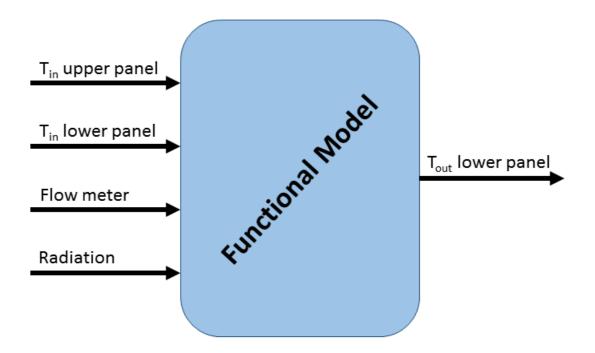
The temperature sensors used are RTD (PT1000) type, and the flow-meter is a Multical®403. For the research we also need to know the solar radiation, that is measured with a PYR-P sensor located outside the house.

Model approach

The aim of this research is to compare different clustering techniques. The way used to choose the best number of clusters is based on different metrics that allow selecting to optimal number of groups. The model approach presented in this research is used to compare the clustering algorithms. Once the optimal number of clusters is chosen, a hybrid model is trained based on the schematic shown in figure 4.

Figure 4 shows the general model, but it is important to highlight that this is a hybrid model created with several local models, as many local models as clusters. The output of the model is the output temperature of the lower string (S4 in figure 3) and the inputs are:

- The inputs temperatures of both strings (S1 and S2 in figure 3).
- The flow rate of the etilenglicol used as thermal fluid throw the panels.



• The solar radiation.

Figure 4. General schema of the functional mode

The procedure to compare the different clusters techniques, was to train one hybrid model for each technique with the same training dataset, and compare the obtained results for each one with a testing dataset. The used regression technique was an Artificial Neural Network for each internal local model.

3. Used techniques

The first step was a preprocessing step in which the data is normalized applying MinMax normalization. After the preprocessing step, four different clustering algorithms have been applied. Three metrics were used to evaluate each of the clustering techniques. The data class assigned by the clustering technique is used as an extra feature. After that, an MLP regressor is used to make predictions.

An LDA technique was used to improve the results visualization. A brief explanation of the implemented techniques is given in the following paragraphs.

3.1. Preprocessing

The MinMax method normalizes the data to fall in the [0,1] interval depending on their maximum and minimum values, according to the following expression 1.

$$\hat{x}_j = \frac{x_j - x_{min}}{x_{max} - x_{min}} \tag{1}$$

This normalization process is recommended to obtain better results when working with Multi-Layer Perceptron or clustering [14] techniques when used for regression analyses [3].

3.2. Linear Discriminant Analysis projection

When using logistic regression, sometimes it is found that, despite classes being well separated, estimation parameters are found to be unstable. In these scenarios, the Linear Discriminant Analysis (LDA) technique is recommended because it is not affected by this kind of problem. With the aid of LDA the classes separability can be maximized.

Moreover, this technique eases data transformation, obtaining the greatest separation between classes, and so it is a good technique for projection. LDA has usually been used as a method for two-dimensional projection, as is the case in this study [27].

3.3. Clustering techniques

Spectral Clustering Spectral Clustering [28] splits a dataset according to its samples' similarity graph. Both the adjacency and the degree matrix can be obtained from this graph indicating, respectively, the relationship between samples and the number of relations. Then, the corresponding Laplacian matrix can be calculated by using the degree matrix. The final step consists in using the Laplacian matrix for applying K-means on its eigenvectors, finding the corresponding clusters of samples. Because of using K-means, the number of centroids must be previously determined.

Gaussian Mixture Clustering This technique [26] takes into account the centroid, the covariance and the weight for defining clusters. These models can be defined as a combination of K Gaussian distributions. An Expectation-Maximization algorithm [13] is used to find the distributions, determining the values for the mean, the covariance and the weight of each distribution. While the K-means technique only uses the mean value, the Gaussian Mixture Clustering also takes into account the variance on the data.

