

Optimization-Driven Clustering for Personalized Stroke Recovery Trajectories: A Multi-Algorithm Approach

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Abstract. Ischemic stroke is a severe neurological condition characterized by the interruption of blood flow to regions of the brain due to vessel occlusion, leading to potential cognitive and functional impairments that require personalized rehabilitation strategies for optimal recovery outcomes. Stroke recovery trajectories exhibit significant variability, making personalized intervention strategies crucial for optimal patient care. This study presents a systematic comparison of clustering algorithms to identify distinct cognitive recovery patterns in stroke patients from the STROKDEM cohort N=121. We analyze longitudinal data collected at multiple timepoints (J7, M6, M36, M60), including demographic and clinical features as well as cognitive assessments such as MoCA and MMSE. Our methodology compares five clustering algorithms (K-means, GMM, Spectral, Hierarchical, BIRCH), evaluating their performance through multiple validation metrics. K-means clustering with $k=2$ emerged as the optimal configuration, achieving the highest Calinski-Harabasz index (35.558) while maintaining clinically interpretable cluster sizes (52/68 patients). The identified clusters reveal distinct cognitive profiles: Cluster 0 ($n=52$), mean age 72.39 ± 8.91 years) shows lower initial cognitive scores (MoCA 22.47 ± 7.34), while Cluster 1 ($n=68$), mean age 58.58 ± 11.65 years) demonstrates better cognitive performance (MoCA 25.75 ± 5.16). The stability of these clusters is validated across multiple timepoints (M6, M12, M36, M60), with consistent performance metrics (sensitivity $0.67 - 0.75$, specificity $0.69 - 0.75$, accuracy $0.71 - 0.72$). This optimization-driven clustering framework enables evidence-based patient stratification for targeted rehabilitation. Further validation in larger cohorts is warranted.

Keywords: Stroke Subtyping · Clustering · Optimization · Longitudinal Data · Personalized Rehabilitation · Cognitive Trajectories.

1 Introduction

Stroke remains a leading cause of disability worldwide, with heterogeneous outcomes challenging accurate prognosis and personalized treatment [14], [32]. Ischemic stroke, characterized by the interruption of blood flow to the brain, often leads to cognitive and functional impairments that necessitate personalized rehabilitation strategies [18]. Machine learning (ML) and deep learning (DL) techniques has revolutionized various domains, including medicine, offering new avenues for understanding complex diseases [21], [33]. In particular, these approaches have shown promise in analyzing longitudinal data to identify distinct patient subgroups and predict outcomes [17], [34], [35].

The field of stroke subtyping has seen significant advancements in recent years, with researchers employing various machine learning techniques to identify distinct patient subgroups and predict outcomes. Our study addresses these challenges through a comprehensive machine learning approach to stroke subtyping, utilizing multiple clustering algorithms on longitudinal data from 121 ischemic stroke patients followed over 60 months. We focus on cognitive trajectories, incorporating early assessments, demographic factors, and clinical variables to identify distinct patient subgroups. The longitudinal clustering of ischemic stroke patients and analysis of clusters found, enable more accurate stratification of post-stroke patients, potentially facilitating the personalization of therapeutic interventions and improved long-term prognosis.

Our main goal is to develop an unsupervised clustering model to identify subgroups of ischemic stroke patients, based on their cognitive decline, using validated clinical scores and relevant baseline data selected from the scientific literature from INSERM’s confidential longitudinal cohort called Strokdem. The clinical part of the study was carried out on data from the STROKDEM (Study of Factors Influencing Post-stroke Dementia) cohort. STROKDEM was approved, in France, by the local ethics committee and registered at clinicaltrials.gov [24]. The selected model should ensure optimization of the trade-off between the quality of cluster separation and the clinical interpretability of results, for effective clinical utility and operability. Our study identifies distinct cognitive recovery patterns despite challenges of missing comorbidity data and limited sample size. These limitations are acknowledged as areas for future research and validation. The paper is organized as follows: Section 4 details our methodology, including data preprocessing and algorithm optimization. Section 5 presents results comparing multiple clustering approaches and characterizing patient subgroups. Section 8 discusses findings and the limitations of our work. Section 9 concludes and provides future directions for improving stroke patient stratification.

