Clustering of Data Streams With Dynamic Gaussian Mixture Models: An IoT Application in Industrial Processes

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Abstract—In industrial Internet of Things applications with sensors sending dynamic process data at high speed, producing actionable insights at the right time is challenging. A key problem concerns processing a large amount of data, while the underlying dynamic phenomena related to the machine is possibly evolving over time due to factors, such as degradation. This makes any actionable model become obsolete and necessary to be updated. To cope with this problem, in this paper we propose a new unsupervised learning algorithm based on Gaussian mixture models called Gaussian-based dynamic probabilistic clustering (GDPC) mainly based on integrating and adapting three well known algorithms for use in dynamic scenarios: the expectation-maximization (EM) algorithm to estimate the model parameters and the Page-Hinkley test and Chernoff bound to detect concept drifts. Unlike other unsupervised methods, the model induced by the GDPC provides the membership probabilities of each instance to each cluster. This allows to determine, through a Brier score analysis, the robustness of the instance assignment and its evolution each time a concept drift is detected. Also, the algorithm works with very little data and significantly less computing power being able to decide whether (and when) to change the model. The algorithm is tested using synthetic data and data streams from an industrial testbed, where different operational states are automatically identified, giving good results in terms of classification accuracy, sensitivity, and specificity.

I. INTRODUCTION

NOWADAYS, Internet of Things (IoT) has opened a wide range of applications, where sensor data and other contextual data combined with computational models are able to produce actionable insights, which can be used as new control or monitoring systems and even new services [1]. Moreover, the application of these technologies to the industry, called industrial IoT (IIoT) [2] presents new opportunities to reduce the downtime of the machines and increase their availability, mainly channeled through predictive maintenance, where the failures need to be predicted as soon as possible [3].

Machine learning is highly used within this context. Algorithms are highly applicable to solve specific predictive maintenance problems, such as the remaining useful life (RUL) for a specific asset, failure diagnosis and prognosis, process optimization, testing, visual inspection, and quality control, among others. As a result, models that are able to predict quality deviations or failures during industrial processes can be obtained. Nevertheless, almost all of them are based on the analysis of datasets, where large and complex data storage and strategies are needed. Industrial problems need an actionable insight at the required time, e.g., to avoid further damage when a failure has occurred in a machine [4]. However, algorithm capabilities are limited by their own complexity and also by the computing power.

Additionally, this type of algorithms relies on the assumption that those models are stable throughout the industrial process (e.g., machining, painting, welding, etc.), basically meaning that the boundary conditions (i.e., degradation, raw material, tooling, temperature, etc.) of the specific problem are invariant over time. This assumption is not held in most industrial applications, where variations of boundary conditions are common across productive assets. Data heterogeneity makes real-time data streams analysis difficult to perform [5]. Therefore, we describe techniques that operate with adaptive window size as a possible solution for improving performance. In this case, data stream analysis is an important tool to detect these variations and the potential implied model in dynamic environments [6]. Machine learning algorithms can then be adapted to this setting.

In clustering, data stream analysis is still under development and limited to specific algorithm types: density- and distance-based [7], [8]. In this survey about data stream clustering, the most relevant algorithms are based on the following.

1) The k-means algorithm, e.g., BIRCH [9], CluStream [10], DGClust [11], StreamKM++ [12], scalable k-means [13], single-pass k-means [14], and SWClustering [15].
2) Hierarchical clustering, e.g., ODAC [16].
3) DBSCAN, e.g., D-Stream [17] and DenStream [18].
4) Hybrid algorithms, e.g., ClusTree [19], which is a hybrid of k-means and DBSCAN.

In all these approaches the underlying algorithm, i.e., k-means, hierarchical and DBSCAN, provides a crisp assignment of instances to clusters without more information about the data and how it is assigned to each cluster. However, Gaussian mixture model (GMM)-based clustering [20] adds the mixture model itself, the posterior probability that an instance has to be assigned to each component (an instance could belong to all components with different probabilities) or the possibility of evaluating the strength of the instance assignment to each component via the Brier score [21].

In the context of real engineering processes, such as machining operation data, [22] showed that GMMs had the ability to clearly discriminate what is happening inside a process much better that the k-means and hierarchical agglomerative algorithms which assume a static picture of the problem, where the variation of clusters (shape, number and position) over time was not taken into account. In this case, partition algorithms, such as k-means, tend to cluster based on underlying variables, which are not included in the analysis, such as angular speed, which do not leverage any new knowledge about the analyzed process. Moreover, agglomerative algorithms perform similarly.

From this point of view, data analysis performed with GMMs has a proven ability to leverage knowledge discovery about the engineering process, giving more information than others. However, the expressiveness of probability-based clustering algorithms is obscured by their inability, to the best of our knowledge, to deal with data streams. For example, GMMs are used as a simple method for dealing with data streams in the absence of concept drift [23]. Additionally, GMMs are usually applied to support other models, such as the evolving Bayesian network capable of dealing with data streams [24]. Nevertheless, the associated distribution parameters are estimated offline, running the EM algorithm on historical data.

A concept drift is defined in [25] as a change over time in the shift of relation between input data and the target variable. This change could be triggered by specific changes in behavior, such as degradation, extremely common in many industrial environments and helpful for early failure detection. It is also useful for detecting sensor failure and degradation [26]. As data streams and the process dynamics are an important consideration for the industry, the work in this paper is driven by the need for GMM-based probabilistic clustering, which is capable of dealing with the dynamic evolution and drifts of the industrial processes, providing a new analytics tool for IIoT.

As one of the objectives of this paper is to detect degradation of the process, a new GMM-based clustering algorithm adapted to dynamic environments called Gaussian-based dynamic probabilistic clustering (GDPC) is proposed. Based on a continuous stream of data, GDPC provides the probabilities of each instance being a member of each cluster. A measure of the cluster assignment robustness and a procedure to detect when data are not well represented by the model, i.e., there is a concept drift. Section II explains the steps of the GMM-based dynamic algorithm from training to component assignment. In Section III, a performance assessment with different parameters is carried out with experimentation to study their influence. In Section IV, a model for parameter selection for high performance is developed to provide the algorithm with real applicability. In Section V, the capacity of the algorithm to assign instances to each component during data stream monitoring is studied. This is in order to assess the ability to increase the adjustment level of the instance to each Gaussian mixture component by updating the Gaussian mixtures model. In Section VII, the algorithm is tested using real data from an industrial testbed and, in Section VIII, the conclusions and further work are given.

II. METHODOLOGY

Inspired by the general data stream algorithm pipeline, described by [10] and [27], the following steps are defined.

