

Application of Mixtures of Gaussians for Tracking Clusters in Spatio-temporal Data

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Abstract: Clustering data based on their spatial and temporal similarity has become a research area with increasing popularity in the field of data mining and data analysis. However, most clustering models for spatio-temporal data introduce additional complexity to the clustering process as well as scalability becomes a significant issue for the analysis. This article proposes a data-driven approach for tracking clusters with changing properties over time and space. The proposed method extracts cluster features based on Gaussian mixture models and tracks their spatial and temporal changes without incorporating them into the clustering process. This approach allows the application of different methods for comparing and tracking similar and changing cluster properties. We provide verification and runtime analysis on a synthetic dataset and experimental evaluation on a climatology dataset of satellite observations demonstrating a performant method to track clusters with changing spatio-temporal features.

1 INTRODUCTION

With the increasing amount of spatio-temporal data researchers across a wide variety of disciplines are facing new challenges mining and analysing datasets. Spatio-temporal data exhibit observations across space and time for example gathered by large sensor networks or satellites providing remote sensing or satellite imagery data. Spatio-temporal clustering is an active research area analysing spatial and temporal data at a higher level of abstraction by grouping data points according to their similarity into meaningful clusters (Trevor et al., 2009). Current approaches leverage variations of well-known methods and algorithms modified to operate on spatio-temporal data (Maciag, 2017). In this context approaches have been adapted for analysing trajectories and moving spatio-temporal clusters (Li et al., 2004; Kalnis et al., 2005). While current approaches often help to reveal and understand potential relationships many proposed methods lack the exploitation of available a priori knowledge that might improve the output quality (Maimon, 2010). Additionally, current algorithms are often limited in detecting substructures in large datasets; especially when clusters are overlapping, for example

when observations are taken continuously at the same locations, as it is often the case with spatio-temporal data. In this paper we propose a data-driven approach of tracking clusters in spatio-temporal data. Our approach is based on the Gaussian mixture model to extract cluster properties that can be analysed for changes over space and time. The concept is evaluated against synthetic data and real world climatology data from satellite observations. We provide a methodology based on well-known algorithms and an interpretation of the algorithmic results in a spatio-temporal context. Future possible extensions and modifications to the applied algorithms will be discussed in the conclusion of the paper.

The remainder of the paper is organized as follows: Section 2 provides the background on the Gaussian mixture model and the Bayesian Information Criterion for model selection while Section 3 presents the proposed concept in detail. Section 4 compares our proposed concept to related work and Section 5 presents the evaluation and exemplification of the concept in the area of climate research. At the end in Section 6 we give a discussion on the results while Section 7 provides the conclusions and outlooks.

All datasets together with the code for

this paper are publicly available online at <https://github.com/bert14398/kdir2019>.

2 PRELIMINARIES

This section shortly recalls the essential basics for this paper: the Gaussian mixture model and the Expectation-Maximization algorithm for fitting Mixture-of-Gaussian models to the given data together with the Bayesian Information Criterion for model evaluation.

2.1 Gaussian Mixture Model

Finite mixtures of distributions provide a sound mathematical-based approach for statistical modelling of a wide variety of random phenomena (McLachlan and Basford, 1988; McLachlan and Peel, 2004). These probabilistic models consist of superposition formed by linear combinations of basic distributions. By using a sufficient number of Gaussian distributions with adjusted means and covariances as well as adjusted contribution to the linear combination, any continuous density can be approximated to arbitrary accuracy with few exceptions. A Gaussian mixture model (GMM) can therefore be formulated as following (Bishop, 2006)

$$p(\mathbf{x}) = \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x}|\mu_k, \Sigma_k) \quad (1)$$

with each Gaussian density components $\mathcal{N}(\mathbf{x}|\mu_k, \Sigma_k)$ having its own mean μ_k and covariance Σ_k . The parameters π_k are called mixing coefficients, satisfying the condition

$$\sum_{k=1}^K \pi_k = 1 \quad (2)$$

Using the maximum likelihood to set the values for μ_k , Σ_k and π_k , the logarithm of the likelihood function from (1) is given by

$$\ln p(\mathbf{X}|\pi, \mu, \Sigma) = \sum_{n=1}^N \ln \left\{ \sum_{k=1}^K \pi_k \mathcal{N}(\mathbf{x}|\mu_k, \Sigma_k) \right\} \quad (3)$$

summing over N number of observations. Because of the summation over the number of clusters k Equation (3) has no closed-form analytic solution but can be estimated with the Expectation-Maximization algorithm.

2.2 Expectation Maximization

The Expectation-Maximization (EM) algorithm finds the local optimum for parameters in the likelihood for models with latent variables from the given data (Dempster et al., 1977).