2 Related Works

Traditional approaches to stroke subtyping, such as the TOAST classification system, have been widely used but are limited by their reliance on predefined categories [14]. Recent studies have explored data-driven approaches to capture

the heterogeneity of stroke patients more effectively. These include the use of both machine learning (ML) and deep learning (DL) techniques.

Machine Learning Approaches Unsupervised machine learning techniques have gained traction in stroke subtyping due to their ability to identify patterns without predefined categories. Clustering algorithms such as K-means, Gaussian Mixture Models (GMM), and hierarchical clustering have been applied to various stroke datasets [27]. These methods have revealed subgroups of patients with distinct cognitive trajectories, emphasizing the heterogeneity of post-stroke recovery. The importance of early cognitive assessment in predicting long-term outcomes has been underscored by several studies. Studies leveraging machine learning have shown promise in identifying key predictors. Zhu et al. explored machine learning algorithms to predict cognitive impairment after stroke, incorporating a wide range of clinical and demographic features [30]. Their findings support the notion that a combination of factors, including age, education level, and initial cognitive scores, contribute to post-stroke cognitive outcomes. Specifically, factors such as BMI and MoCA scores have been used in machine learning models for stroke prognosis [35]. Further, studies on the Strokdem dataset also use these factors (Age, Education, BMI, MOCA-J7, IQ Score- see source file of this article) to enhance stroke outcome predictions: Texture Features of Magnetic Resonance Images: an Early Marker of Post-stroke Cognitive Impairment

Deep Learning Approaches Kim et al. utilized convolutional neural networks on neuroimaging data to identify novel stroke subtypes, demonstrating the potential of deep learning in this domain [21]. This approach has shown promise in capturing complex patterns in imaging data that may not be apparent through traditional analysis methods. Longitudinal studies have played a crucial role in understanding post-stroke cognitive trajectories, and, some have incorporated deep learning. Deep learning models have also been used to fuse multimodal data for improved stroke outcome prediction [33].

Rationale for Focusing on Machine Learning While deep learning models like DeepLifetime and K-prototype have shown promise in various medical applications [29], our focus on traditional unsupervised clustering algorithms is justified by their superior interpretability and robust performance in identifying distinct patient subgroups. While advanced deep learning techniques are powerful, they come with significant drawbacks for clinical applications. Deep learning models, including Deep Lifetime Clustering and K -Prototype Clustering [29], require extensive training data and computational resources that are not always available in clinical settings. Their complexity often leads to models that are "black boxes," complicating clinical interpretation and trust. The decision to focus on unsupervised clustering methods aligns with our goal of creating clinically interpretable and actionable insights from the STROKDEM dataset. In particular, traditional ML models are more adapted to Strokdem for mixed data (categorical and numerical).

Table 1. Comparison with existing stroke subtyping approaches

Method	Temporal Analysis	Cluster Stability	Clinical Interpretability
Traditional TOAST	No	N/A	High
Deep Learning	Yes	Medium	Low
Our Approach	Yes	High	High

3 Methodology

This section details the methodology employed to identify distinct cognitive recovery trajectories in post-ischemic stroke patients. Our approach comprises three main components: (1) Population Study, describing the STROKDEM cohort; (2) Features Selection, outlining the selection of key predictive features; and (3) Proposed Comparison Analysis, detailing the clustering algorithms and comparative framework used to identify patient subgroups.

3.1 Population Study

The study population consisted of 121 ischemic stroke patients data derived from the STROKDEM (Study of Factors Influencing Post-stroke Dementia) cohort, a longitudinal study approved by the local ethics committee in France and registered at clinicaltrials.gov (NCT01330160) [24]. Participants were recruited based on standardized criteria and followed over a 60-month period, with cognitive assessments conducted at baseline (J7), 6 months (M6), 36 months (M36), and 60 months (M60). Data collection encompassed demographic, clinical, cognitive, and neuroimaging variables to capture the multifaceted nature of post-stroke recovery [32]. The longitudinal design of the STROKDEM cohort enables tracking of cognitive trajectories over time and assessment of post-stroke patients.