Agglomerative Clustering Agglomerative Clustering [12] is a technique included in the so-called Hierarchical Clustering family of algorithms. It builds clusters by means of a number of splitting and merging processes, starting with a unique sample per cluster. In each iteration of the algorithm, a merge between the most similar clusters is performed. The process ends when all the samples belong to the same clustering.

K-Means algorithm This algorithm is one of the most popular clustering techniques. K-Means tries to separate the data minimizing the inertia of the groups [17]. This method requires to set the number of clusters before training. Each cluster is represented by its centroid μ_i , which represents the mean values of all its elements (see equation 2).

$$\sum_{i=0}^{n} \min_{\mu_{j} \in C} (||x_{i} - \mu_{j}||^{2}) (2)$$

3.4. Cluster Error metrics

The unsupervised metrics Silhouette coefficient, Calinsky-Harabasz and Davies-Bouldin have been studied for evaluating the clustering methods.

Silhouette The Silhouette coefficient is a score for evaluating the goodness of clustering algorithms, with the objective of identifying the most adequate number of clusters.

The number of clusters when using unsupervised learning algorithms may be an input parameter or may be automatically established by the algorithm itself. When included as a parameter, as is the case with K-Means algorithm, an external score must be used to find the most adequate number of clusters. The Silhouette coefficient can be used as an indicator for estimating the ideal number of clusters, where a higher coefficient means a better quality using this number of clusters.

For an observation j, the Silhouette coefficient is denoted as s(j) and calculated as:

$$s(j) = \frac{y - x}{\max(x, y)} (3)$$

Where:

• *x* is the average of distances (or dissimilarities) of observation *j* respect to the rest of observations in the cluster which *j* belongs to.

y is the minimum distance to a different cluster (not the same as observation j).
 The cluster meeting this requirement is known as "the neighbourhood of j", and would be the second-best option for j.

The Silhouette score takes values between -1 and 1.

When observation *j* is on the boundary of two clusters the value of s(j) will be close to zero.

When s(j) takes a negative value, the *j* observation must be assigned to the closest cluster.

In short:

- $s(j) \approx 1$, the assignation of the *j* observation to the cluster is correct.
- $s(j) \approx 0$, the *j* observation lies between two different clusters.
- $s(j) \approx -1$, the assignation of *j* observation to the cluster is wrong.

Calinski-Harabasz The Calinsky-Harabasz score can be obtained using the following expression (4):

$$CH = \frac{\frac{BGSS}{K-1}}{\frac{WGSS}{N-K}} = \frac{N-K}{K-1} \frac{BGSS}{WGSS} (4)$$

begin N the number of observations and K the number of clusters and with

$$BGSS = \sum_{j=1}^{K} n_j ||G^j - G||^2 (5)$$

(where G^{j} denotes, for each cluster, the dispersion of the barycenters, and G is the barycenter of the set of data as a whole. The number of samples in the cluster C_{j} is represented as n_{j})

$$WGSS = \sum_{j=0}^{K} WGSS^{j} (6)$$
$$WGSS^{j} = \sum_{i \in I_{j}} ||M_{i}^{j} - G^{j}||^{2} (7)$$

 (M_i^j) are the coefficients for the i-th row in the data matrix for cluster C_j , while I_j represents the set of indices of the observations for the C_j cluster).

