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*Article*

# Proposal of a Mathematical and Computational Method for Determining the Optimal Number of Clusters in the K-Means Algorithm

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**Abstract:** (1) Background: The rapid evolution of the internet and technological infrastructure has led to a surge in data generation across various contexts, increasing the use of machine learning tools to extract valuable information. Clustering, particularly using the K-means algorithm, is a common technique. However, determining the optimal number of clusters in K-means is challenging, as traditional methods like the elbow method can be imprecise and subjective. This study proposes a more accurate and objective method to identify the optimal number of clusters. (2) Methods: The proposed method utilizes the numerical derivative of cluster inertias and the maximum value of the ratio between contiguous derivatives. Implemented in Python using sklearn, numpy, and matplotlib, the method was validated with synthetic datasets generated by artificial intelligence, where cluster numbers are clearly distinguishable. (3) Results: The method proved to be more precise and less subjective than the traditional elbow method, accurately identifying the optimal number of clusters in all tested synthetic datasets. Additionally, it demonstrated computational efficiency with minimal RAM usage and execution time, making it suitable for practical data analysis applications. (4) Conclusions: This new mathematical and computational method significantly improves the determination of the optimal number of clusters in K-means, offering a more accurate and objective alternative to traditional techniques. Future work will extend this method to hierarchical clustering and develop a cloud service for wider accessibility

**Keywords:** machine learning; kmeans; mathematical method; elbow method

## 1. Introduction

With the growth of the internet and the extensive dissemination and development of information and communication technologies, a data revolution has been fostered, leading companies from diverse application contexts to generate vast amounts of data. Consequently, these companies face the significant challenge of transforming such data into information for decision-making through artificial intelligence and machine learning [1–6]. In a similar vein, thanks to the availability of not only data but also computational resources and tools, there has been a renewed interest in applying machine learning techniques to solve problems where conventional methods exhibit shortcomings [7–9].

Machine learning is a technique within artificial intelligence that seeks to emulate the human brain's capability by extracting generalized knowledge or patterns from a set of historical data to make predictions when faced with new data [10,11]. Machine learning can be categorized into two types: supervised and unsupervised. In supervised learning, the model is trained with labeled data, meaning examples that include both inputs and their correct outputs. In unsupervised learning, the data is unlabeled, and the goal is to identify patterns and ratios within the data to group them into categories [7,12–14], which are typically determined through distance or similarity metrics [15].

One of the most widely utilized unsupervised learning models is K-means, where specifying the number of groups or clusters to form from the dataset instances based on distance metrics is essential as an input parameter [16,17]. K-means aims to organize the instances of a dataset into K groups, minimizing the sum of squared distances between instances and their centroids [18,19]. To determine the optimal number of clusters, the elbow method or elbow criterion is frequently used, which involves plotting the inertias obtained for different numbers of clusters and identifying the inflection point or elbow, where the decrease in inertias becomes less pronounced [20–24].