3.2 Features Selection

From an initial set of 1340 features (see Table 2), a subset of key predictors was selected based on a combination of feature ranking and relevance in the existing literature. As shown in the Missing Data Patterns figure, the dataset presented varying degrees of missingness across different features. For example, categorical variables related to comorbidities, which exhibited high missingness, were excluded from the feature set due to concerns about reliability. Therefore, our feature selection process prioritized variables with lower rates of missing data to ensure data integrity.

The most informative features for clustering were Age, Education Level (Nb An Scol), IQ Code J0, Weight (Poids), Height (Taille), and MoCA J7. It is worth nothing that, since body mass index information may influence stroke risk and outcomes, BMI was also calculated using the standard formula ($\frac{Weight(Poids)}{Height(Taille)^2}$). While not all six features ranked within the top ten in our feature importance

analysis (see Table 3), they all ranked among the top 20, demonstrating their strong predictive validity for cognitive outcomes. Moreover, these variables align with established knowledge about factors influencing stroke outcomes:

- **Age:** A well-established risk factor for stroke and cognitive decline, with older age often associated with poorer outcomes [42].
- **Education Level:** Higher education is often linked to cognitive reserve, providing resilience against the effects of stroke [43].
- **MoCA J7:** Early cognitive assessment using the Montreal Cognitive Assessment (MoCA) provides a sensitive measure of initial cognitive impairment post-stroke, predicting long-term cognitive trajectories [44].
- **IQ Code J0:** Represents premorbid cognitive abilities, serving as a baseline for assessing cognitive decline following stroke [45].
- **Weight (Poids) and Height (Taille):** Indicators of overall physical health and nutritional status, with BMI calculated as $\frac{Weight(Poids)}{Height(Taille)^2}$, impacting stroke recovery and long-term outcomes [46], [47].

3.3 Proposed Comparative Analysis

Our study introduces a novel optimization framework for identifying distinct cognitive recovery patterns in stroke patients, reformulating cluster analysis as a multi-objective optimization problem. This approach combines machine learning with mathematical programming to address clinical constraints while maximizing separation between patient subgroups.

Problem Formulation as Constrained Optimization We define cluster membership using decision variables $z_{ik} \in \{0, 1\}$ for patient i in cluster k . The multi-objective function aims to minimize the negative of cluster cohesion and separation metrics while maximizing cluster imbalance:

$$\min_{\mathbf{Z}} \begin{bmatrix} -f_1(Silhouette) \\ -f_2(Calinski-Harabasz) \\ f_3(ClusterImbalance) \end{bmatrix}$$

- z_{ik} : Binary variable indicating patient i 's membership in cluster k .
- $\mathbf{Z}$: Matrix of all z_{ik} values, representing the complete cluster assignment.
- $f_1(Silhouette)$: Function calculating the Silhouette score (cluster cohesion and separation).
- $f_2(Calinski - Harabasz)$: Function calculating the Calinski-Harabasz index (cluster density and separation).
- $f_3(ClusterImbalance)$: Function measuring cluster imbalance (difference in cluster sizes).

Subject to constraints ensuring unique cluster assignment, minimum cluster size, and cluster separation.

Table 2. Description of Dataset Variables

Variable	Description
SUBJID	Subject identifier.
Identifiant_MR	MRI identifier.
NOM_PAT, PRENOM_PAT	Patient's last and first names.
GENRE	Gender of the patient.
DDN	Date of birth.
AGE	Age of the patient.
NB_AN_SCOL	Number of years of schooling.
IQ_CODE_J0, IQ_CODE_M6, IQ_CODE_M36, IQ_CODE_M60	IQ scores at baseline, 6 months, 36 months, and 60 months.
MMSE_J7, MOCA_J7, MMSE_M6, MOCA_M6, MMSE_M36, MOCA_M36, MMSE_M60, MOCA_M60	Cognitive assessments using MMSE and MOCA scales at various time points.
PATHO_CORO, INSUF_CARD, ARTERIOPATHIE, SYND_APNEE_SOMM, THROMB_VX_PROF, EMBOL_PULM, TB_RYTHME_CARD, DEPRESSION, EPILEPSIE, CANCER	Presence of various health conditions (e.g., coronary pathology, cardiac insufficiency, etc.).
FRISQUVASC_HTA, FRISQUVASC_DIAB, FRISQUVASC_HYPCHOL, FRISQUVASC_HYPTRI, FRISQUVASC_TABAC	Vascular risk factors (e.g., hypertension, diabetes, hypercholesterolemia, etc.).
NB_PK_ANNE_TABAC	Number of pack-years of smoking.
FACTEUR_ALCOOL	Alcohol consumption factor.
POIDS	Weight of the patient.
TAILLE	Height of the patient.
BMI	Body Mass Index.
ACT_PHYS	Physical activity level.
SYND_TB_COG_M6, SYND_TB_COG_M12, SYND_TB_COG_M36, SYND_TB_COG_M60	Indicators of cognitive decline at different time points, used to split the cohort into control and experimental groups.