Davies-Bouldin The Davies-Bouldin index is a score used for the evaluation of clustering algorithms. It uses characteristics and quantities that are inherent to the data set, and is defined as the mean value of the samples M_k (among all the clusters), as is represented in 8.

$$DB = \frac{1}{\kappa} \sum_{j=1}^{\kappa} M_j$$
(8)

where δ_j represents the mean value distance from the points belonging to the C_j cluster to their barycenter G_j , while $\Delta_{jj'}$ is the distance between barycenters G^j and $G^{j'}$ (equation 10).

$$\Delta_{jj'} = d\left(G^{j}, G^{j'}\right) = ||G^{j} - G^{j'}|| (9)$$

When the clusters are compact, smalls values are obtained for the DB index, and their corresponding centers are well separated. For this reason, the optimum number of clusters is chosen when the DB index is minimized.

$$DB = \frac{1}{\kappa} \sum_{j=1}^{K} M_j (10)$$

3.5. Regression Error metrics

The different regression models used in the study are compared using the fol- lowing error metrics (for all of them, the observed value is denoted by Y_j and the foretold value by \hat{Y}_j):

 M.A.E.: Mean Absolute Error. This metric measures differences between the real and the predicted values, having some advantages over other error scores [37].

$$MAE = \frac{1}{m} \sum_{j=1}^{m} |Y_j - \widehat{Y}_j| \quad (11)$$

• LMLS: Least Mean Log Squares. It is used as a logistic error function for both the training process and the validation error [7], equation 12.

$$LMLS = \frac{1}{m} \sum_{j=1}^{m} \log \left(1 + \frac{1}{2} \left(Y_j - \widehat{Y}_j \right)^2 \right)$$
(12)

• SMAPE: Symmetric Mean Absolute Percentage Error. The objective of this metric is to give an explanation for relative errors by using percentages [19], equation 13.

$$SMAPE = \frac{2}{m} \sum_{j=1}^{m} \frac{|Y_j - \hat{Y}_j|}{|Y_j + \hat{Y}_j|} \quad (13)$$

• MSE: Mean Squared Error. This metric can be applied in different forecast- ing problems, it can include the error variance [35] equation 14.

$$MSE = \frac{1}{m} \sum_{j=1}^{m} \left(Y_j - \widehat{Y}_j \right)^2 \quad (14)$$

• MAPE: Mean Absolute Percentage Error. This metric is one of the most usual ones for measuring the accuracy of regression problems [24], equation 15.

$$MAPE = \frac{100\%}{m} \sum_{j=1}^{m} \frac{|Y_j - \hat{Y}_j|}{Y_j} \quad (15)$$

• NMSE: Normalised Mean Square Error. This metric estimates the overall deviation between predicted and observed values [29], equation 16.

$$NMSE = \frac{1}{m} \sum_{j=1}^{m} \frac{(Y_j - \hat{Y}_j)^2}{mean(\hat{Y}_j) * mean(Y_j)} \quad (16)$$

3.6. Regression method

Multi-Layer Perceptron: A Multi-layer Perceptron (MCP) was implemented to obtain a metric for the evaluation of the previously mentioned clustering algorithms.

MLP is one of the most commonly used supervised learning techniques. The learning function for this algorithm is: $Fun(\cdot) : X^n \to X^0$. The Scikit-Learn library for Python was used to implement this technique.

A cross validation procedure was used for obtaining the optimal number of neurons for the hidden layer and the best activation function for each one. With the aid of this procedure, the MLP was trained with different parameters (number of neurons and activation function) to obtain the most suitable regression model [33, 15, 8, 2].

4. Experiments and results

This section addresses the results from clustering and regression point of view. The first one makes reference to how clustering methods have working based on a set of measurements. On the other hand, the second one, defines how a regression technique as MLP works with the clustering procedure applied previously.

4.1. Cluster

Four different clustering techniques have been evaluated with the aim of determining possible groupings of the unsupervised data. These techniques are: Spectral Clustering, Gaussian Mixture Clustering, Agglomerative clustering and K-Means. After the clustering step, the assigned group of each sample is used as an extra feature for a supervised regression. A hyperparameters study was carried out varying the number of clusters and finally, to determine which is the best configuration for the presented problem, three different unsupervised metrics were taken into account: Silhouette,

Calinski-Harabasz and Davies-Bouldin scores. In table 1 we can see the results achieved with the selected hyperparameter.