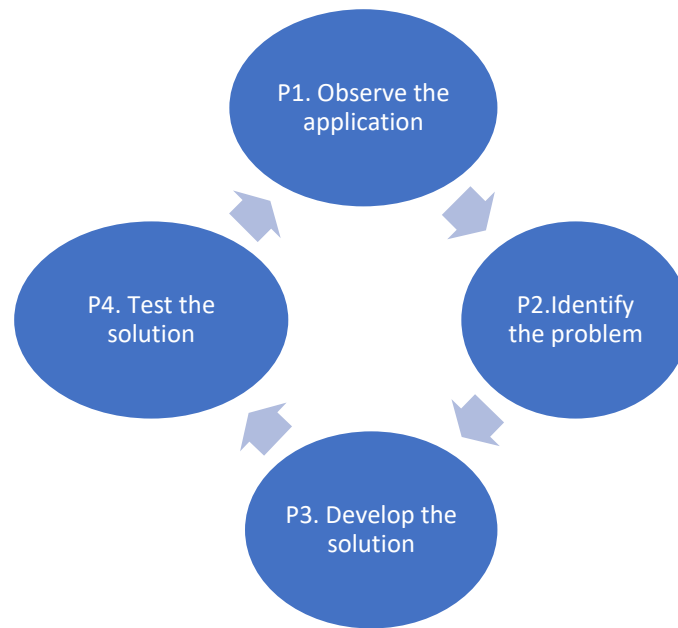
Despite the widespread use of the elbow method, one of its major challenges is visually determining the point where the variation is least pronounced, especially in datasets where data separation is not clear, as seen in Figure 3, where this selection criterion could be quite subjective. Therefore, it is necessary to have mathematical tools that enable a more precise and less subjective determination of the optimal number of clusters [25,26]. In this work, we propose a mathematical and computational method for determining the optimal number of clusters in unsupervised learning, supported by the KMeans model. This method is based on the mathematical analysis of the inertias or sums of intra-cluster squared distances for different numbers of clusters, thereby obtaining the most probable number of clusters to use for grouping the data. This method serves as an alternative to the shortcomings of the elbow method when it is difficult to visually distinguish the optimal number of clusters. It is also noteworthy that the proposed method can be used with datasets where the optimal number of clusters can be easily differentiated using the elbow method, thereby serving as a means to corroborate these results.

The method was implemented using the advantages provided by the sklearn, numpy, and matplotlib libraries [27]. Additionally, it was validated with five datasets generated through artificial intelligence, where the number of clusters is clearly distinguishable, allowing for verification of the method's relevance. This method aims to serve as a reference for use in various work contexts involving supervised learning supported by K-means, enabling more efficient and effective determination of the optimal number of clusters.

The remainder of this work is organized as follows: Section 2 presents the methodological phases considered for the development of this study. Section 3 describes the results obtained, including the design and implementation of the method, as well as the evaluation of its efficacy in determining the optimal number of clusters and its computational efficiency. Additionally, this section discusses the proposed method in comparison to other state-of-the-art approaches. Finally, Section 4 presents the conclusions and future work derived from this research.

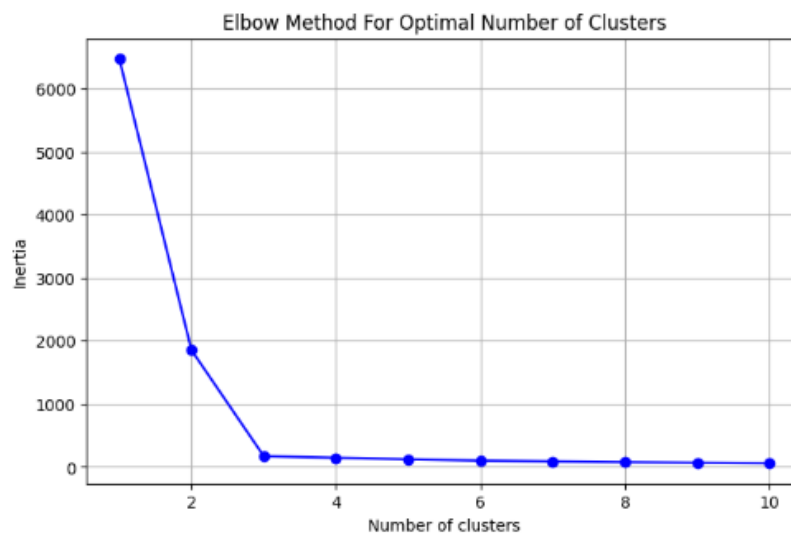
## 2. Materials and Methods

For the development of this research, the four methodological phases of Pratt's iterative research pattern were considered [28]: P1. Observe the application, P2. Identify the problem, P3. Develop the solution, P4. Test the solution (see Figure 1).



**Figure 1.** Methodology considered.

In the first phase of the methodology, the elbow method was characterized, which is employed in the K-means model to determine the optimal number of clusters into which the dataset should be divided for a more effective and precise interpretation of each group. For instance, Figure 2 shows the plot of inertia calculated from 1 to 10 clusters for a dataset of 100 instances. It is evident that starting from the third cluster, the inertia does not show significant variations, indicating that the optimal value is  $k=3$ .