1) GMM offline training with an initial dataset \( D = \{x_1, \ldots, x_N\} \) of size \( N \), where each instance has \( d \) components, \( x_i = (x_{i1}, \ldots, x_{id}) \) (see Section II-A).
2) Fitting test of new incoming instances from the data stream to the current model trained in step 1 or 7, measured by the mean log-likelihood (see Section II-B).
3) Outlier detection (see Section II-C).
4) Concept drift detection with a cumulative value of outliers found, detect if there is a concept drift (see Section II-D).
5) If a concept drift is detected go to step 7, otherwise continue with step 6 for clustering.
6) If no concept drift is detected, assign the instance to each component of the GMM depending on its posterior probability (see Section II-E).
7) If concept drift is found, the model is readjusted using the last Chernoff bound window \( D' = \{x_1, \ldots, x_n\} \) of size \( n \) (see Section II-D).

The sequence of the algorithm steps is described in Fig. 1. A detailed description of these steps is given in the following sections.

A. GMM Offline Training

This step may be considered static since the algorithm must be trained with a data batch of size \( N \). The idea is to create an initial GMM from these data to be used as reference.

Assuming the density in the 4th cluster is given by \( f_k = (x; \theta_k) \) the GMM model is given by

\[
f(x; \Psi) = \sum_{k=1}^{K} \pi_k f_k(x; \theta_k)
\]

with parameters

\[
\Psi = (\pi_1, \ldots, \pi_K, \theta_1, \ldots, \theta_K)
\]

where \( \pi_k \) are the specific weights of each component \( k \) within the mixture, \( \theta_k = (\mu_k, \Sigma_k) \) with \( k = 1, \ldots, K \) and \( \mu_k, \Sigma_k \)
are the vector of means and the covariance matrix of each component \( k \), respectively.

The unknown parameter \( \Psi \) is estimated by maximum likelihood method. This estimation can be approached using the EM algorithm [20], [28]. The EM algorithm is an iterative method that has two main steps: 1) expectation or E-step, where a function for the expectation of the log-likelihood is created and 2) maximization or M-step, where the parameters are computed maximizing the expected log-likelihood. These steps are summarized below.

1. **E-Step:** Having the likelihood for \( \Psi \) defined by

\[
L(\Psi) = \prod_{i=1}^{N} \left( \sum_{k=1}^{K} \pi_k f_k(x_i; \theta_k) \right)
\]

then, the log-likelihood is

\[
\log L(\Psi) = \sum_{i=1}^{N} \log \left( \sum_{k=1}^{K} \pi_k f_k(x_i; \theta_k) \right) = \sum_{i=1}^{N} \sum_{k=1}^{K} z_{ki} \{ \log \pi_k + \log f_k(x_i; \theta_k) \}
\]

where \( z_{ki} \) contains 0 or 1 depending on the assignment of each input instance \( i \) to each component. Its corresponding random variable \( Z_{ki} \) is defined as

\[
Z_{ki} = \begin{cases} 
1, & \text{if } x_i \text{ is assigned to component } k \\
0, & \text{otherwise.}
\end{cases}
\]

The value of \( Z_{ki} \) is estimated by the E-Step at each iteration.

The subsequent iterations for E-Step only require finding the posterior probability that the \( i \)-th instance is assigned to the \( k \)-th component, i.e., \( \tau_k^{(t)}(x_i; \Psi^{(t)}) \).

Therefore,

\[
E_{\Psi^{(t)}}[Z_{ki} | \mathcal{D}] = P_{\Psi^{(t)}}[Z_{ki} = 1 | \mathcal{D}] = \tau_k^{(t)}(x_i; \Psi^{(t)})
\]

where

\[
\tau_k^{(t)}(x_i; \Psi^{(t)}) = \frac{\pi_k f_k(x_i; \theta_k^{(t)})}{f(x_i; \Psi^{(t)})}.
\]

For a GMM, (3) is defined as

\[
\tau_k^{(t)}(x_i; \Psi^{(t)}) = \frac{\pi_k^{(t)} N(x_i | \mu_k^{(t)}, \Sigma_k^{(t)})}{\sum_{j=1}^{K} \pi_j^{(t)} N(x_i | \mu_j^{(t)}, \Sigma_j^{(t)})}.
\]

Consequently, using (2) and (3) it is possible to define the conditional expectation of \( \log L(\Psi) \) given \( \mathcal{D} \) as

\[
E_{\Psi^{(t)}}[\log L(\Psi) | \mathcal{D}] = Q(\Psi; \Psi^{(t)}) = \sum_{k=1}^{K} \sum_{i=1}^{N} \tau_k^{(t)}(x_i; \Psi^{(t)}) \times \{ \log \pi_k + \log f_k(x_i; \theta_k) \}.
\]

Once the expectation is found at iteration \( t \), the M-Step is applied.

2. **M-Step:** The \( t + 1 \) iteration for the M-Step searches for the maximization of \( Q(\Psi; \Psi^{(t)}) \) with respect to \( \Psi \) in order to update the value of \( \Psi^{(t+1)} \). The estimated specific weights, \( \pi_k^{(t+1)} \), are found independently of the vector of unknown parameters \( \theta^{(t+1)} \) by means of

\[
\pi_k^{(t+1)} = \frac{1}{N} \sum_{i=1}^{N} \tau_k^{(t)}(x_i; \Psi^{(t)}), \quad k = 1, \ldots, K.
\]

To obtain \( \theta^{(t+1)} \) we solve

\[
\sum_{k=1}^{K} \sum_{i=1}^{N} \tau_k^{(t)}(x_i; \Psi^{(t)}) \frac{\partial \log f_k(x_i; \theta_k)}{\partial \theta_k} = 0.
\]

As the likelihood increases monotonically with the iterations of the EM algorithm, a standard stopping criterion is to cease when

\[
L(\Psi^{(t+1)}) - L(\Psi^{(t)}) \quad \text{is small enough to indicate the convergence of the sequence of parameters } \{\Psi^{(t)}\}.
\]

For GMM

\[
f_k(x_i; \theta_k) = f_k(x_i; \mu_k, \Sigma_k) = (2\pi)^{-\frac{d}{2}} |\Sigma_k|^{-\frac{1}{2}} \exp \left\{ -\frac{1}{2} (x_i - \mu_k)^T \Sigma_k^{-1} (x_i - \mu_k) \right\}
\]

where parameters \( \mu_k^{(t+1)} \) and \( \Sigma_k^{(t+1)} \) \((k = 1, \ldots, K)\) are estimated using (4)

\[
\mu_k^{(t+1)} = \frac{\sum_{i=1}^{N} \tau_k^{(t+1)}(x_i)}{\sum_{i=1}^{N} \tau_k^{(t+1)}(x_i; \Psi_k^{(t+1)})}, \quad k = 1, \ldots, K
\]
and
\[
\Sigma_k^{(i+1)} = \frac{\sum_{i=1}^{N} r_k^{(i+1)} (x_i; \Psi_k^{(i)}) (x_i - \mu_k^{(i+1)}) (x_i - \mu_k^{(i+1)})^T}{\sum_{i=1}^{N} r_k^{(i+1)} (x_i; \Psi_k^{(i)})},
\]
where \( k = 1, \ldots, K \).