Clustering	Best number of clusters	Silhouette	Calinski-Harabasz	Davies-Bouldin
Gaussian Mixture	4	0.4450	32735.4139	0.7654
Spectral Clustering	3	0.4936	40391.5038	0.6354
Agglomerative Clustering	4	0.5279	41354.7560	0.6359
K-Means	4	0.5374	47787.0924	0.6338

Table 1. Best hyperparameter scoring using for the clustering techniques implemented In order to get a projected visualization of the data, a 2D mapping was done by training a LDA model using the cluster assigned to each sample as its class. In figures
5-8, we can see the 2D projection for all clustering techniques evaluated.

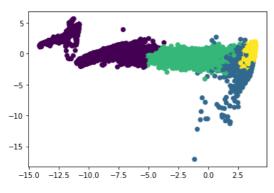


Figure 5. 2D representation of the dataset for Gaussian Mixture technique.

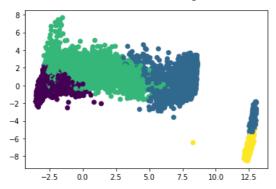


Figure 7. 2D representation of the dataset for Agglomerative clustering technique.

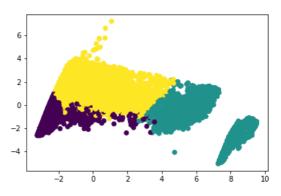


Figure 6. 2D representation of the dataset for Spectral clustering technique.

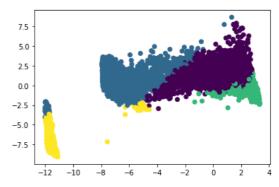


Figure 8. 2D representation of the dataset for K-Means clustering technique.

4.2. Regression

The main challenge of this work is to find the best clustering method for developing hybrid models, therefore it is necessary to complement the clustering metrics, due to this kind of metrics gives an idea about how the cluster processes is working, from unsupervised learning point of view. For this reason, in order to achieve efficient hybrid models applying clustering techniques, is essential to join them with a regression method. In this case MLP architecture is used.

An optimal behavior of MLP is based in the correct election of parameters for each cluster extracted by each clustering method applied. With this purpose, a Grid Search join to Cross-validation procedure has been designed in order to get the best parameters [1], with Mean Squared Error [32] like measure for defining the optimal model in the training process. The combination of parameters tested are showing following:

- Number of neurons in the hidden layer: from 12 to 30 neurons.
- Activation function: the hyperbolic tan function ("tanh") or the rectified linear unit function ("relu").
- Solver: optimizer in the family of quasi-Newton methods ("lbfgs"), stochastic gradient descent ("sgd") or stochastic gradient-based optimizer ("adam").

Final results show the four different approaches that have been implemented, based on the four clustering methods addressed previously. The tables 2, 3, 4 and 5 reflect each error measurement based on a weighted average, proportional for each error measure to the size of each grouping. The validation split is formed by the 20% of the total number of cases (5333).

Cluster	1	2	3	4	Weighted
					average
MSE	24.444	0.576	60.793	15.529	28.898
MAE	3.074	0.563	6.014	2.793	3.321
LMLS	1.386	0.201	2.468	1.334	1.423
MAPE	0.283	0.010	0.269	0.124	0.218
MASE	0.184	0.036	0.597	0.280	0.262
SMAPE	0.112	0.010	0.234	0.125	0.123