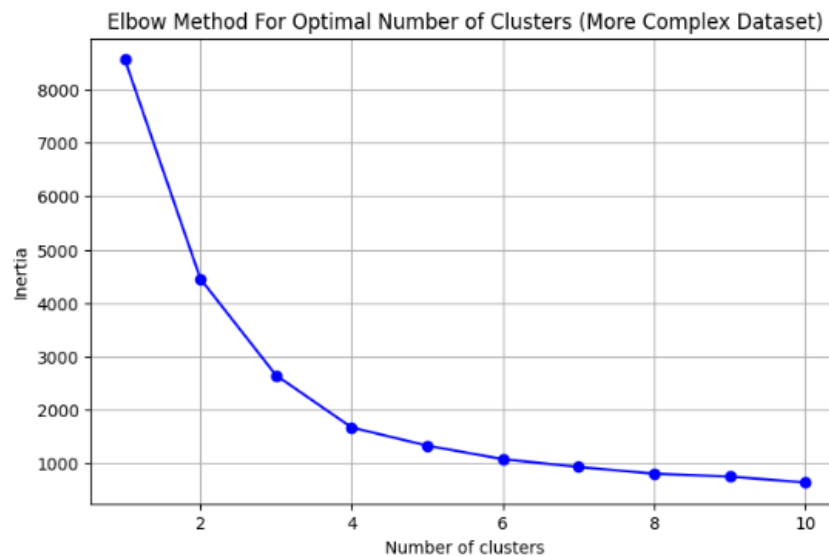
**Figure 2.** Example of the elbow method in KMeans

It is worth noting that each point in the graph in Figure 2 corresponds to the inertia for a given number of clusters  $k$ , which is determined by equation (1).

$$Inertia = \sum_{i=1}^n \min_{\mu_j \in C} \|x_i - \mu_j\|^2 \quad (1)$$

Where  $n$  is the total number of instances in the dataset,  $x_i$  corresponds to the  $i$ -th instance of the dataset,  $\mu_j$  is the centroid of the  $j$ -th cluster,  $C$  is the set of all cluster centroids, and  $\|x_i - \mu_j\|^2$  is the squared Euclidean distance between instance  $x_i$  and centroid  $\mu_j$ .

On the other hand, within phase 2, the limitation of the elbow method was identified, which can be summarized as the difficulty in visually differentiating the point where the cluster inertia is less pronounced [25]. For instance, Figure 3 presents a graph of the calculated inertia from 1 to 10 clusters for a dataset with 100 instances, in which it is not easy to discern the optimal number of clusters.



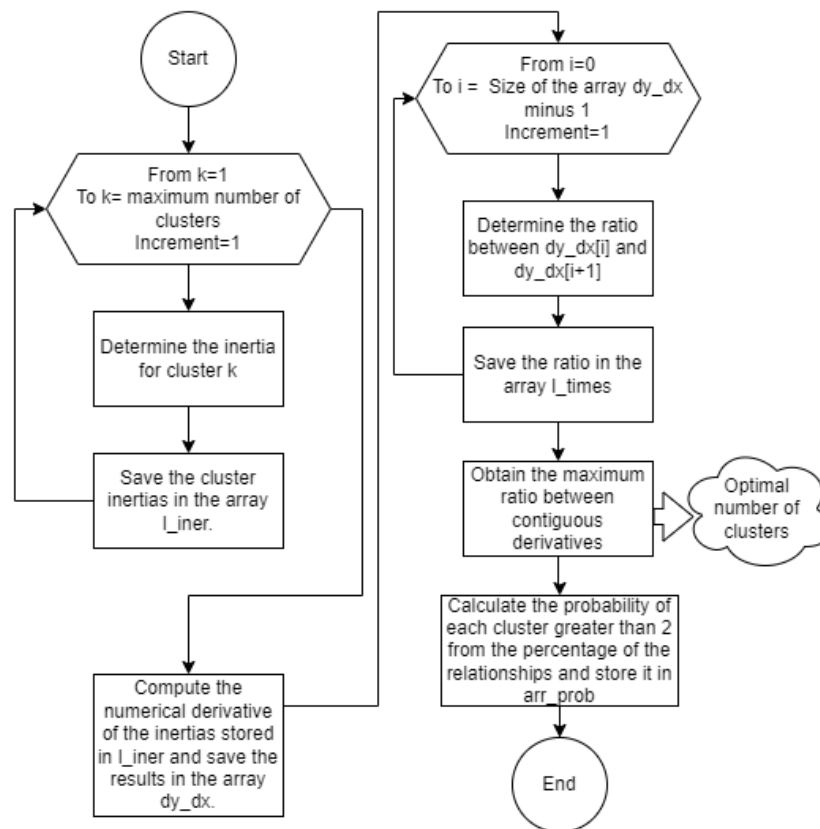
**Figure 3.** Limitation of the elbow method