For the Gaussian mixture model the algorithm first initializes the means μ_k , covariances Σ_k and mixing coefficients π_k and evaluates the initial value of the log likelihood. After initializing the parameters the algorithm iterates over expectation and maximization steps until convergence for maximizing the likelihood function.

In the expectation step posterior probabilities for each *responsibility* that the i -th data point x_i belongs to the k -th component of the mixture are computed as follows.

$$p(z_k = 1|x_i) = \frac{p(z_k = 1)p(x_i|z_k = 1)}{\sum_k p(z_k = 1)p(x_i|z_k = 1)} \quad (4)$$

Where z_k is the k -th element of the binary random variable \mathbf{z} with $\sum_k z_k = 1$ and $z_k \in 0, 1$, specified in terms of the mixing coefficients π_k such that $p(z_k = 1) = \pi_k$.

In the maximization step the parameters μ_k , Σ_k and π_k can be re-estimated using the computed responsibilities.

$$\begin{aligned} \mu_k &= \frac{\sum_i \{p(z_k = 1|x_i)x_i\}}{\sum_i p(z_k = 1|x_i)} \\ \Sigma_k &= \frac{\sum_i \{p(z_k = 1|x_i)(x_i - \mu_k)(x_i - \mu_k)^T\}}{\sum_i p(z_k = 1|x_i)} \\ \pi_k &= \frac{\sum_i p(z_k = 1|x_i)}{N} \end{aligned} \quad (5)$$

If the convergence criterion is not satisfied the algorithm returns to the expectation step.

2.3 Bayesian Information Criterion

Selecting the number of components in the Gaussian mixture model can be done in an efficient way with the Bayesian Information Criterion (BIC) (Chen and Gopalakrishnan, 1998) penalizing the likelihood by the number of clusters C_k . The number of parameters for each cluster is $d + \frac{1}{2}d(d+1)$ with data points $x_i \in \mathbb{R}^d$.

$$\text{BIC}(C_k) = \sum_{j=1}^k \left\{ -\frac{1}{2}n_j \log|\Sigma_j| \right\} - Nk(d + \frac{1}{2}d(d+1)) \quad (6)$$

By selecting the Gaussian mixture model with the lowest BIC we can ensure to choose the true number of clusters according to the intrinsic complexity present in a particular dataset in the asymptotic regime. This means, that the probability that BIC will select the correct model approaches one with increasing number of models up to infinity, but will tend to choose models that might be too simple, because of the penalty on complexity (Maimon, 2010).

3 PROPOSED MODEL

In this paper we propose a data-driven model to track clusters and their changing properties over space and time, based on the Gaussian mixture model and Bayesian Information Criterion as introduced in the previous Section 2. The model can be outlined in four basic steps.

1. Splitting the Data in Spatial Regions of Interest:

This first step needs careful consideration if splitting the data has a significant effect on the mixture components by including or excluding certain observations ("edge-effects"). Also splitting the data in spatial regions of interest (ROI) might not always be applicable, for example if all the clusters are tracked within one spatial area.

However, if the nature of the dataset allows the statistical evaluation and mitigation of edge-effects, for example by assuming or proving complete spatial randomness (Diggle, 2013), and if the analysis is conducted over multiple spatial areas, this step significantly increases the scalability of the model.

Each mixture model for any ROI can be computed in parallel, which decreases the overall runtime, see Section 5. Selection criteria for the size of the ROI typically depends on domain knowledge or the premise of the analysis.

2. Modelling the Observed Data in each ROI with Mixtures of Gaussians:

For each spatial region a range of Gaussian mixture component models with different maximum number of mixture components are fitted to the data. The model with the lowest Bayesian Information Criterion score determines the number of clusters for the most suitable mixture model, as described in the preliminaries in Section 2.

An alternative approach to infer the number of clusters of the most suited mixture model is the variational Bayes approach (Attias, 1999). This approach however requires additional fine tuning of hyperparameters and introduces additional computa-

tional overhead.

3. Extracting Cluster Parameters for each Gaussian Mixture Component:

Following a data summarization approach, we propose to extract the ellipsoid properties of the multivariate Gaussian distribution, the center and principal axes given by the mean and length of the eigenvectors of the covariance matrix as well as the polar angle of the major axis, since these properties most prominently describe the underlying distribution. However, different cluster properties can be selected according to the nature of the data and the expected behaviour of spatio-temporal changes of the clusters.

4. Comparison of Cluster Parameters for Spatio-temporal Changes:

The comparison can be done by a range of suitable methods, which demonstrates the flexibility of the proposed model. For the model evaluation and results in this paper, we used the DBSCAN clustering algorithm (Ester et al., 1996) on the extracted cluster properties to find similar clusters and account for spatio-temporal changes.

