

A Unified Framework with Difference of Convex Decomposition and Sparse Topological Learning for Skeleton-Based Clinical Assessment

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Abstract

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2 1 Introduction

3 In recent decades, the global demand for physical rehabilitation services has increased dramatically, driven by demographic changes and the rising prevalence of chronic diseases [Jesus *et al.*, 2022]. It is estimated that in 2019, more than 2.4 billion people worldwide were living with health conditions that could benefit from rehabilitation services, a significant increase since 1990. This increase is largely due to factors such as an aging population and the increasing prevalence of non-communicable diseases, including musculoskeletal disorders, which have been identified as the leading cause of disability worldwide [Chen *et al.*, 2022]. Despite this growing demand, there remains a significant gap between the supply and demand for rehabilitation services, particularly in low and middle income countries where resources are often limited. This imbalance not only affects the quality of life of those requiring rehabilitation, but also imposes a significant economic burden on health systems globally. For example, a study by Soberg, H. L. *et al.* have examined rehabilitation demand and service provision highlighted the significant costs associated with rehabilitation in the first year after injury, underscoring the financial burden on health care infrastructure [Soberg *et al.*, 2022]. The integration of artificial intelligence (AI) into physical rehabilitation offers a promising avenue to address these challenges by improving the efficiency, accessibility, and personalization of care [Calderone *et al.*, 2024]. AI-based technologies, such as machine learning algorithms, can analyze complex patient data to develop individualized treatment plans, predict recovery trajectories, and monitor patient progress in real time. For example, AI applications have been used to improve motor function in patients with neurological disorders by tailoring interventions to specific needs. The application of AI in rehabilitation also extends to administrative and operational aspects, streamlining processes such as scheduling, documentation, and resource allocation. By automating these tasks, healthcare providers can devote more time and attention to direct patient care, ultimately improving service delivery and patient satisfaction.

2 Related Works

40 Several studies have highlighted the efficiency of deep learning in this domain. In 2020, Yao *et al.* proposed a deep learning framework for assessing the quality of physical rehabilitation exercises [Liao *et al.*, 2020a]. Their approach utilized performance metrics based on the log-likelihood of a Gaussian mixture model, combined with deep neural networks to generate quality scores. In recent years, Mennella *et al.* have introduced a novel system for home-based, remote, and unsupervised rehabilitation exercise monitoring, leveraging deep learning for real-time evaluation [Mennella *et al.*, 2023]. The system focuses on two components with range of motion (ROM) classification and compensatory pattern recognition, achieving mean accuracy of 89% and 98%, respectively, with a unique dataset of six resistance training exercises. Furthermore, Zhu *et al.* proposed a multipath convolutional neural network (MP-CNN), comprising a dynamic convolutional neural network (D-CNN) and a state transition probability CNN (S-CNN) [Zhu *et al.*, 2019]. The D-CNN uses Gaussian mixture models to capture sensor data distributions, while the S-CNN extracts transition probabilities using a modified Lempel–Ziv–Welch algorithm, achieving an average accuracy of 97.8% for recognition and 96.5% for evaluation. To further increase the advancement, this work further increase the robustness for skeleton-Based rehabilitation assessment, the main contributions of this work are as follows:

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The main challenge lies in capturing the subtle biomechanical deviations that distinguish correct from compensatory movements. Traditional skeleton-based action recognition methods focus predominantly on discriminating between action categories, treating skeletal sequences as spatial-temporal graphs and applying graph convolutional networks to learn discriminative representations [Yan *et al.*, 2018b; Shi *et al.*, 2019]. However, movement quality assessment requires a fundamentally different paradigm: rather than learning categorical distinctions, we must capture fine-grained continuous variations in movement execution that correlate with clinical expertise. Existing approaches often rely solely on Euclidean geometric features, failing to capture the rich geometric structure inherent in skeletal motion. Furthermore, they typically employ single-stream architectures that cannot simultaneously model multi-scale spatial dependencies, geometric constraints, and topological relationships essential for comprehensive quality assessment.