In phase 3 of the methodology, a mathematical method was designed based on obtaining the first numerical derivative of each inertia point and the subsequent numerical ratio between the derivatives of consecutive inertia points. This was done to determine the maximum ratio between contiguous derivatives and to assess the probability of each cluster being the optimal number. Subsequently, the method was implemented computationally using the sklearn, numpy, and matplotlib libraries. The sklearn library was used to determine the cluster inertias, the numpy library was utilized to calculate the numerical derivatives and identify the maximum variation, and finally, the matplotlib library was employed to generate the corresponding graphs for the clusters and the elbow method.

Finally, in phase 4, once the mathematical method was implemented, its validation was carried out using 5 datasets generated by AI, in which the optimal number of clusters is 2, 3, 4, 5, and 6 clusters. This allowed for verifying the effectiveness of the method in correctly identifying the number of clusters. Additionally, the performance of the method was evaluated by monitoring RAM consumption and execution time in a Google Colab cloud environment.

### 3. Results and Discussion

This section may be divided by subheadings. It should provide a concise and precise description of the experimental results, their interpretation, as well as the experimental conclusions that can be drawn.



**Figure 4.** Mathematical method implemented

According to Figure 4, an iteration is first performed from 1 to the maximum defined number of clusters to determine the inertia for each cluster count, with these inertia values stored in an array named `l_iner`. Once the different inertias for the various clusters are calculated, the method computes the numerical derivative of the values stored in the `l_iner` array, with the resulting derivatives stored in the `dy_dx` array. Subsequently, an iteration is performed over the `dy_dx` array from index 1 to the array's length minus 1, calculating the ratio between the derivative `dy_dx[i]` and the subsequent derivative `dy_dx[i+1]`. These ratios are then stored in the `l_times` array. The maximum ratio from the `l_times` array corresponds to the optimal number of clusters. Additionally, the selection probability for each cluster is determined based on the percentage of each ratio in the `l_times` array.

The method designed and presented in Figure 4 was implemented in Python (see Figure 5) using the `sklearn` and `numpy` libraries. The `sklearn` library was used to determine the inertias associated with each considered number of clusters (`inertia_` attribute). Similarly, the `numpy` library was used to compute the numerical derivative of the cluster inertias (`diff()` function), as well as the maximum ratio between contiguous derivatives and the selection probability for each cluster greater than 2 (`argmax()` function). Additionally, it is important to highlight that the `matplotlib` library was used to generate the elbow and clustering graphs.



```

def k_method(data, max_cent):
    #Determination of inertias for the n defined clusters
    l_iner = []
    num_clus = range(1, max_cent+1)
    for k in num_clus:
        km = KMeans(n_clusters=k)
        km = km.fit(data)
        l_iner.append(km.inertia_)

    x=np.arange(1,max_cent+1)
    y=np.asarray(l_iner)

    # Calculation of the numerical derivative of the inertias
    dx = np.diff(x)
    dy = np.diff(y)
    dy_dx = dy / dx
    dy_dx = abs(dy_dx)
    print(f"Numerical derivative: {dy_dx}")

    # Calculation of the relationship between contiguous slopes or derivatives
    l_times=[]
    for i in range(0,len(dy_dx)-1):
        n_times=(dy_dx[i]/dy_dx[i+1])
        l_times.append(n_times)

    #Obtaining the maximum ratio between contiguous derivatives
    arr_n_times=np.asarray(l_times)
    n_clus=np.argmax(arr_n_times)+2

    #Obtaining the probability for the second cluster onward
    arr_prob= (arr_n_times / arr_n_times.sum())

    print(f"Array of relations: {arr_n_times}")
    print(f"Optimal number of clusters: {n_clus}")
    print(f"Probability array: {arr_prob}")