By comparing the extracted cluster parameters between clusters in different spatial regions over time we can identify similar clusters and further review these occurrences to investigate any underlying motion.

4 RELATED WORK

Using Gaussian mixture models for image matching has been proposed in a continuous probabilistic framework by Greenspan et al. (2001). Greenspan et al. propose a transition of the image pixels to coherent regions in feature space via Gaussian mixtures to apply a probabilistic measure of similarity between the Gaussian mixtures. To determine the number of mixture components Greenspan proposes the minimum description length principle compared between cluster sizes ranging from three to six. Similar images are identified by the Kullback–Leibler divergence (Kullback, 1997) of their mixing components. Compared to our approach we are not relying on the Kullback–Leibler divergence measure to identify similar mixing components, but extract cluster features that can be further clustered for similarity with different cluster algorithms.

A dynamic model-based clustering method for spatio-temporal data with finite mixtures of Gaussians with spatio-temporal varying mixing weights was presented by Paci and Finazzi (2018). In their work,

Paci and Finazzi adjusted the mixing coefficients in a way that similar observations at near space and time points are assigned with similar cluster membership's probabilities. However, the model is only formulated for the univariate case of observations and also the Bayesian approach has a higher model bias than the proposed model in this paper.

Jin et al. (2005a,b) proposed a clustering system based on the Gaussian mixture model with independent attributes within clusters. They modified the EM-algorithm introduced in Section 2 to include the clustering features of sub-clusters. Jin et al. use on the one hand a grid-based method for cluster features extraction and on the other hand the BIRCH clustering algorithm (Zhang et al., 1996). In our proposed model, we use the Gaussian mixture model for data summarization and an additional clustering algorithm to group similar cluster properties together. The advantage of our approach is that we do not need to modify the Expectation Maximization algorithm and allow the application of different clustering algorithms that might be better suited for data with outliers or noise.

Providing a more formal definition of moving clusters, Kalnis et al. (2005) present three algorithms based on the DBSCAN clustering algorithm to identify moving clusters over a period of time. Moving clusters are identified over consecutive time slices if the ratio of their intersect density to their joint density is greater or equal as a specified threshold. While allowing to track clusters based on their densities over time, the presented work by Kalnis does not provide the necessary means to monitor changing cluster properties over space and time. Our approach does not demand intersecting densities to identify moving clusters and is further able to identify reappearing clusters and can track clusters with changing properties over space and time as well.

5 MODEL EVALUATION

In this section we evaluate our proposed model on a synthetic dataset and a real dataset of satellite observations.

The synthetic dataset has been generated of similar size and structure as our real dataset, but with already known distributions in feature space. The spatial points are generated from a *N*-conditioned *Complete Spatial Randomness* (CSR) process (Diggle, 2013). A *N*-conditioned CSR process is defined as: *Given the total number of events N occurring within an area A, the locations of the N events represent*

Table 1: Dataset Descriptions.

Time steps	Real data # points	Synthetic data # points
2014-02-12	154,031	160,001
2014-02-13	157,719	160,000
2014-02-14	167,473	160,000
2014-02-15	161,441	160,000
2014-02-16	155,187	160,000
2014-02-17	164,059	159,999
Total	959,910	960,000
Features	{x, y}	{ln(H ₂ O), δD}
Spatial extent	162 clusters	162 20x20 boxes

an independent random sample of N locations where each location is equally likely to be chosen as an event (Rey and Anselin, 2007). The two dimensional feature patterns are well separated isotropic Gaussian blobs and assigned to k-means clustered spatial regions. The spatial partitioning in the data generation process (k-means clustered spatial regions) is different from the spatial partitioning in the analysis (regular lattice) to better evaluate possible edge-effects. The number of k-means clusters is equal to the number of spatial grid cells, 162, so that clusters and grid cells have similar spatial extent, but some data points at the cluster edges will be assigned to different grid cells during the analysis.

The satellite observations have been gathered by Metop-A and Metop-B satellites with the IASI (Infrared Atmospheric Sounding Interferometer) instrument (EUMETSAT, 2018). IASI measures in the infrared part of the electromagnetic spectrum at a horizontal resolution of 12 km over a swath width of about 2,200 km, providing information on the vertical structure of the atmospheric temperature and humidity in an unprecedented accuracy of 1 K and a vertical resolution of 1 km. Compared to the synthetic data points, the real observations are less dense due to different filters, missing observations and processing errors. Ideally, we would have global coverage with no missing data points and only variations in the distributions of latitude and longitude of the satellite measurements as in our synthetic dataset. Table 1 outlines the two datasets, while a visualization of both datasets is provided in Figure 1 and 2. Additionally, all data is provided publicly available online.