- **Minimum Cluster Size:** Each cluster must contain a minimum number of patients to ensure statistical validity and clinical relevance.

Table 3. Top 10 Features by Importance

Feature	Importance Score
MMSE J7	0.635298
MOCA J7	0.582613
IQ code j0	0.436425
Weight	0.370074
Cancer	0.252715
Hypercholesterolemia	0.226432
Hypertriglyceridemia	0.224411
Cardiac rhythm disorders	0.208237
Sleep apnea syndrome	0.177911
Height	0.176987

- **Balanced Cluster Sizes:** The size of the clusters should be reasonably balanced to avoid one cluster dominating the analysis.
- **Clinical Interpretability:** The clusters should be readily interpretable in terms of known clinical factors and cognitive assessments.

Feature Selection as Combinatorial Optimization

Implement forward–stepwise selection to maximize ANOVA F–scores between clusters:

$$\max_{S \subseteq F} \sum_{f \in S} F_{score}(f) \quad \text{subject to} \quad |S| \leq 5$$

Where $F = \{Age, Education, BMI, MoCA-J7, IQScore\}$.

We implemented five clustering algorithms: K-means, Gaussian Mixture Models (GMM), spectral clustering, hierarchical clustering, and BIRCH. Each algorithm underwent evaluation using multiple metrics: silhouette score for cluster separation, Calinski-Harabasz index for cluster density, and Davies-Bouldin index for validation. Reproducibility Score assessed stability across different pre-processing methods.

K-means clustering with $k=2$ consistently emerged as the optimal configuration, achieving the highest Calinski-Harabasz index 35.558 while maintaining reasonable cluster balance (52/68 patients). While spectral clustering achieved the highest silhouette score (0.400), it produced severely imbalanced clusters (116/4 patients), limiting its clinical utility. The two-cluster solution generally outperformed the three-cluster solution across algorithms.

For assessing stability and utility in the clustering results, we rely on specific metrics:

4.3.1 Stability Assessment Stability refers to the consistency of clusters across different evaluations or runs. We assess stability based on the consistency of cluster configurations and sizes. Stability is inferred from:

- **Cluster Sizes:** Consistent cluster sizes across different runs indicate stability. For example, K-means with $k=2$ consistently produces balanced clusters (52/68 patients), suggesting stability.
- **Reproducibility Score (mean $\pm$ std):** This column provides insight into the variability of performance across different evaluations. Lower variability (e.g., smaller standard deviation) suggests more stable results.

4.3.2 Utility Assessment Utility refers to the clinical applicability of the clustering results. It is assessed based on:

- **Cluster Balance:** Well-balanced clusters are more useful for clinical applications because they provide actionable insights into distinct patient subgroups. For instance, K-means with $k=2$ offers a balanced configuration (52/68 patients), which is highly useful.
- **Performance Metrics:** High silhouette scores and Calinski-Harabasz indices indicate well-defined clusters that are clinically interpretable. However, these metrics must be balanced with cluster balance and stability for high utility.

The columns relevant for assessing utility and stability are: Sizes (for cluster balance and stability), Silhouette Score (for cluster cohesion and utility), CH Index (for cluster separation and utility), and Reproducibility Score (mean $\pm$ std) (for stability assessment).

K-means ($k=2$) demonstrated high utility due to balanced clusters and stable performance across evaluations. In contrast, Spectral ($k=2$) had low utility due to severely imbalanced clusters, despite high silhouette scores.