```

**Figure 5.** Mathematical method implemented in Python

To evaluate the effectiveness of the proposed method, 5 datasets of 100 instances each were generated with the aid of AI. Each dataset shows a clear division into groups of 2, 3, 4, 5, and 6, respectively (see Figure 6). These datasets were used to assess the proposed method, aiming to determine if the number of clusters obtained by the method matches the groups observed in each dataset.

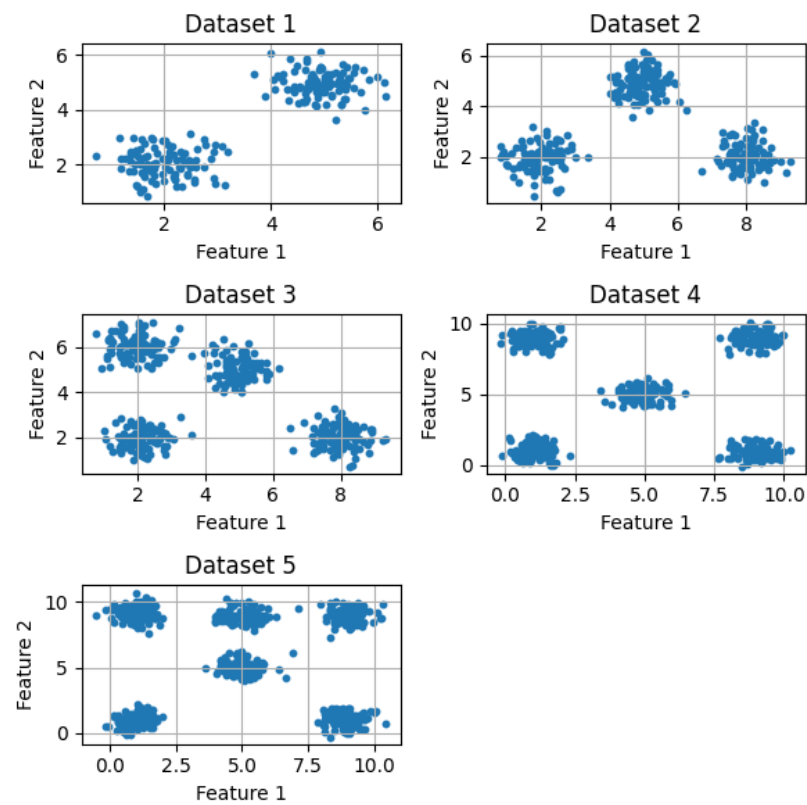
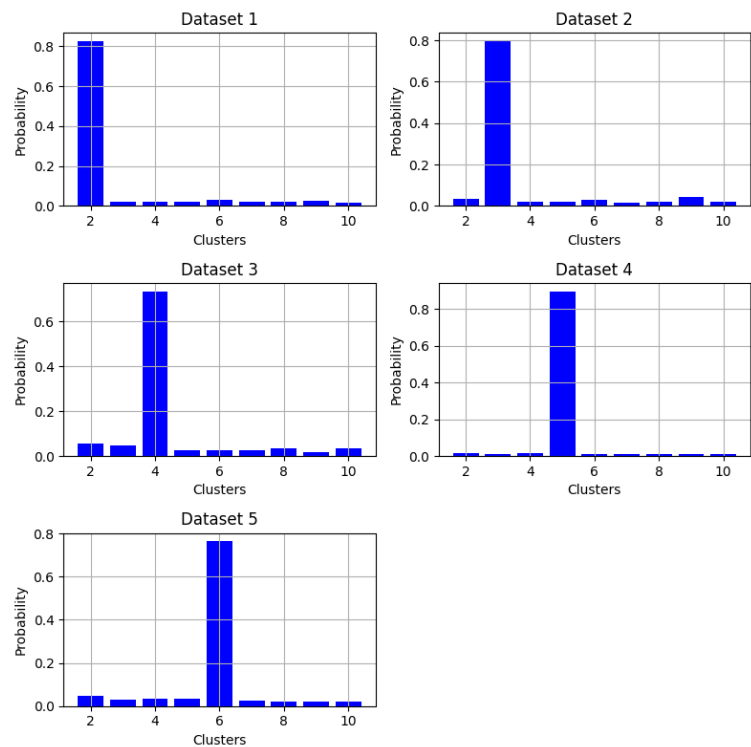