We applied our proposed model described in Section 3 on each of the six consecutive days in Table 1 as follows:

1. Splitting the Data in Spatial Regions of Interest:

We split the data into geospatial regions on a regular grid with grid size of 20 x 20 degrees for longitude

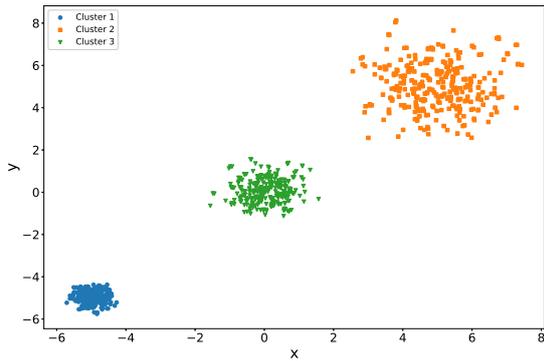


Figure 1: Feature distribution of a random grid cell for the synthetic dataset. Markers and colors correspond to the GMM clusters identified.

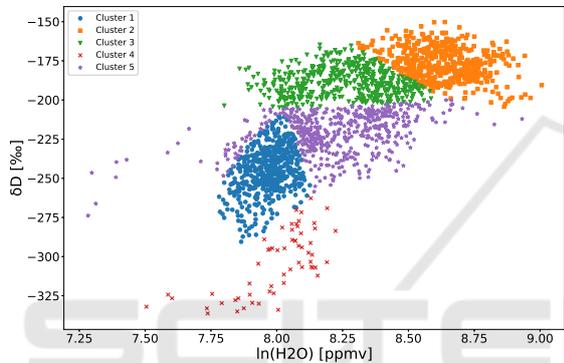


Figure 2: Feature distribution of a random grid cell for the climatology dataset. Markers and colors correspond to the GMM clusters identified.

180 degrees West to 180 degrees East and latitude 90 degrees South to 90 degrees North. The size of the ROIs here is postulated by the domain expert to conduct an analysis of far-reaching climatological events. This step results into 162 ROIs, which we impose on our real and synthetic dataset.

2. Modelling the Observed Data in each ROI with Mixtures of Gaussians: For each spatial region we fit multiple Gaussian mixture models with at least two observations per cluster and a maximum number of 10 components. Each model is evaluated according to its BIC score and the model with the lowest BIC score is selected as the best model. The maximum number of 10 components has been selected out of multiple experimental runs, where we determined the trade-off between searching for models with a higher number of components and lower BIC scores and the number of observations per cluster.

3. Extracting Cluster Parameters for each Gaussian Mixture Component: For each Gaussian

mixture component identified in step 2, we extracted the ellipsoid properties of the multivariate Gaussian distribution, the center and principal axes given by the mean and eigenvectors of the covariance matrix as well as the angle of the major axis. More specifically we extracted the major and minor axis length and the axis angle in degrees for the bivariate contour where 95% of the probability falls.

4. Comparison of Cluster Parameters for Spatio-temporal Changes: In the last step we used the DBSCAN algorithm with a maximum distance between two samples $Eps = 0.3$ and the number of minimum points $MinPts = 2$ in each Eps-neighbourhood. Each Eps-neighbourhood is defined by the specified radius Eps and the number of minimum points $MinPts$ within Eps from a point under consideration, so that this point can be identified as a core point. For further details on the DBSCAN algorithm we refer to the paper of Ester et al. (1996). Similar to step 2, the DBSCAN parameters have been assessed and evaluated on the data through empirical analysis.

5.1 Verification

To verify our model we take a look at the feature distributions and the corresponding GMM clusters together with the identified groups of clusters by the DBSCAN algorithm.

We can visualize the identified groups of cluster by the DBSCAN algorithm by plotting the mean ellipsoids of each DBSCAN cluster. Figure 3 and Figure 4 show the ellipses defined by the extracted properties, the major and minor axis length and the axis angle in degrees for the bivariate 95% contour. The number of clusters for the synthetic dataset shown in Figure 3 indicates three groups of similar clusters in accordance with our initially defined three separated clusters for each spatial region. A fourth group is dedicated to DBSCAN outlier results.

For the real dataset we can see much more overlapping groups of clusters depicted in Figure 4, in total 250 groups including the dedicated outlier group. This result is consistent with the feature distribution in the dataset, exemplified in Figure 2, and will be further discussed in the next subsection.