Having the reference model (induced from initial dataset \( \mathcal{D} \)), a fitting test is done with new instances coming from the data stream.

**B. Fitting Test**

The fitting test checks if the new incoming data stream instance is well represented by the current GMM. To measure this, we use the mean log-likelihood criterion [29], which is calculated from (2) and (5)
\[
\overline{\log L}(\Psi) = \frac{1}{N} \sum_{i=1}^{N} \log \left( \sum_{k=1}^{K} \pi_k f_k (x_i; \mu_k, \Sigma_k) \right)
\]
and compare it with the log-likelihood \( \log L_n(\Psi) \) of the new instance \( x_n \). A significant small value of \( \log L_n(\Psi) \) in comparison with \( \overline{\log L}(\Psi) \) indicates that the new data is badly represented by the current mixture model. Next, the values of \( \log L_i(\Psi) \) are used to detect outliers (see the following section).

**C. Outlier Detection**

Once the fitting test result for the data stream is found, i.e., the value of (2), it is necessary to know the value of \( \log L_i(\Psi) \) that shows if the instance \( x_i \) is correctly represented by the model or not. When an instance is not represented by the current GMM, it is labeled as an outlier, and it is used afterward in the concept drift detection step (Section II-D).

Sebastiao and Gama [27] and Borchani et al. [30] successfully applied the Page-Hinkley test [31], sequential analysis of the data stream, taking the sample variance of the analyzed data as the key measure. As explained by [32], the test associates the sample variance with the normal behavior of the data stream. Therefore, any disturbance, such as degradation, in this variance is associated with an abrupt change in the data stream behavior. In this case, the null hypothesis of the test is that the instance \( x_i \) is correctly represented by the GMM.

To perform the Page-Hinkley test, as new instances arrive from the data stream process, \( \log L_i(\Psi) \) is calculated from (6) and stored on a vector defined by
\[
\log L_i(\Psi) = \{ \log L_1(\Psi), \ldots, \log L_n(\Psi) \}
\]
where \( n \) is explained by [33]–[35], as an adaptive window size bounded by
\[
n \leq \frac{3(1 + \epsilon)}{(1 - \epsilon) e^2} \ln \left( \frac{2}{\delta} \right)
\]
and the size \( n \) changes inversely to \( p \), meaning that if a new instance is well represented (large \( p \)) or badly represented (small \( p \)), the window size is small or large, respectively. This adaptive window helps to optimize the amount of data stored, i.e., if a new data instance fits the model, there is no need to have a large amount of historical data.

Next, the accumulated difference between mean log-likelihood and the accumulated mean value (\( \text{CUM}_j \)) is defined by
\[
\text{CUM}_n = \frac{1}{n} \sum_{j=1}^{n} (\log L_j(\Psi) - \text{mean}_j) - \delta
\]
where \( \text{mean}_j = (1/j) \sum_{i=1}^{j} \log L_i(\Psi) \) is the mean value of the first \( j \) components of the vector defined in (7), and \( \delta \) is a positive random parameter of tolerance for the maximum magnitude of accepted changes. The way to select this value is described in Section III. Additionally, a value \( \text{MAX}_j \) is defined as
\[
\text{MAX}_j = \max(\text{CUM}_j, j = 1, \ldots, n).
\]

Therefore, threshold \( \text{PH}_j \) is defined as \( \text{PH}_j = \text{MAX}_j - \text{CUM}_j \) \((j = 1, \ldots, n)\), and so
\[
\text{reject}_j = \begin{cases} 
1, & \text{if } \text{PH}_j > \lambda \\
0, & \text{otherwise}
\end{cases}
\]
where \( \lambda > 0 \) is the value that controls the null hypothesis, where 1 means that instance \( x_i \) is labeled as an outlier (the hypothesis is rejected). Therefore, a vector \( \text{reject}_j \), of size \( n \) stores 1 or 0 if instance \( x_j \) is an outlier or not, respectively.

**D. Concept Drift Detection**

The aim of this step is to differentiate between outliers produced by noise in the data stream (data acquisition system uncertainty) and those related to the GMM, where a group of instances, whose size is larger than a defined threshold, no longer fit properly. The latter case is defined as concept drift.

In order to detect a concept drift, we adapt the approach proposed by [35] to apply Chernoff bounds to define the minimum number of instances \( s \) that are not outliers in a defined window, according to
\[
s = \frac{3(1 + \epsilon)}{e^2} \ln \left( \frac{2}{\phi} \right)
\]
and the size of adaptive data stream windows from (8) to control the amount of stored instances for training if a concept drift is detected. The main idea of this approach is to determine the total amount of outliers needed to indicate that the GMM has expired and a new one should be learned from data. Now, if \( r = \sum_{i=1}^{n} (1 - \text{reject}_i) < s \), a concept drift is detected.

As soon as this model change is detected, a new GMM is estimated from the adaptive window data using the algorithm described in Section II-A, with \( N \) defined as the lowest \( n \), \( n \) being defined by (8). \( r \) is initialized as 0 and increases its value each time an instance is correctly represented by the GMM. \( p \) is \( r/n \) and the initial value for \( p = 0 \) and it is updated each cycle.
E. Instance Assignment

The next step in the GDPC algorithm is to assign an instance to each component via the posterior probability that instance $x_i$ belongs to component $k$ of the mixture, as reflected in (3). That is

$$z_{kj} = \begin{cases} 1, & \text{if } k = \arg \max_h \tau_h(x_j; \Psi) \\ 0, & \text{otherwise.} \end{cases}$$

F. Algorithm Wrap-Up

It is now necessary to wrap-up the isolated modules described in Sections II-A–II-E to provide a clear view of their operation and interactions that make up the GDPC algorithm backbone.

Once the modules are bonded together, the algorithm is ready to receive a data stream coming from a specific process and initialize the parameters related with each module: $\delta$, $\lambda$, $\epsilon$, $\phi$, and $N$. These parameters are selected following the approach presented in Section IV. Using these values, the minimum allowed window size, $s$, is calculated using (10). Other parameters, such as the probability, $p$, and the number of instances without outliers, $r$, are initialized to zero.

After initialization, a training data set $\mathcal{D}$ is filled with data stream until the number of instances is equal to the parameter $N$. Then, the initial GMM can be trained with $\mathcal{D}$ using the EM algorithm to estimate the Gaussian mixture parameters. From this step, the algorithm starts to analyze data stream, processing one instance at a time using the GMM as the statistical summary that represents the analyzed data.