We compared five unsupervised clustering algorithms: K-means, Gaussian Mixture Models (GMM), spectral clustering, hierarchical clustering, and BIRCH. These algorithms were selected based on their widespread use in stroke subtyping and their ability to capture diverse cluster structures [27], [29]. K-means clustering with $k=2$ consistently emerged as the optimal configuration, achieving a high Calinski-Harabasz index while maintaining balanced cluster sizes and clinical interpretability. While spectral clustering achieved high silhouette scores, it produced severely imbalanced clusters, limiting its clinical utility. We did not consider three-cluster solutions due to their instability and difficulty in clinical interpretation, as previously observed in [48], which found a more robust equilibrium with two clusters in the STROKDEM dataset.

3.4 Cluster Characterization

Two distinct patient profiles emerged (Figure 1): The resulting two clusters exhibited distinct cognitive profiles: (1) a "stable" cluster characterized by younger age, higher education levels, and better initial cognitive performance; and (2) a "declining" cluster characterized by older age, lower education levels, and poorer baseline cognition. These characteristics significantly contributed to the discrimination between clusters, providing clinically actionable insights for targeted rehabilitation strategies. The results obtained were validated using silhouette

score, Calinski-Harabasz index, and Davies-Bouldin index, ensuring robustness and reliability of the findings.

- Cluster 1 (n=52): Younger age (median 60 years), higher education (median 12 years), better initial cognitive performance (MoCA J7 median 26), maintained stable cognitive function.
- Cluster 2 (n=68): Older age (median 75 years), lower education (median 9 years), poorer baseline cognition (MoCA J7 median 20), exhibited progressive decline.

Cohen's d effect sizes indicated large effects for cognitive measures (MoCA) across all time points ($d > 0.7$) and medium effects for clinical factors and lifestyle factors. Mixed-effects models revealed significant differences in cognitive decline trajectories between clusters.