Figure 6. Validation Datasets

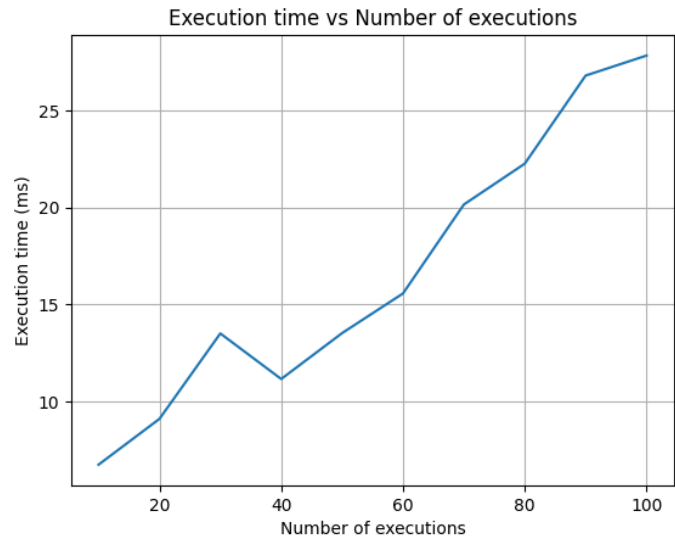
Once the validation datasets were defined, the proposed method was applied to each of them with a maximum of 10 clusters. The probability of being an optimal cluster within each dataset was determined for each cluster, yielding the results presented in Figure 7. As shown in Figure 7, the optimal cluster identified by the model matches the correct optimal cluster for each dataset provided by the AI, with each case achieving a probability above 0.6, while the other clusters did not exceed 0.1.





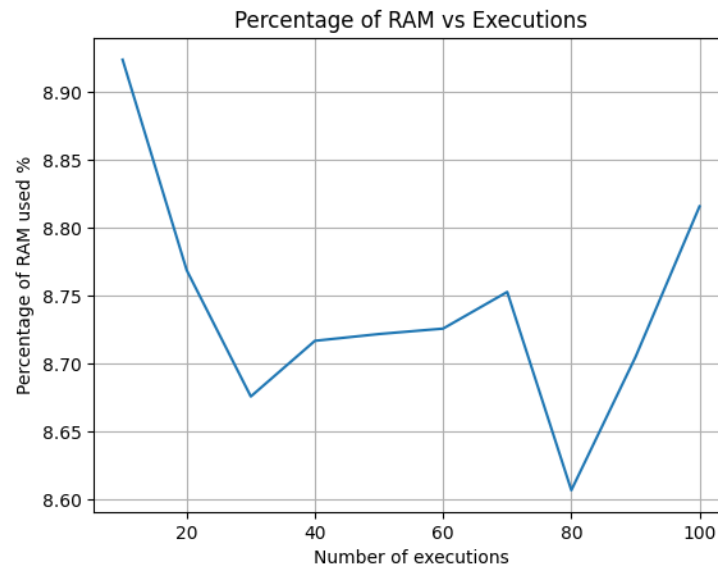
**Figure 7.** Results obtained in the validation **with** the test datasets.