The data stream analysis start by finding the adaptive window of size $n$ using Chernoff bound that works as a temporary buffer. It stores the newest $n$ instances incoming from the data stream. Meanwhile, each data stream instance is checked against the current GMM using the mean log-likelihood. This mean value is monitored by the Page-Hinkley test that searches for outliers. If this hypothesis test is not rejected, the instance is correctly represented by the GMM and it is recorded by updating $r$.

The next mainly compares $r$ against $s$. If the number of correctly represented instances $r$ is larger than $s$, a concept drift turning point in the data stream is detected. If $r$ remains smaller than $s$ (i.e., no concept drift) the probability, $p$, is updated with $r/n$ and the instance is assigned to a GMM component using its posterior probability and continues with the next instance from the data stream.

The complete process is summarized in Algorithm 1.

### Algorithm 1 GDPC

<table>
<thead>
<tr>
<th>Requirement</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>$\mathcal{D}$</td>
<td>Data stream</td>
</tr>
<tr>
<td>$N$</td>
<td>Number of initial training instances</td>
</tr>
<tr>
<td>$\delta$ and $\lambda$</td>
<td>Positive threshold that defines the value from which the null hypothesis of an instance $x_i$ to be correctly represented by the GMM is rejected.</td>
</tr>
<tr>
<td>$\epsilon$ and $\phi$</td>
<td>Positive tolerance to changes that are to be absorbed.</td>
</tr>
</tbody>
</table>

1) GMM offline training (Section II-A).
   a) $N$: Number of initial training instances.
2) Page-Hinkley hypothesis test (Section II-C).
   a) $\lambda$: Positive threshold that defines the value from which the null hypothesis of an instance $x_i$ to be correctly represented by the GMM is rejected.
   b) $\delta$: Positive tolerance to changes that are to be absorbed.
3) Chernoff bound (Section II-D).
   a) $\epsilon$: Additive error bound ($0 < \epsilon < 1$).
   b) $\phi$: Constant to control the probability of an instance to be successfully clustered ($0 < \phi < 2$).

A. Experimental Datasets

To study the parameter effect on the concept drift detection capability, synthetic datasets are created by simulating random values from a mixture of Gaussian distributions. As the principal objective of these experiments is to calibrate the GDPC's concept drift detection capability, it is critical to know the concept drift locations within the data stream beforehand. This outputs a reference data stream, with a predefined variety...
of concept drifts, useful for comparing the algorithm results in terms of concept drift detection and its detection delay.

Therefore, data streams are created concatenating different subsets of instances each of which is simulated from GMMs with different parameter value combinations. Due to the variation of the parameters, different GMMs along the dataset simulate the concept drifts.

In the synthetic model with \( K = 3 \) components for this particular scenario

\[
\theta = (\pi_1, \pi_2, \pi_3, \mu_1, \mu_2, \mu_3, \Sigma_1, \Sigma_2, \Sigma_3)
\]

where \( \mu_k \) (\( k = \{1, 2, 3\} \)) is the parameter vector of mean values and \( \Sigma_k \) (\( k = \{1, 2, 3\} \)) is the parameter matrix of variances.

Different values of offsets and Gaussian noise, selected to artificially produce different models, are applied equally to all the parameter dimensions: offset = ±5%, ±10%, and ±25%. A random variable is defined as noise, where \( \sigma = \{0.05, 0.10, 0.25\} \) when its random value is simulated with Gaussian distributions: \( N(0, 0.05) \), \( N(0, 0.10) \), and \( N(0, 0.25) \), respectively. Parameter magnitudes are changed in order to study detection capabilities depending on the similarity of the models, i.e., offset = ±5% or \( N(0, 0.05) \) means that the drift is more similar to the original model than offset = ±25% or \( N(0, 0.25) \).

Once the parameters of the mixtures are varied, for example: using \( \Sigma_i = \text{offset} \cdot \Sigma \) instances are simulated producing a subset of length \( \text{offset} \). Accordingly, an original subset (without variations) plus nine subsets of instances with different lengths and variations of their parameters are generated. The description of the way parameters are modified for obtaining different subsets is shown in Table I.

The concatenation of the different subsets of instances described in Table I enables different data streams to be obtained for experimentation. In this way, there are three data streams created from these concatenations shown in Table II. Data streams have positive and negative offsets and Gaussian noise depending on the magnitude defined in Table I. These data streams have an equal number of instances, that is, \( l_{\text{datastream}} = 37,000 \) with three additional concept drifts (CDset).

### B. Design of Experiments

To efficiently explore the parameter effect on model behavior, we require a fractional factorial and orthogonal design in which factors (variables for analysis) and levels (value ranges of each factor) are balanced. This type of designs are called Taguchi methods. The entry key input to select the Taguchi design is the required number of parameters. In this case, we studied five GDPC parameters: \( \lambda, b, c, \phi, \) and \( N \); each with four levels. Therefore, the design that best fits this number of parameters and levels is the orthogonal Taguchi L16 (type B) design [36] with five four-level parameters, i.e., a 4\(^5\) design with 16 observations. Table III shows the parameters and levels to be used in the experimentation.

To study the influence of these parameters, we determine the following measures to be extracted from the experiment.

1. **Detected Concept Drifts (CDdet):** The amount of concept drifts detected within a data stream.
2. **Instance Marked As Concept Drift Turning Point (nCD):** The first instance, where the algorithm detects a concept drift CDdet, defined as a turning point.
3. **True Concept Drifts (CDtrue):** From CDdet, the amount of true concept drifts measured by their location \( n_{\text{CD}} \) compared to CDset in Table II.
4. **Processing Time in Seconds (t):** The time required by the algorithm to process the data stream.

From observations, performance is assessed using the figures of merit derived from the concept drift detection confusion matrix, i.e., accuracy, sensitivity, and specificity. The confusion matrix is detailed in Table IV, where a comparison between a true concept drift class \( c_i \) and concept drift detected by the model class \( c_m \) can be made.

1. **\( c_i \) Is Positive and \( c_m \) Is Positive:** The instances detected by the model are true concept drifts, i.e., CDtrue.
2. **\( c_i \) Is Negative and \( c_m \) Is Positive:** Not all the instances detected by the model are true concept drifts, i.e., CDdet − CDset.
3. **\( c_i \) Is Positive and \( c_m \) Is Negative:** Not all the instances with true concept drifts are detected, i.e., CDset − CDtrue.
4. **\( c_i \) Is Negative and \( c_m \) Is Negative:** The amount of instances not detected as concept drift, i.e., \( N − \text{CDdet} \).
From Table IV, the figures of merit are defined as follows.
1) Accuracy: \([\frac{(CD_{\text{true}} - CD_{\text{det}} + N)}{N}]\).
2) Sensitivity—True Positives (TP): \((CD_{\text{true}}/CD_{\text{det}})\).
3) Specificity—True Negatives: \([N - CD_{\text{det}}]/(N - CD_{\text{set}})]\).
4) Recall: \([TP/(TP + FN)]\).
5) F-Score: \([2TP/(2TP + FP)]\).