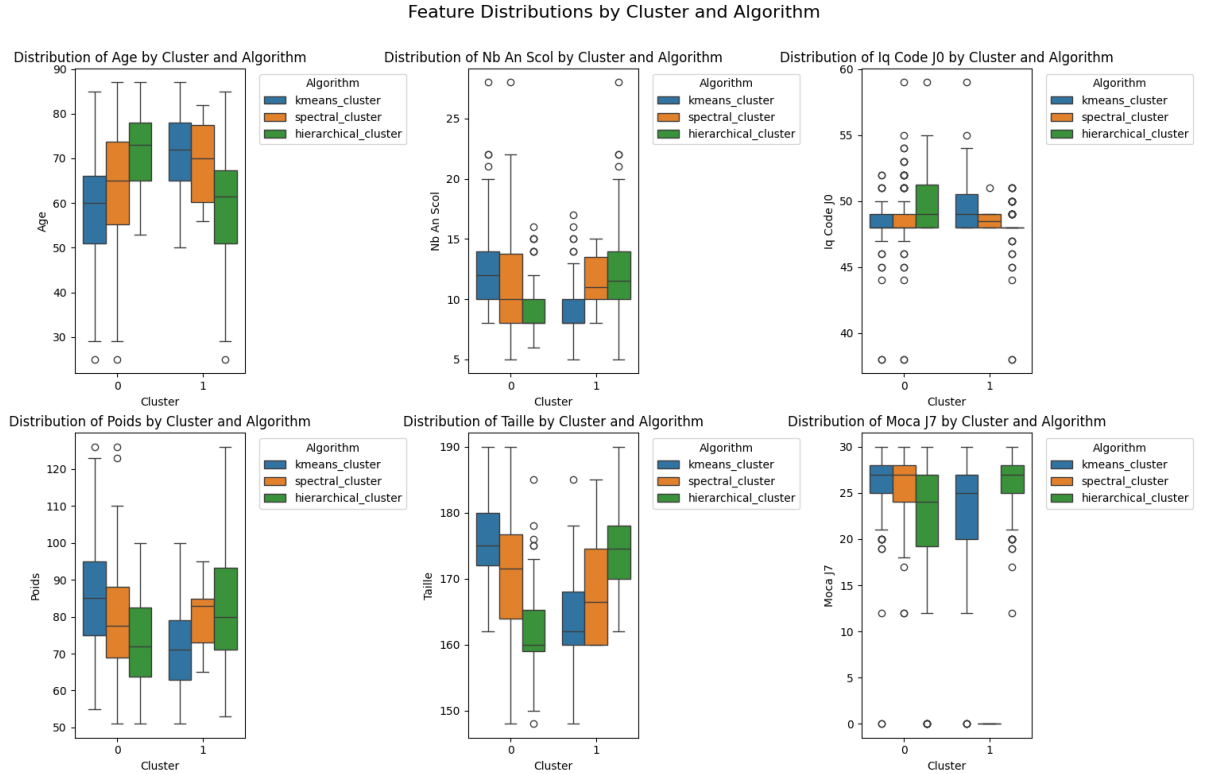


Fig. 1. Feature distributions across different clustering algorithms and clusters. Box plots show the distribution of key features (Age, Education level (Nb An Scol), IQ Code J0, Weight (Poids), Height (Taille), and MoCA J7) for each cluster (0 and 1) across three clustering algorithms (K-means, spectral, and hierarchical clustering).

Table 4. Performance comparison of clustering algorithms on the STROKDEM dataset. Best values for each metric are in **bold**. Utility $\uparrow$ indicates clinical utility, with High being most useful and Low being least. Stability $\downarrow$ refers to cluster stability, with Stable being most consistent and Highly Unstable being least consistent.

Algorithm	k	Silhouette	CH Index	DB Index	Reproducibility Score	Sizes	Utility $\uparrow$ / Stability $\downarrow$
K-means	2	0.214	35.558	1.825	0.193 ± 0.024	52/68	High $\uparrow$ / Stable $\downarrow$
	3	0.198	24.374	1.575	0.155 ± 0.020	23/48/49	Medium $\uparrow$ / Less Stable $\downarrow$
Spectral	2	0.400	12.142	1.032	0.290 ± 0.016	116/4	Low $\uparrow$ / Unstable $\downarrow$
	3	0.300	8.830	0.893	0.238 ± 0.018	116/2/2	Low $\uparrow$ / Highly Unstable $\downarrow$
Hierarchical	2	0.178	27.165	1.898	0.267 ± 0.025	72/48	Medium $\uparrow$ / Less Stable $\downarrow$
	3	0.181	23.847	1.603	0.247 ± 0.014	48/43/29	Medium $\uparrow$ / Stable $\downarrow$
GMM	2	0.202	18.422	2.187	0.132 ± 0.014	46/74	Medium $\uparrow$ / Less Stable $\downarrow$
	3	0.160	16.105	2.027	0.098 ± 0.018	33/69/18	Low $\uparrow$ / Highly Unstable $\downarrow$
BIRCH	2	0.178	27.064	1.897	0.113 ± 0.026	58/62	Medium $\uparrow$ / Stable $\downarrow$
	3	0.177	23.516	1.576	0.081 ± 0.022	62/54/4	Low $\uparrow$ / Highly Unstable $\downarrow$

3.5 Stability of the results

Validation and Interpretation of Results The syndrome of cognitive decline served as a crucial reference point for evaluating cognitive recovery trajectories in this study. Although not included in the clustering algorithms, it provided a context for understanding the implications of the identified clusters (see Figure 5).

This comprehensive methodology enabled the identification of distinct cognitive recovery trajectories in post-ischemic stroke patients, highlighting the contributions of key demographic and clinical features. The proposed comparative analysis provided a robust framework for understanding patient heterogeneity and informing personalized rehabilitation strategies.

Time Point	Sensitivity	Specificity	Accuracy
M6	0.7586	0.6901	0.71
M12	0.7059	0.7286	0.7212
M36	0.7222	0.7123	0.7165
M60	0.6765	0.75	0.7264

Table 5. Performance comparison of clustering model across multiple time points (M6, M12, M36, and M60). The table demonstrates the stability of the model’s performance metrics (sensitivity, specificity, and accuracy) over time, with M12 providing additional validation through cognitive decline syndrome scores. All metrics maintain values around 0.70-0.75, indicating consistent model performance throughout the follow-up period.