MAE 3.283 0.891 5.870 LMLS 1.474 0.373 2.4400.080 0.323 MAPE 0.025 MASE 0.277 0.046 0.743 0.079 0.261 SMAPE 0.025

1

27.356

Cluster

MSE

Table 2. MLP error for Gaussian Mixture clustering with 4 clusters

Table 3. MLP error for Spectral clustering with 3
clusters

2

2.358

3

57.282

Weighted

average

35.635

3.948

1.677

0.176

0.441

0.151

Cluster	1	2	3	4	Weighted	Cluster	1	2	3	4	Weighted
					average						average
MSE	56.655	1.191	60.793	11.128	27.447	MSE	5.2092	59.4698	21.2747	16.5433	36.7701
MAE	5.8624	0.699	2.897	2.578	3.331	MAE	0.8098	6.0651	2.9002	3.0794	4.0445
LMLS	2.437	0.279	1.317	1.295	1.446	LMLS	0.305	2.5043	1.321	1.492	1.704
MAPE	0.324	0.013	0.070	0.135	0.134	MAPE	0.014	0.326	0.069	0.136	0.187
MASE	0.692	0.049	0.252	0.479	0.346	MASE	0.056	0.752	0.277	0.4	0.470
SMAPE	0.254	0.013	0.070	0.125	0.113	SMAPE	0.014	0.263	0.068	0.137	0.155

Table 4. MLP error for Agglomerative clustering with 4 clusters

Table 5. MLP error for K-Means clustering with 4 clusters

Tables 6 shows the best parameters for each MLP model extracted from the list implemented on Grid Search procedure.

Figures 9, 10, 11 and 12 shows the graphical representation for each group where MLR regressor was applied, being the "Y" axis the output value, which refers to the output temperature of the lower solar panel situated in the output system. Each graphic displays the predicted output represented in red and the real output represented in blue. Only 100 elements from each data sample have been displayed for visualization purposes, when the size of the cluster is large enough, due to there are several clusters than contain minus than 100 data cases.

Gaussian clustering							
Grid Parameter / Cluster	1	2	3	4			
Number of neurons	25	23	30	30			
Activation function	tanh	tanh	tanh	tanh			
Solver	lbfgs	lbfgs	lbfgs	lbfgs			
Spectral clustering							
Grid Parameter / Cluster	1	2	3				
Number of neurons	27	27	27				
Activation function	tanh	tanh	tanh				
Solver	lbfgs	lbfgs	lbfgs				
Agglomerative clustering							
Grid Parameter / Cluster	1	2	3	4			

Number of neurons	30	30	29	18			
Activation function	tanh	tanh	tanh	tanh			
Solver	lbfgs	lbfgs	lbfgs	lbfgs			
K-Means clustering							
Grid Parameter / Cluster	1	2	3	4			
Number of neurons	21	27	24	25			
Activation function	tanh	tanh	tanh	tanh			
Solver	lbfgs	lbfgs	lbfgs	lbfgs			