C. Experimental Results

Three experimental repetitions are run with each data stream defined in Table II. The results of these experiments are shown in Tables V–VII.

In Table V, the results with data stream 1 are shown. In this case, the algorithm is able to detect the first concept drift with delays not larger than 591 instances for experiment 14 (turning point detected in instance 15591) in the complete set of experiments. Experiments 1 and 12, where the only experiments without a correct true concept drift (CD_{true}) detection, with only one and two detections, respectively. On the other hand, experiments 6–11 and 13–16 achieved 100% accuracy without any FPs. Therefore, 87.5% of the experiments have parameter combinations that are able to detect true concept drifts (100% sensitivity), but only 62.5% of them with 100% accuracy.

Table VI shows the results for data stream 2, and the situation is slightly different: 62.5% of the experiments have 100% sensitivity and only 50% of them have 100% accuracy (experiments 6, 9, 10, 12, and 14), meaning that fewer combinations of parameters are able to detect the predefined concept drifts. These results show that the algorithm is sensitive to the difference between GMM, simulated by the offset and noise changes (concept drifts) are small.

In the experiment with data stream 3 (Table VII), the results are more critical because some parameter combinations are not able to detect any concept drift, i.e., experiments 13–16. In this case, 56% of the experiments have 100% sensitivity with only 55% of them with 100% accuracy, i.e., experiments 6, 8–10, and 12. For this data stream, the experiments that could not detect any concept drift are differentiated by the parameter \(\lambda\), which has the largest level (100). As a preliminary analysis for parameter selection, large levels of \(\lambda\) are not optimal when GMM changes (concept drifts) are small.

Regarding processing time, experiments 1–4 with \(\lambda = 0.1\) from any data stream have the highest value, not always guaranteeing the concept drift detection. As a preliminary insight, \(\lambda > 1\) is able to give similar results with 50% of processing time. Other experiments need around 7.21 s to process 37 000 instances independently from the data stream used.

In summary, parameter combinations 9, 10, and 12 are the only combinations able to detect concept drifts in all data streams, meaning that it is possible to use optimal parameters in the proposed algorithm. It is interesting to note that these combinations have the same \(\lambda = 10\), giving some clues to understand the optimal magnitude for this parameter.

However, these are intuitive results that can help to understand the parameter influence in the algorithm performance. As a result, in Section IV a complete study, using the experimental results to build a supervised model to select parameters to improve the figure of merit, is presented.

IV. PARAMETER SELECTION FOR HIGH PERFORMANCE

From the experimentation results shown in the previous section, optimal parameter selection is studied to find the underlying common characteristics between parameter combinations 9, 10, and 12. Therefore, data from experiments is fed into traditional models for experiment result analysis, such as linear or polynomial regressions or the signal-to-noise ratio test. However, those models have not given the desired result in relation to their accuracy due to a low coefficient of determination.

One of the principal issues that could affect these underperforming models is the lack of enough data. Consequently, taking advantage of algorithm efficiency and its ability to obtain good performance with small amounts of data, naïve Bayes classifier [37] is used. This classifier is built upon the assumption of conditional independence of the predictive variables given the class, where the maximum posterior probability assignment of the class variable is calculated as prediction.

Three classifiers are designed for each figure of merit: accuracy, sensitivity, and specificity, taken as the class variables. The predictor variables are the five univariate Gaussian parameters \(\lambda, \delta, \epsilon, \phi,\) and \(N\).

Because class variables are continuous, the values in each experiment are categorized using three different groups.
1) Low, for values smaller than 50%.
2) Average, for values that are greater than 50% and less than 99%.
3) High, for values greater than 99%.

Therefore, the naïve Bayes model is defined by

\[
c^* = \arg \max_c p(c|\text{par}) = \arg \max_c p(c) \prod_{i=1}^{5} f(\text{par}_i|c)
\]

where \(c^*\) is the Bayesian classifier selected as the most probable hypothesis with the maximum a priori or MAP and \(\text{par} = (\lambda, \delta, \epsilon, \phi, N)\), and \(c\) is the class variable of each figure of merit previously described.

A. Parameter Ranking by Relevance

The first step in the experimental results analysis with the naïve Bayes classifier is to rank the relevance of each parameter for each figure of merit. Experimental data from Section III-C is fed into the model and the posterior probabilities for high value of class variable are shown.

<table>
<thead>
<tr>
<th>Predicted</th>
<th>Truth</th>
<th>CD_{true} - CD_{det}</th>
<th>CD_{set} - CD_{true}</th>
<th>N - CD_{set}</th>
</tr>
</thead>
<tbody>
<tr>
<td>+</td>
<td>+</td>
<td>CD_{true}</td>
<td>CD_{det} - CD_{true}</td>
<td>CD_{set} - CD_{true} + N - CD_{set}</td>
</tr>
<tr>
<td>-</td>
<td>-</td>
<td>CD_{true} - CD_{true}+ N - CD_{true}</td>
<td>CD_{true} + CD_{true} - N - CD_{true}</td>
<td>CD_{true} + CD_{true} - N - CD_{true}</td>
</tr>
</tbody>
</table>

| TABLE IV | CONFUSION MATRIX |

<table>
<thead>
<tr>
<th></th>
<th>+</th>
<th>CD_{true} - CD_{det}</th>
<th>CD_{det} - CD_{true}</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Predicted</strong></td>
<td><strong>Truth</strong></td>
<td>CD_{true} - CD_{det} + N - CD_{det}</td>
<td>CD_{true} + CD_{true} - N - CD_{true}</td>
</tr>
</tbody>
</table>

TABLE IV

CONFUSION MATRIX
in Table VIII. The independent probability for each class to take high, average, and low values is shown, along with the prediction accuracy as naïve Bayes validation in Table IX.