As exemplified by the feature distributions of a random spatial grid cell in Figure 1 and 2, the GMM clusters in the synthetic dataset in Figure 1 are identified as clearly separable as generated, although the imposed spatial lattice is different from the initial spatial k-means clustering in the generation process. We can conclude that as stated in Section 3 the synthetic dataset allows the statistical evaluation and mitigation

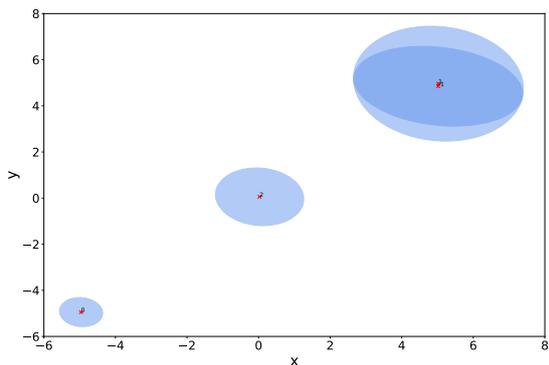


Figure 3: Mean ellipsoids of each DBSCAN cluster group of clusters for the synthetic dataset.

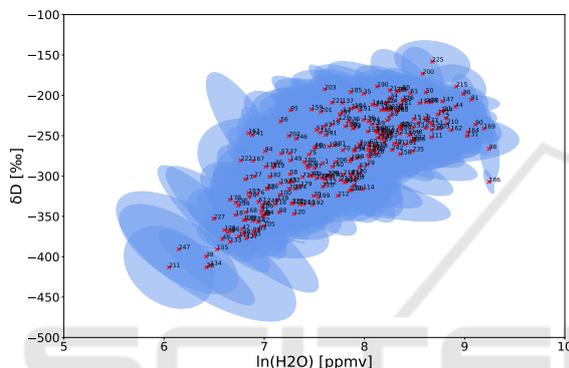


Figure 4: Mean ellipsoids of each DBSCAN cluster group of clusters for the climatology dataset.

of edge-effects. Data points that initially belong to a different spatial region than our imposed lattice are merely contributing to the feature distribution but do not alter the distribution. The feature distribution in the real dataset illustrated in Figure 2 do in contrast not show any apparent separable clusters, however the properties of the GMM clusters fitted to the data allows to track moving clusters, emerging clusters and cluster changes.

5.2 Application to Climate Research

Our real dataset consists of spectral data gathered from Metop-A and Metop-B satellites that have been processed for the water vapour H_2O mixing ratio and water isotopologue δD depletion for air masses at 5 km height with most sensitivity. The water isotopologue in question is HDO , which differs only in the isotopic composition compared to H_2O . Isotopologues of atmospheric water vapour can make a significant contribution for a better understanding of atmospheric water transport, because different water transport pathways leave a distinctive isotopologue fingerprint (Schneider et al., 2017).

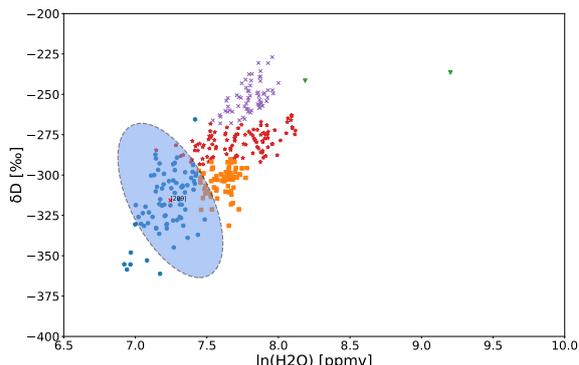


Figure 5: Grid cell at latitude 50 degrees South to 30 degrees South and longitude 120 degrees East to 140 degrees East at 2014-02-14. The ellipse visualizes the cluster properties of the moving cluster.

5.2.1 Tracking Moving Clusters

As an example of tracking moving clusters we are looking at all GMM clusters that DBSCAN has assigned to the same group in neighbouring grid cells. Here we identify moving clusters as clusters that appear in neighbouring grid cells with one day time delay. This example has been selected for demonstration purpose, but the proposed method allows to analyse similar clusters with different time lags and paths across the imposed lattice as well. If two GMM clusters are similar according to DBSCAN and one is present in grid cell A at day 1 and present in grid cell B, a neighbour grid cell to A, at day 2 we conclude that the cluster or cluster generating process has moved from cell A to cell B. Our results show multiple moving clusters according to the above definition; specifically we could identify 105,616 occurrences that comply with the above definition of moving clusters. Figures 5 and 6 give one example of two neighbouring grid cells with observations from 2014-02-14 and 2014-02-15, at latitude 50 degrees South to 30 degrees South and longitude 120 degrees East to 140 degrees East and 140 degrees East to 160 degrees East respectively. Both cells contain a specific cluster that DBSCAN has assigned to the same group and which is highlighted with the corresponding ellipses.