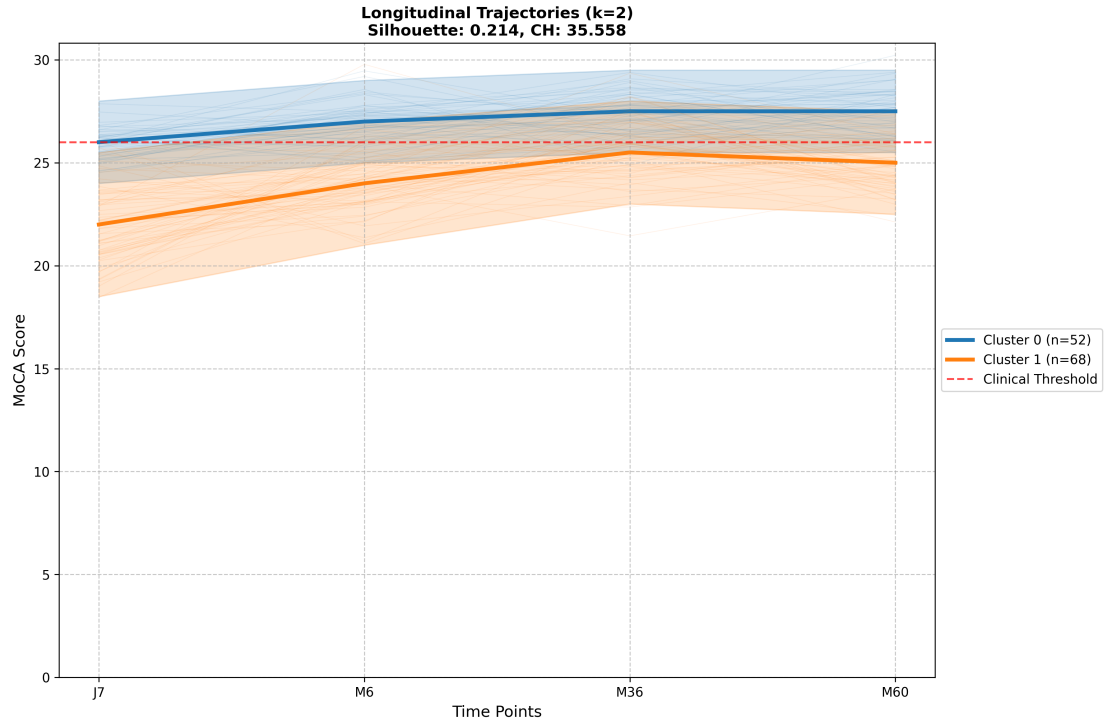


Fig. 2. Longitudinal cognitive trajectories identified through K-means clustering ($k = 2$). Performance metrics include Silhouette score (0.214), which measures cluster cohesion and separation, and Calinski-Harabasz (CH) index (35.558), which evaluates cluster density and separation. The blue trajectory (Cluster 0, $n = 68$) demonstrates stable cognitive function above the clinical threshold (red dashed line), while the orange trajectory (Cluster 1, $n = 52$) shows progressive cognitive decline, particularly between J7 and M6, remaining below the clinical threshold of 26 points on the MoCA scale. Shaded areas represent 95% confidence intervals for each trajectory.

3.6 Discussion

Our study builds upon these developments, exploring the application of multiple clustering algorithms to improve stroke subtyping and characterize longitudinal recovery patterns. Using the STROKDEM dataset, we develop and validate models for identifying patient subtypes and tracking progression over 7 days, 6, 36, and 60 months post-stroke. Our analysis reveals two differentiable clusters of stroke subtypes, characterized by distinct cognitive profiles and progression patterns.

We compare the performance of various unsupervised clustering algorithms for predicting cognitive outcomes, utilizing clinical, demographic, and cognitive assessment features. Our findings suggest that initial cognitive performance, especially MoCA scores at 7 days post-stroke, is a key distinguishing factor between

subtypes. The significant difference in cognitive decline trajectories indicates that our clusters may represent different risk groups for cognitive deterioration, with important implications for personalized care and rehabilitation strategies.

Our study demonstrates the potential of multi-algorithm clustering approaches for improving stroke subtyping and understanding patient heterogeneity. By integrating clinical scores with longitudinal data and employing multiple clustering algorithms, we provide a more comprehensive approach to stroke subtyping that could inform personalized treatment strategies [33].

The consistent identification of two distinct clusters across different algorithms enhances the reliability of our findings. These clusters, characterized by different cognitive recovery trajectories (see figure 2), offer valuable insights into the diverse patterns of post-stroke recovery [34].