Table 6. MLP best parameters for each clustering algorithm

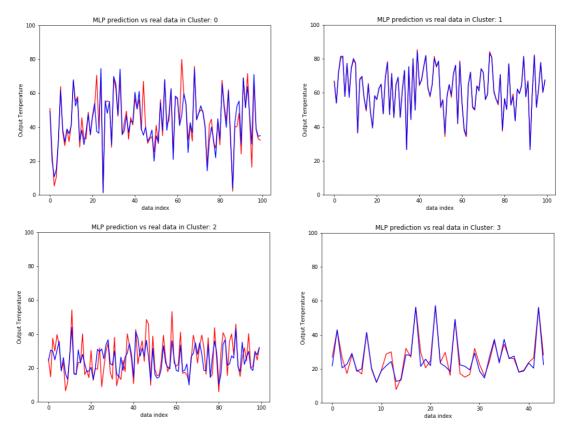


Figure 9. Real data vs. MLP predictions for Gaussian Mixture clustering

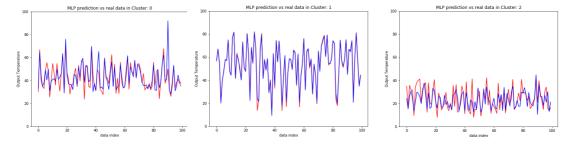


Figure 10. Real data vs. MLP predictions for Spectral clustering

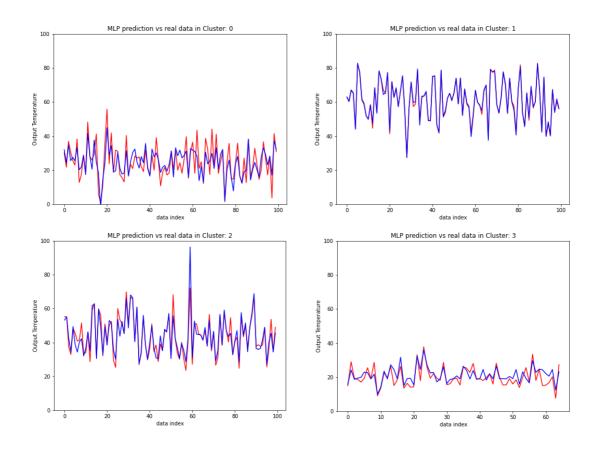


Figure 11. Real data vs. MLP predictions for Gaussian Mixture clustering

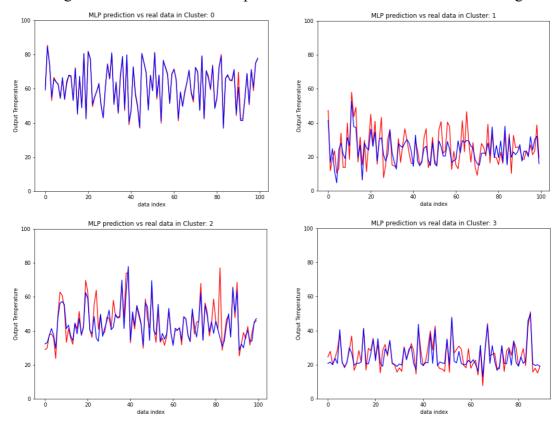


Figure 12. Real data vs. MLP predictions for Gaussian Mixture clustering

4.3 Discussion

Showing the unsupervised clustering results, it can be concluded that in all cases except Spectral Clustering the best number of clusters is 4. K-Means algorithm shows the best performance for the three evaluated metrics. Although all the results are quite similar, the worst results are achieved by Gaussian Mixture with the worst values for all the metrics. Agglomerative clustering and Spectral Clustering shown a similar clustering results with a slightly better values in first one taking into account that Spectral Clustering just groups the data in 3 clusters while Agglomerative Clustering groups it in 4.

As we can see in tables 2-5, the results achieved on the MLP show two different groups of performance. The better results are achieved with Gaussian Mixture and Agglomerative clustering with a similar errors in all the metrics evaluated. However, Agglomerative clustering improves Gaussian mixture in a 5% taking into account the Mean Squared Error. In the other point, both spectral clustering and K-Means show a lower performance than the other two clustering methods being outperformed by Agglomerative in more than a 25%.

Taking into account the results of the clustering with Shilouette, Calinski-Harabasz and Davies-Bouldin, agglomerative clustering has demonstrated to obtain good clustering power and a good regression power in combination with MLP. In contrast, K-Means achieved a great performance in clustering but showing high errors in regression.

5. Conclusions and future works

The paper address four possible clustering methods: Gaussian Mixture Clustering, Spectral Clustering, Agglomerative Clustering and K-means, in order to achieve the best one for implementing a robust hybrid MLP regression model of a thermal solar system. Based on the typical regression error metrics and specific clustering errors metrics such as Silhouette, Calinski-Harabasz and Davies-Bouldin, authors can conclude that the best method is Agglomerative Clustering, being the optimal number of cluster four.

While it is true that four groups could limit the effective operations of MLP, the combination of this technique with Agglomerative Clustering could even be better working with bigger datasets.

Future works will be oriented to apply other regression techniques such as Support Vector Machines, Extra Tree Regressor and Polynomial Regression. On the other hand, authors will work with new real datasets from bio-climatic field.

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