The presented result has been selected as a representative example for detecting moving clusters between spatial regions. The clusters identified in Figure 5 and Figure 6 show close similarities and allow additional visual verification. However, the proposed method allows also to track clusters with varying cluster properties, which might not be immediately apparent.

Our proposed model is not limited to the definition of moving clusters used in this example. As men-

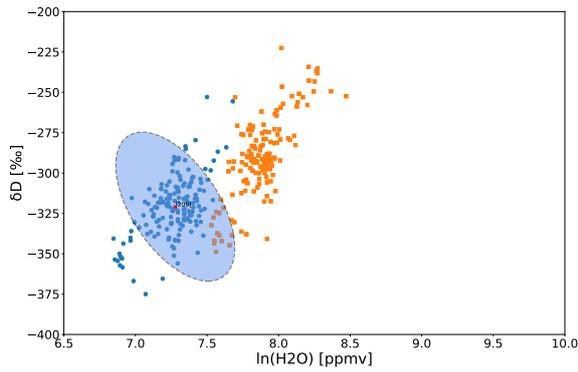


Figure 6: Grid cell at latitude 50 degrees South to 30 degrees South and longitude 140 degrees East to 160 degrees East at 2014-02-15. The ellipse visualizes the cluster properties of the moving cluster.

tioned at the beginning of this section, the definition of a moving cluster has been generalized to one spatial neighbour and one time lag, in our case one day. But each step in our model can be adjusted to the premise of the analysis; for example by imposing different spatial structures and temporal slices (Step 1), varying the number of mixture components (Step 2), applying and adjusting different cluster algorithms for clustering the mixture components properties (Step 3) and analysing the clusters of cluster properties according to the definition of the motion (Step 4), including more complex search patterns such as trajectories across multiple spatial regions with varying timestamps, which will be explored in future work.

5.2.2 Tracking Emerging Clusters

By comparing clusters of the same group in the same spatial region over time we can also identify emerging and disappearing clusters. For example if we search for occurrences over three consecutive days, where a cluster has been identified on the first and third day, but not on the second day. One example of an emerging cluster is given in Figure 7, 8 and 9.

Detecting emerging, disappearing and reappearing clusters with varying cluster properties in climatology data can be a strong indicator for emerging, disappearing and reappearing climatology events. In the presented case the emerging and disappearing cluster in the $\{H_2O, \delta D\}$ feature space can be associated with atmospheric water transport due to mixing of air masses with distinctive isotopologue fingerprints.

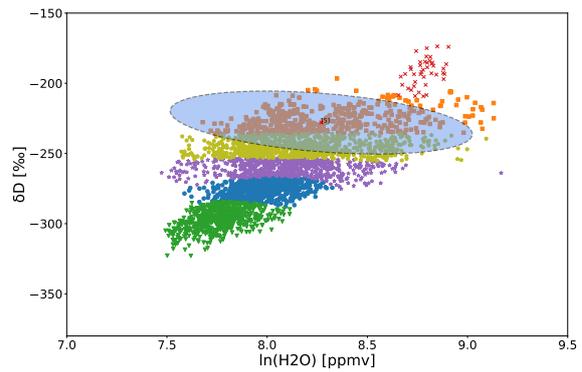


Figure 7: Grid cell at latitude 30 degrees South to 10 degrees South and longitude 160 degrees West to 140 degrees West at 2014-02-15. Cluster is present.

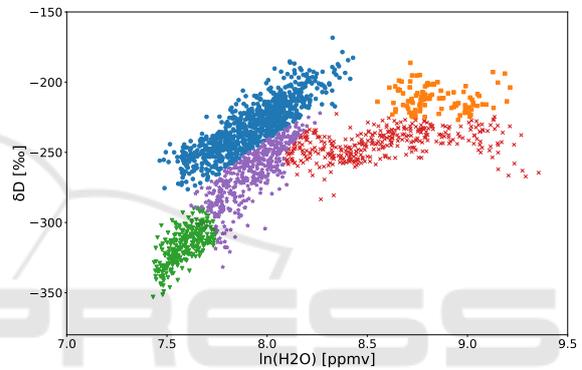


Figure 8: Grid cell at latitude 30 degrees South to 10 degrees South and longitude 160 degrees West to 140 degrees West at 2014-02-16. Cluster is absent.