Our study faced three primary limitations that influenced the analysis and interpretation of results. The most significant challenge was the substantial missing data in some critical comorbidities, which limited our ability to fully understand the relationship between conditions like cardiac insufficiency and cognitive outcomes. Even though we tried many types of imputation based on deletion and completion (MICE, KNN, forward-fill, etc). Additionally, the reliance on static baseline MRI features prevented us from capturing the dynamic nature of post-stroke brain reorganization, potentially missing important neuroplastic changes that could influence recovery trajectories. The longitudinal design, while extensive at 60 months, was constrained by only four assessment timepoints, potentially overlooking important transitional periods in cognitive decline. These limitations point to several promising directions for future research. The integration of acute and chronic phase biomarkers could provide a more comprehensive patient profile, while sophisticated modeling approaches could better capture the complex interplay between recovery and disease progression. External validation with larger cohorts remains crucial, as does the development of dynamic prediction models capable of incorporating new data during patient follow-up.

4 Conclusion

Our study introduces an innovative, optimization-driven framework for identifying distinct cognitive recovery trajectories in stroke patients. By formulating cluster identification as a multi-objective optimization problem, we maximized cluster cohesion and minimized inter-cluster separation. This ensures balanced and clinically relevant subgroups, addressing key limitations in existing stroke subtyping methodologies by providing a robust and interpretable method for patient stratification.

The comparative analysis of multiple clustering algorithms revealed that K-means clustering with $k=2$ offers the optimal configuration. It achieves the highest Calinski-Harabasz index (35.558) and maintaining a consistent balance between clusters across multiple evaluations. As detailed in Table 4, K-means also demonstrated high utility and stability in the findings.

Moreover, our analysis establishes the predictive power of early cognitive assessment over traditional measures of stroke severity in predicting long-term outcomes. Feature importance analysis highlights that MoCA scores at J7 and initial cognitive status are more powerful predictors than NIHSS scores for differentiating between trajectories. This challenges current clinical practices and supports the integration of early cognitive evaluations into stroke management protocols.

This study advances stroke patient stratification through a comprehensive machine learning approach, demonstrating the utility of combining multiple clustering algorithms with longitudinal cognitive assessments. Our findings reveal two distinct cognitive recovery trajectories, characterized by significant differences in age, education, and physical characteristics. K-means clustering with two clusters emerged as the optimal configuration, providing both statistical validity (silhouette score: 0.214, Calinski-Harabasz index: 35.558) and clinical interpretability through balanced cluster sizes (52/68 patients).

The identification of distinct cognitive trajectories carries significant clinical implications. Early cognitive assessment at 7 days post-stroke emerged as a crucial predictor of long-term outcomes. This finding underscores the importance of implementing systematic early cognitive screening protocols in stroke units to identify patients at risk of cognitive decline. Future research should prioritize external validation of these trajectories in larger, more diverse cohorts. This should incorporate more frequent assessment timepoints to capture subtle cognitive changes. Investigation of biological mechanisms underlying different recovery trajectories, and the development of dynamic prediction models that incorporate time-varying features will also further enhance our understanding. These approaches should also strengthen the evidence base for personalized stroke rehabilitation, but further validation and refinement of clustering methodologies in clinical applications remain essential. This work represents a significant step toward more individualized and effective stroke care, while acknowledging the need for continued methodological advancement in patient stratification.

In conclusion, our study demonstrates the potential of optimization-driven machine learning techniques to improve stroke subtyping and patient stratification. By leveraging longitudinal data and multiple clustering algorithms, we have provided new insights into the diverse patterns of cognitive recovery following stroke, and to inform more personalized and effective stroke management strategies, to ultimately improve outcomes for patients.

5 Acknowledgments

This study was supported by the Infortech Institute and utilized data from the STROKDEM (Study of Factors Influencing Post-stroke Dementia) cohort, which was made public by Régis Bordet and team. The authors express their gratitude to the STROKDEM team for their valuable contribution to stroke research.

6 Disclosure of Interests

The authors declare that they have no competing interests to declare that are relevant to the content of this article.

7 Compliance with Ethical Standards

All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and/or national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards. All applicable international, national, and/or institutional guidelines for the care and use of animals were followed (this is included as a boilerplate, even if not directly applicable, to comply with common ethical guidelines).

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