<table>
<thead>
<tr>
<th>Test</th>
<th>(\lambda)</th>
<th>(\delta)</th>
<th>(\epsilon)</th>
<th>(\phi)</th>
<th>(N_{\text{train}})</th>
<th>(t)</th>
<th>(C_{\text{DET}})</th>
<th>(C_{\text{TRUE}})</th>
<th>(n_{\text{CD}})</th>
<th>Accu.</th>
<th>Sens.</th>
<th>Spec.</th>
<th>Recall</th>
<th>F-score</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>0.1</td>
<td>0.01</td>
<td>0.10</td>
<td>0.01</td>
<td>10</td>
<td>15.00</td>
<td>1.00</td>
<td>1.00</td>
<td>1.00</td>
<td>0.899</td>
<td>0.333</td>
<td>1.000</td>
<td>1.000</td>
<td>0.500</td>
</tr>
<tr>
<td>2</td>
<td>0.1</td>
<td>0.10</td>
<td>0.25</td>
<td>0.27</td>
<td>10</td>
<td>18.72</td>
<td>230</td>
<td>3</td>
<td>32100</td>
<td>0.994</td>
<td>1.000</td>
<td>0.994</td>
<td>0.994</td>
<td>0.997</td>
</tr>
<tr>
<td>3</td>
<td>0.1</td>
<td>0.75</td>
<td>0.54</td>
<td>100</td>
<td>3</td>
<td>18.45</td>
<td>717</td>
<td>3</td>
<td>32021</td>
<td>0.981</td>
<td>1.000</td>
<td>0.981</td>
<td>0.981</td>
<td>0.981</td>
</tr>
<tr>
<td>4</td>
<td>0.1</td>
<td>0.90</td>
<td>0.80</td>
<td>10000</td>
<td>16.40</td>
<td>320</td>
<td>3</td>
<td>32043</td>
<td>0.992</td>
<td>1.000</td>
<td>0.991</td>
<td>0.991</td>
<td>0.995</td>
<td></td>
</tr>
<tr>
<td>5</td>
<td>0.1</td>
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<td>0.01</td>
<td>100</td>
<td>9.66</td>
<td>35</td>
<td>3</td>
<td>32145</td>
<td>0.999</td>
<td>1.000</td>
<td>0.999</td>
<td>0.999</td>
<td>0.999</td>
<td></td>
</tr>
<tr>
<td>6</td>
<td>1.0</td>
<td>0.10</td>
<td>0.10</td>
<td>0.80</td>
<td>10</td>
<td>8.23</td>
<td>4</td>
<td>3</td>
<td>32155</td>
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<tr>
<td>7</td>
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<td>0.90</td>
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<td>3</td>
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<td></td>
</tr>
<tr>
<td>8</td>
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<td>0.75</td>
<td>0.27</td>
<td>10</td>
<td>8.28</td>
<td>12</td>
<td>3</td>
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<td>1.000</td>
</tr>
<tr>
<td>9</td>
<td>1.0</td>
<td>0.01</td>
<td>0.75</td>
<td>0.80</td>
<td>100</td>
<td>8.17</td>
<td>3</td>
<td>3</td>
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<tr>
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<td>3</td>
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<td>1.000</td>
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<tr>
<td>11</td>
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<td>0.10</td>
<td>0.27</td>
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<td>9.36</td>
<td>3</td>
<td>3</td>
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</tr>
<tr>
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<td>0.10</td>
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<td>8.18</td>
<td>2</td>
<td>2</td>
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<td>1.000</td>
</tr>
<tr>
<td>13</td>
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<td>0.01</td>
<td>0.90</td>
<td>0.27</td>
<td>100</td>
<td>7.91</td>
<td>3</td>
<td>3</td>
<td>32370</td>
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<td>1.000</td>
</tr>
<tr>
<td>14</td>
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<td>0.75</td>
<td>0.01</td>
<td>100</td>
<td>9.66</td>
<td>3</td>
<td>3</td>
<td>32544</td>
<td>1.000</td>
<td>1.000</td>
<td>1.000</td>
<td>1.000</td>
<td>1.000</td>
</tr>
<tr>
<td>15</td>
<td>1.0</td>
<td>0.01</td>
<td>0.25</td>
<td>0.80</td>
<td>10</td>
<td>9.29</td>
<td>3</td>
<td>3</td>
<td>32057</td>
<td>1.000</td>
<td>1.000</td>
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<td>1.000</td>
</tr>
<tr>
<td>16</td>
<td>1.0</td>
<td>0.00</td>
<td>0.10</td>
<td>0.54</td>
<td>100</td>
<td>9.97</td>
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<td>3</td>
<td>32260</td>
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<td>1.000</td>
<td>1.000</td>
<td>1.000</td>
<td>1.000</td>
</tr>
</tbody>
</table>

From Table VIII, the most relevant parameter, with the maximum posterior probability for high figures of merit (accuracy, sensitivity, and specificity) is naïve Bayes validation (Section II-C) parameters \(\delta\) and \(\lambda\) are the second most relevant
and Chernoff bound (Section II-D) parameters \( \phi \) and \( \epsilon \) are the least relevant. Analyzing these results, there is a sharp dependency of the algorithm on the size of the training window to be able to increase the figures of merit. However, parameter \( N \) is the most relevant in terms of training time, as the algorithm has to wait until the window is filled to continue with the data stream analysis. Additionally, large amounts of \( N \) could cause over-fitting, where concept drifts could be masked during the training.

These results are important to conclude that previous knowledge about the analyzed processes is needed to calibrate the existence of concept drifts. For example, if the RUL is expected to be 8 and 2000 h for a tool or a motor, respectively, the size of \( N \) should be lower for the tool.

With a clear understanding of each parameter role within the overall dynamic clustering algorithm given by experimentation and posterior analysis with naïve Bayes, mean values and standard deviation for each parameter are given in terms of the figure of merit that is optimized. Therefore, we estimate the mean and the standard deviation of the Gaussian distribution of each parameter for each figure of merit at its best value, i.e., the mean and the standard deviation of the distribution of any parameter given that the figure of merit takes its high value. A summary of the parameter values for high accuracy, sensitivity, and specificity is given in Table X.

From Section III-C, where the most accurate experiments (9, 10, and 12) were found, a comparison with the optimal parameters described in Table X can be made. Thus, almost all parameters from those experiments in Section III-C are within the values shown in Table X, meaning that it is possible to optimally tune the algorithm and control the initial training size.

To check the performance of the model built with the optimal parameters, an experiment is designed selecting the mean values for each parameter (Table X). Then, the parameters are introduced into the naïve Bayes model to check the classification process in terms of low, average, or high categories for each figure of merit with its corresponding posterior probability.

The experiment results are shown in Table XI, where the selected parameters are able to predict the highest possible values related to the corresponding figure of merit. In fact, as shown in the table, the posterior probability for the classification is 1.

Next, the parameters are checked using predefined data streams 1–3. Results are shown in Tables XII–XIV for data streams 1 and 2, 100% for accuracy, sensitivity, and specificity is obtained. It is important to stress that the largest detection delay is 135, meaning that it only needs 135 instances to detect the concept drift.

However, for data stream 3 (Table XIV), the parameters for high sensitivity have problems to detect the last concept drift located at instance 32 000. This last concept drift is related to a normal distribution noise with standard deviation 0.05 (Table I), explaining the difficulties for the algorithm to detect those small changes with these parameters.