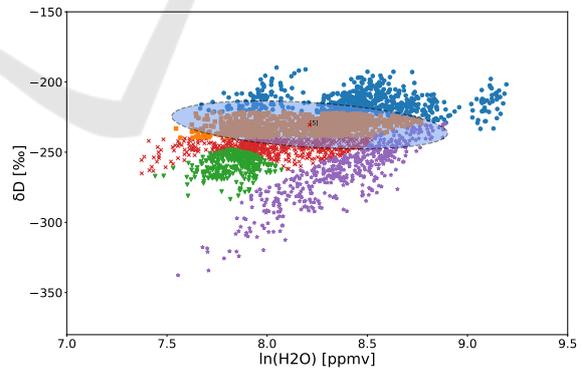


Figure 9: Grid cell at latitude 30 degrees South to 10 degrees South and longitude 160 degrees West to 140 degrees West at 2014-02-17. Cluster is present again.

5.2.3 Tracking Changing Clusters

The proposed approach allows additionally to compare statistics of cluster properties within the same group and across cluster groups. By looking at the mean, standard deviation, minimum, maximum

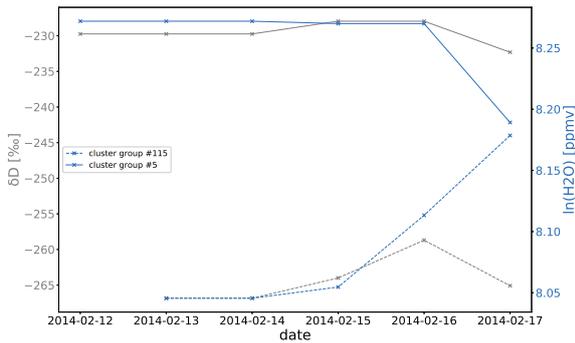


Figure 10: Temporal changes of $\ln(H_2O)$ as an example variable over five consecutive days within different cluster groups. Continuing trends across cluster groups give indications of cluster evolution.

and percentiles of the identified clusters within the same group and across groups, these statistical measures can provide valuable insight into the variability of cluster properties and possible inference between clusters and associated climatology events.

We can look for example at the temporal changes of cluster properties of any cluster group to identify trends in the cluster features. If the trends are continuously decreasing or increasing, we can analyse if the next closest cluster group can be considered an evolution of the original cluster group. Figure 10 illustrates this on the basis of the temporal changes of $\ln(H_2O)$ as an example variable over five consecutive days within two different cluster groups. While the variable δD for both example cluster groups (#115 and #5) does not show an apparent trend, the $\ln(H_2O)$ variable for both cluster groups seem to converge to the same level. This can give rise to further in-depth analysis of both cluster groups to explore possible interactions between them.

5.3 Runtime Measurements

As stated in Section 3 each mixture model for any ROI can be computed in parallel, which decreases the overall runtime significantly. To demonstrate the scalability of the model, we run tests on both the real data and synthetic data sequentially with increasing number of maximum mixture components and in parallel. Figure 11 and Figure 12 illustrates the increasing amount of computational time necessary with the increasing number of maximum mixture components, starting from one up to 20 components.

The measurements have been taken on a ordinary workspace computer with eight Intel©Xeon©CPU E3-1246 v3 cores and 32 GB main memory. For the sequential execution of fitting the GMM models to one spatial region after another one CPU core was

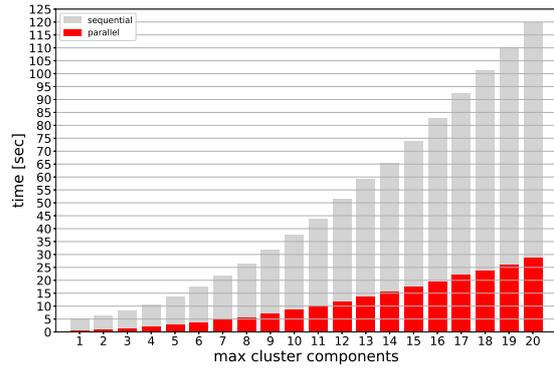


Figure 11: Sequential and parallel runtime with increasing maximum number of GMM cluster components for the climatology dataset. The runtime plotted is the time measured for the best run out of seven runs.

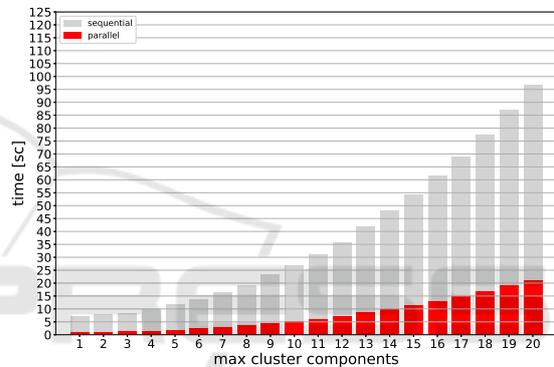


Figure 12: Sequential and parallel runtime with increasing maximum number of GMM cluster components for the climatology dataset. The runtime plotted is the time measured for the best run out of seven runs.

busy for up to around two minutes for the real dataset and around 95 seconds for the synthetic dataset. Running the GMM fitting for eight spatial regions in parallel by distributing the work as separate processes on all eight cores reduced the runtime significantly by at least a factor of four, providing the same overall results.