Finally, to evaluate the computational efficiency of the model, the processing time of the computational method was measured by executing it from 10 to 100 runs in the Google Colab environment (with 12.67 GB of available RAM). The resulting graph is presented in Figure 8.



**Figure 8.** Executions time vs Number of executions

From Figure 8, it can be observed that in the generic environment of Google Colab, for every additional 3 executions, the method execution time increases by approximately 1 millisecond. Additionally, when measuring the percentage of RAM usage after performing invocations in increments of 10 up to 100, it is found that the percentage of RAM remains between 8.6% and 8.9% (see Figure 9). This allows us to conclude that the method for calculating the optimal number of clusters in KMeans does not require a significant computational load and can be executed without difficulty in generic environments such as those provided by the Google Colab platform.



**Figure 9.** Percentage of RAM vs Number of executions

As a discussion point, it is important to mention that the proposal presented in this article contributes to the issue addressed in [25] and [26], where it is noted that the visual inspection method of the elbow is not always an effective solution for identifying the optimal number of clusters, especially in datasets lacking clear group divisions. In this regard, the proposed method allows determining the probability that a number of clusters is optimal for a particular dataset, based on the use of numerical derivatives and the relationship between contiguous numerical derivatives. Thus, this proposal contributes to reducing subjectivity in calculating the optimal number of clusters compared to the approaches presented in [29,30], where the elbow method is used for obtaining the optimal number of clusters with datasets from political and marketing contexts.

#### 4. Conclusions

One of the most widely used approaches in machine learning is unsupervised learning, which begins with an unlabeled dataset to identify relationships between instances and determine possible categories for grouping the data. In this context, one of the most popular methods in unsupervised learning is KMeans, where it is necessary to specify the number of clusters to group the data as an input parameter. Identifying the optimal number of clusters is a significant challenge. This article proposes a mathematical method that allows for a more precise determination of the optimal number of clusters compared to the visual inspection method of the elbow, which is inaccurate when the dataset does not present well-defined groups.

The method proposed in this article is based on the numerical derivative of the inertia for each cluster count and the ratio between contiguous derivatives, such that from the maximum ratio between these derivatives, it is possible to determine the optimal number of clusters. For the implementation of this method, open-source tools and/or libraries such as scikit-learn, numpy, and matplotlib were utilized, which proved to be suitable for obtaining cluster inertia, determining numerical derivatives, obtaining the maximum ratio between contiguous derivatives, and generating the corresponding graphs for clusters and the elbow method.

The proposed method was validated using 5 datasets of 100 instances, each containing well-defined groups (2, 3, 4, 5, and 6 clusters), generated by AI. The evaluation conducted enabled the determination that the optimal clusters calculated by the proposed method coincide with the groupings observed in each dataset. As an additional contribution, the method not only determines the optimal cluster but also the probability associated with each cluster being optimal.

Regarding efficiency evaluation, the conducted tests led to the conclusion that in a generic environment such as Google Colab, for every 3 method executions, the processing time increases by 1 millisecond, implying that the proposed method uses an average of around 0.3 milliseconds per

execution. Similarly, when conducting a varying number of executions incremented by 10 up to 100, it was observed that the percentage of RAM usage remained between 8.6% and 8.9%. This indicates that the proposed method does not require significant computational overhead and can be executed without difficulty in generic environments like those provided by the Google Colab platform.

As a future work stemming from the present research, the first step is to implement a cloud service for identifying the optimal number of clusters in KMeans, based on the proposed computational method. Similarly, there is an intention to extrapolate the proposed method to the domain of hierarchical clustering, thereby providing an alternative to the visual dendrogram method.

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