As a conclusion of this section, it can be seen that algorithm parameters are sensitive to the analyzed scenario, where a tuning is needed the first time it is used. The idea is to select parameters, where concept drifts are detected. Therefore, as a procedure to fine-tune the algorithm parameters depending on the application, a Taguchi L16 design is used to analyze the response and to build a model. In this case, a naïve Bayes is used to select the best parameters for high accuracy, sensitivity, and/or specificity.

Another aspect is the ability of the algorithm to assign the instances to each component of the mixture model (cluster), which is the ultimate goal. The goodness of this assignment process is analyzed in the following section.

### V. Model Adjustment Measurement

The clustering step in the GDPC assigns an instance to each component via the posterior probability that the instance \( x_i \) belongs to component \( k \) of the mixture, as reflected in (3).

With the aim of studying how the GDPC performs with data streams containing concept drifts during the cluster assignment, we measure the posterior probability that instance \( x_i \) belongs to the evolution of component \( k \) during concept drift detection. Therefore, it is important to monitor how the model assigns instances to each component and how it begins to mismatch due to a concept drift.

Intuitively, one expects to find a better adjustment once the concept drift is detected and the model is recalculated, i.e., the posterior probabilities that an instance belongs to each

### Table VIII

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Accuracy</th>
<th>Sensitivity</th>
<th>Specificity</th>
</tr>
</thead>
<tbody>
<tr>
<td>( \phi )</td>
<td>0.0002</td>
<td>0.0002</td>
<td>0.0001</td>
</tr>
<tr>
<td>( \epsilon )</td>
<td>0.0002</td>
<td>0.0002</td>
<td>0.0002</td>
</tr>
<tr>
<td>( \delta )</td>
<td>0.0013</td>
<td>0.0013</td>
<td>0.0011</td>
</tr>
<tr>
<td>( \lambda )</td>
<td>0.0079</td>
<td>0.0076</td>
<td>0.0101</td>
</tr>
<tr>
<td>( N )</td>
<td>0.9904</td>
<td>0.9907</td>
<td>0.9885</td>
</tr>
</tbody>
</table>

### Table IX

<table>
<thead>
<tr>
<th>Figure of Merit</th>
<th>High</th>
<th>Average</th>
<th>Low</th>
<th>Prediction Accuracy</th>
</tr>
</thead>
<tbody>
<tr>
<td>Accuracy</td>
<td>0.63</td>
<td>0.25</td>
<td>0.12</td>
<td>58.3%</td>
</tr>
<tr>
<td>Sensitivity</td>
<td>0.69</td>
<td>0.10</td>
<td>0.22</td>
<td>68.8%</td>
</tr>
<tr>
<td>Specificity</td>
<td>0.92</td>
<td>0.08</td>
<td>0.00</td>
<td>93.8%</td>
</tr>
</tbody>
</table>

### Table X

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Accuracy</th>
<th>Sensitivity</th>
<th>Specificity</th>
</tr>
</thead>
<tbody>
<tr>
<td>( \lambda )</td>
<td>22.6 ± 38.7</td>
<td>20.6 ± 37.5</td>
<td>29.6 ± 43.0</td>
</tr>
<tr>
<td>( \delta )</td>
<td>3.65 ± 4.79</td>
<td>3.41 ± 4.63</td>
<td>2.89 ± 4.34</td>
</tr>
<tr>
<td>( \epsilon )</td>
<td>0.52 ± 0.32</td>
<td>0.54 ± 0.31</td>
<td>0.48 ± 0.34</td>
</tr>
<tr>
<td>( \phi )</td>
<td>0.46 ± 0.29</td>
<td>0.46 ± 0.28</td>
<td>0.39 ± 0.30</td>
</tr>
<tr>
<td>( N )</td>
<td>2837 ± 4310</td>
<td>2675 ± 4143</td>
<td>2896 ± 4347</td>
</tr>
</tbody>
</table>
component \( k \) \((k = 1, \ldots, K)\) will be more similar between them (close to a uniform distribution over the components), so that allocation is no longer so clear.

A. Brier Score

In order to assess the assignment performance of instances to clusters, the Brier score [21] is used. It allows us to measure the deviation of posterior probabilities from those that are expected. Therefore,

\[
\text{Brier} = \frac{1}{n} \sum_{j=1}^{K} \sum_{i=1}^{n} (P_{ij} - e_{ij})^2
\]

where \( P_{ij} \) is the calculated posterior probability of instances \( x_i \) and component \( j \), predicted by the model, and \( e_{ij} \) takes a value of 1 or 0 if it is the expected component to be allocated or not. For example, if the \( P_{ij} \) values of an instance \( x_i \) for three components are \([0.33, 0.34, 0.33]\), then \( e_{ij} \) is \([0, 1, 0]\). Accordingly, using (11)

\[
\text{Brier} = \frac{1}{n} \left[ (0.33 - 0)^2 + (0.34 - 1)^2 + (0.33 - 0)^2 \right] = 0.65.
\]

The lower the values the better, i.e., the assignment probability is more concentrated and less distributed between components. The value of the \( n \)-sized window, with \( n \) obtained from (8) of Chernoff bound, will be used as the calculation window of the Brier score.

To study how the concept drift affects the Brier score, the data streams used in the previous sections are used below.

B. Experimentation Results and Analysis

The Brier score is monitored over the three data streams to analyze its behavior regarding how the algorithm adjusts a better model after a concept drift detection. For these experiments, the parameters for high accuracy obtained in Section IV are taken \((\lambda = 22.6, \delta = 3.65, \epsilon = 0.52, \phi = 0.46, N = 2837)\) with the three data streams defined in Table II.

The results of the experiments are shown in Fig. 2, where the GDPC is able to detect the three concept drifts. In each case, every time the model changes (instance 15'000, 22'000, and 32'000), the Brier score changes its behavior and a general stabilization of the Brier score is obtained after the concept drift turning point is detected. As shown in the figures, the Brier score is stable at each leg of different variations described in Table I, which means that the model is constantly assigning, with the same confidence, the instances to specific components of the mixture, only changing if a concept drift is detected.

Therefore, the Brier score can be used to assess the quality of the new model detected after a concept drift, where the idea of dynamic behavior is to adjust the model during data stream processing to guarantee high clustering quality taking into account system changes over time, e.g., degradation.

VI. COMPARISON OF GDPC WITH STATE-OF-THE-ART ALGORITHMS

Using the optimal parameters defined for high accuracy, labeled \( \lambda, \delta, \epsilon, \phi, \) and \( N \) (Table XI) and data stream no. 3 (Table II), we compared GDPC against some well-known...
data stream clustering algorithms described in Section I: CluStream ($k$-means-based), D-Stream (DBSCAN-based), and ClusTree (hybrid algorithm with $k$-means and DBSCAN). These algorithms are available at the massive online analysis framework [38].