Figure 13 shows the decreasing runtime with increasing number of CPUs. The number of Gaussian mixture components has been set to 20 for all runs. We can see, that the maximum gain has been achieved with four cores, while adding more cores decreases the runtime more slowly towards the minimum time required to analyse one single spatial cell.

By looking at the runtime measurements, we can conclude that our approach scales well with the number of spatial regions computed in parallel. While an increasing number of maximum mixture components requires an almost quadratic increasing amount of time, the overall computational time can be signif-

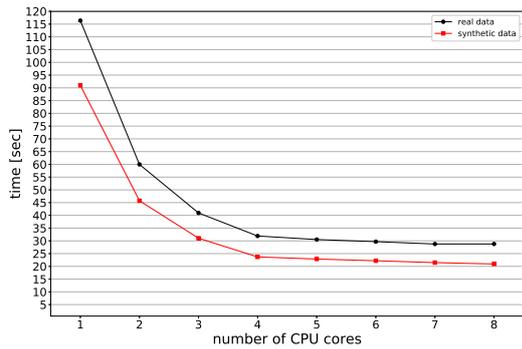


Figure 13: Runtime with different numbers of CPU cores for the real and synthetic dataset. The maximum number of GMM components is fixed to 20. The runtime plotted is the time measured for the best run out of seven runs.

icantly decreased by running the GMM fits for each spatial region in parallel.

These results highlight the scalability of our approach which becomes more and more important with the increasing amount of data that needs to be analysed. In fact, even if splitting the data in spatial regions of interest (ROI) might not be applicable, analysing sub-samples can be done in parallel as well. By uniform random sampling or biased sampling the operation of general data mining tasks like clustering can be significantly speed up (Kollios et al., 2003).

6 DISCUSSION

The initial step of our proposed model, dividing the data in spatial regions of interest, allows the parallel computation of Gaussian mixture models per area, which considerably increases the scalability of the approach. Even if dividing the data into spatial regions before applying the Gaussian mixture model might not always be applicable, for example if splitting the data has a significant effect on the mixture components by including or excluding certain observations. In these cases an analysis of the point generating process can help to determine, if the edge-effects can be statistically evaluated and mitigated (Diggle, 2013).

To further improve the scalability of our model, each Gaussian mixture model could theoretically be run in parallel. Instead of sequentially evaluating the BIC score, the model with the best BIC score could be selected from parallel computations. Future work will take this into account to analysis datasets spanning over several years as compared to several days presented in this paper.

The results of the experimental evaluation in Section 3 show that our approach is able to successfully

detect changes in cluster properties over space and time. The examples illustrated in Section 5.2 identify clusters moving between neighbouring spatial regions with similar cluster properties according to the applied DBSCAN algorithm on all extracted cluster properties. Although the proposed approach is not restricted to any method for comparing the Gaussian mixture components properties, DBSCAN shows good results and has the advantages that the number of clusters has not to be specified as a model parameter and the *Eps* and *MinPts* parameters can be fine-tuned according to the cluster properties.

While identifying spatio-temporal changes successfully, one drawback of our proposed approach is that modelling the data with mixtures of Gaussians with the number of components dependent on the model BIC score is not necessarily describing the underlying process that generated the data accurately. Therefore the interpretation of the results additionally relies on domain knowledge associating cluster and cluster changes to processes.

The utilization of more specific models that can also capture the underlying data generating processes more accurately is part of ongoing research.

7 CONCLUSION

In this article we present a scalable approach for tracking clusters in spatio-temporal data. The presented method models the observed data in each spatial region with a mixture of Gaussians and compares extracted cluster properties over time. By dividing the initial dataset into spatial regions or applying sampling techniques such as random uniform sampling or biased sampling, each subset can be processed independently and in parallel, which significantly improves the overall runtime of the proposed model. As verified on synthetic data with known data generating processes and applied to real world climatology data, the proposed model can reliably detect cluster changes over space and time, indicating moving clusters, appearing and disappearing events as well as evolution of clusters over time.

In the future, we plan to apply different Bayesian models in exchange for the Gaussian mixture model to incorporate more a priori domain specific knowledge to better catch the underlying processes that generated the data. Also a more in-depth evaluation in regard to the scalability of the proposed concept is planned to be discussed in follow-up work, together with a detailed evaluation of the clustering results and their application to climate research.

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