The analysis was performed on the synthetic data stream for three drifts at instances 15,000, 22,000, and 32,000. True instance cluster labels were output from the synthetic data stream. From the clustering results, the recall and F-score were computed in 1000-instance batches and then plotted to check the evolution throughout the data stream. The analysis results are shown in Fig. 3(a) and (b).

Looking at Fig. 3, D-Stream performance is comparable to GDPC in terms of recall during the first part of the stream. After the first concept drift occurs, the performance drops below the other algorithms, meaning that D-Stream does not have robust capabilities for concept drift detection. In terms of the F-score, the value for D-Stream is below par throughout the entire data stream with the lowest value of all four analyzed algorithms.

CluStream and ClusTree performed similarly, exhibiting low true positive rates throughout the entire data stream. However, ClusTree does react when the concept drift occurs, although it misses the concept drift located at instance 15,000. Both algorithms performed the best during the last leg of the data stream, though it is unclear whether this behavior is because the algorithm is not concept drift sensitive.

The GDPC algorithm does react to the concept drift, adapting online to the data stream to maintain the highest true positive rate. The algorithm detects the concept drift located at instance 15,000 perfectly, though it is noted that its performance drops in some instances. Once a new GMM model is calculated, the recall increases again, i.e., the algorithm adapts the model to account for the new data from the stream after the concept drift occurs. The next concept drift, at instance 22,000 is also detected, and a new GMM model is launched. This further increases the true positive rate to almost one. The last concept drift, instance 32,000, is detected and the GMM is launched again. In this case, the new model performs worse than CluStream and ClusTree.

Table XV provides a summary of these results.

VII. DYNAMIC PROBABILISTIC CLUSTERING APPLIED TO DATA STREAMS

A. Data Acquisition

In order to analyze the algorithm performance with data streams using a real machinery, an experimental testbed is used. This testbed, shown in Fig. 4, has the ability to reproduce the real behavior of a machine and all its subsystems from data acquisition to analytics as described in [39]. To simulate concept drifts produced by the machinery, the testbed is programmed to perform a servo-motor movement with different steps of angular velocity, expressed in revolutions per minute (RPM), in order to recreate different states or phases. Thus, the change in angular velocity defined as a concept drift is located every 4500 instances, totaling 25.
algorithm shows a high degree of accuracy for concept drift detection (98.7%). In this case, parameters for high accuracy and sensitivity have the best algorithm performance, achieving 98.7% accuracy and sensitivity of 96%, meaning that almost all detections are TP. However, there are about twice the amount of instances detected as turning points for concept drift. Further work on the algorithm development should be oriented to find an effective way to reduce that detection in real/unknown environments.

From the time \( t \) required to process the complete data stream of 117,822 instances and 25 concept drifts, the total effective processing time (required time upon data instance arrival) is 22 s, which means that the algorithm needs an average of 185 \( \mu \text{s} \) per instance, including the time taken to recalculate the model due to concept drift. In this particular scenario, where monitoring is done at 100 ms, real-time performance is achieved.

Additionally, taking the parameters for high accuracy, the Brier score is monitored (Fig. 6) to check how the concept drift is detected and then the model is adjusted decreasing the value of the score, so there is a high confidence in the assignment of instances to each component.

From the results, it can be seen that the algorithm has the capability to cluster the data stream based on a GMM with a
Fig. 6. Testbed results of the Brier score.

reasonable accuracy of around 99% and at the required time. This has been achieved with only 2% of the total available instances. Almost all of the 25 induced concept drifts have been detected along with others changes. These detected concept drifts could alert a monitoring system that the process is changing over time probably due to degradation or some other abnormal behavior. The knowledge obtained from these concept drifts and their evolution over time could be used for further analysis related to the aforementioned RUL.

VIII. CONCLUSION

The main conclusions of this paper are as follows.

1) A new algorithm called GDPC has been developed, which is capable of dealing with data streams with a limited amount of historical data and time needs. It detects concept drifts and adjusts the current model with fitting guarantee.

2) The easiest concept drift detected in a data stream is the first change, regardless of its type. From this point onward, the following changes represent a detection challenge. This behavior is caused by the window size used by the Chernoff bound, which tends to be smaller than the first training set as the algorithm always uses the smallest possible window. This effect has to be controlled using minimal values of ε and φ to maximize the window size.

3) The algorithm parameters, especially the training window size N, have a critical effect on the result, basically, in its ability to recognize concept drift but also in its accuracy. A slight variation in some of the parameters can cause big changes in the figures of merit.

4) The existing relationship between the algorithm parameters does not allow a simple representation with traditional tools for experimental analysis, this is due to the high dependency of all of them. That is why a naïve Bayes classifier can help to find a good parameter configuration in an automatic way.

5) Regardless of the model used, the size of the training set is the most critical factor and Chernoff bound parameters is the least critical factor.

6) Algorithm performance is related to the nature of the concept drift. Small changes, such as the introduction of noise at low standard deviation, are difficult to detect.

7) GMM clustering is a really powerful tool that is able to give useful information for posterior analysis, such as Brier score, mixture parameters, probability of assignment to the cluster, among others.

8) The Brier score is a proper tool to evaluate the ability of dynamic clustering algorithms to adjust the models during the data stream analysis. If the new model is not able to reduce the Brier score, the algorithm performance is not good enough.

9) Real-time applicability of the algorithm to the industry is possible because there is a small footprint in terms of the amount of data needed and processing time. That is, the algorithm uses a small size of data stream window that is continuously adapted to the data stream in order to increase efficiency. From this window, the algorithm is able to update the model if a change occurs without taking excessive processing time, as with the traditional static GMM. This represents a first step to deploying a probability-based clustering technique at the edge, where actionable insights are needed at the right time and accessibility to large computing systems is limited.

Further work related to the development of the algorithm is directed to parameter selection improvement with a more advanced classifier than naïve Bayes, which does not take into account the relationship between variables (parameters). Additionally, specific work should be done in order to optimize whether small changes, such as Gaussian noise with low standard deviation must be detected. On the other hand, work has to be done to increase algorithm features, such as the estimation for the number of components K depending on the training data, where new components could arise or be deleted during the data stream. This new step can leverage the algorithm to be used in novelty detection. Furthermore, the different Gaussian mixtures models produced during concept drift need be studied to differentiate between drift magnitudes, i.e., measure the distance between models. This is especially useful when the algorithm is used in real unsupervised environments, where the concept drift turning point is unknown.

As this algorithm is designed primarily for deployment in a IoT platform, further research has to be
conducted with regard to implementation in terms of computing power footprint. Some preliminary tests with GMMs have been conducted on system-on-chip devices, such as Zynq®UltraScale™MPSoC, where programmable logic can accelerate part of the algorithm without the need of complex distributed infrastructures.

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REFERENCES