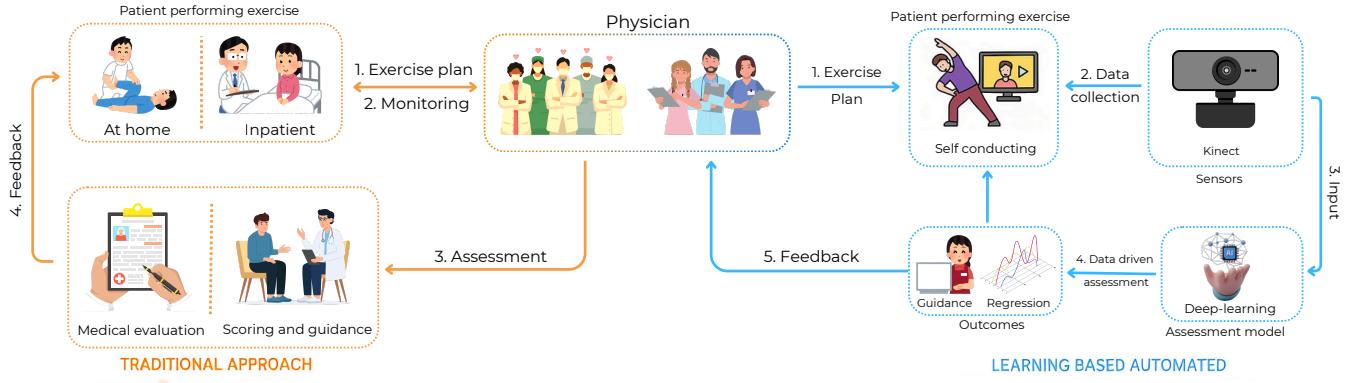


Figure 1: Enter Caption

Recent work in biomechanics and motor control theory suggests that movement quality emerges from the complex interplay of multiple factors: the geometric configuration of body segments, the differential relationships between joint positions, and the local topological structure of the kinematic chain [Bernstein, 1967]. This observation motivates our design of a multi-stream architecture that explicitly models these complementary aspects. Additionally, while graph neural networks have demonstrated success in capturing skeletal connectivity, they often treat all joints uniformly, neglecting the varying importance of different body regions in clinical assessment. Similarly, standard distance-based features employ simplistic metrics that do not exploit the underlying mathematical structure of distance functions, limiting their representational capacity.

3 Methodology

3.1 Problem Formulation and Notation

Using skeletal-based movement quality assessment in clinical rehabilitation presents a fundamental challenge in quantifying the quality of human motion based on temporal skeletal information. We begin by formally defining the skeletal representation and graph structure that underlies our approach. A human skeleton at time t is represented as a graph $\mathcal{G}_t = (\mathcal{V}, \mathcal{E}, \mathbf{F}_t)$, where $\mathcal{V} = \{v_1, v_2, \dots, v_N\}$ denotes the set of N joints, $\mathcal{E} \subseteq \mathcal{V} \times \mathcal{V}$ represents the natural skeletal connectivity defined by the human kinematic chain, and $\mathbf{F}_t \in \mathbb{R}^{N \times D}$ contains the feature vectors for all joints. Each joint v_i is associated with a feature vector $\mathbf{f}_{t,i} = [\mathbf{q}_{t,i}; \mathbf{p}_{t,i}] \in \mathbb{R}^D$, where $\mathbf{q}_{t,i} \in \mathbb{R}^4$ represents the quaternion orientation and $\mathbf{p}_{t,i} \in \mathbb{R}^3$ denotes the 3D spatial position in the coordinate system.

The adjacency matrix $\mathbf{A} \in \{0, 1\}^{N \times N}$ encodes the skeletal connectivity, where $A_{ij} = 1$ if joints v_i and v_j are connected by a bone, and $A_{ij} = 0$ otherwise. We denote the degree matrix as \mathbf{D} , where $D_{ii} = \sum_j A_{ij}$, and we define the normalized adjacency matrix following by [Kipf and Welling, 2017] as $\tilde{\mathbf{A}} = \mathbf{D}^{-1/2} \mathbf{A} \mathbf{D}^{-1/2}$. The temporal sequence of skeletons forms a spatial-temporal graph $\mathcal{G} = \{\mathcal{G}_1, \mathcal{G}_2, \dots, \mathcal{G}_T\}$. Given a temporal sequence of 3D skeleton data $\mathbf{X} = \{\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_T\} \in \mathbb{R}^{T \times N \times D}$, where T denotes the number of temporal frames, N represents the

number of skeletal joints, and D is the feature dimension encompassing both joint positions and orientations, our objective is to learn a mapping function $f : \mathbb{R}^{T \times N \times D} \rightarrow \mathbb{R}^K$ that predicts clinical assessment scores $\mathbf{y} \in \mathbb{R}^K$ reflecting movement quality across K evaluation criteria.

3.2 Overview of the Proposed Approach

We propose a novel three-stream GNN architecture that addresses the limitations of existing approaches through complementary feature extraction pathways, each designed to capture distinct aspects of movement quality. Our method, termed DC²-STG (Difference of Convex-inspired Sparse Topological Graph Convolutional Network), integrates three parallel processing streams: (i) a skeleton-geometry stream with Selective Kernel (SK) attention [Li *et al.*, 2019] for adaptive multi-scale spatial feature learning, (ii) a geometric relationship stream employing difference-of-convex decomposition [Le Thi and Pham Dinh, 2005; Le Thi and Pham Dinh, 2018] with sparse regularization inspired by DC programming principles, and (iii) a sparse topological stream that captures local neighborhood structures through multi-scale k-nearest neighbor analysis.

The three streams operate in parallel on the input skeletal sequence, extracting features at different levels of abstraction. These features are subsequently fused through a learnable fusion layer, followed by temporal modeling via Gated Recurrent Units (GRU) to capture the sequential dynamics of movement execution. The entire architecture is trained end-to-end using movement quality scores annotated by clinical experts for the target regression assessment task. The proposed DC²-STG architecture overview is illustrated in Figure ??.

3.3 Skeleton-Geometry Processing Stream

The first stream processes the raw skeletal features through a sequence of spatial-temporal graph convolution layers augmented with adaptive multi-scale feature selection. We employ Shift-GCN [Cheng *et al.*, 2020] as the fundamental building block, which extends standard graph convolution by proposed an temporal shift operations to model motion dynamics effectively without additional parameters. To enable adaptive multi-scale spatial feature extraction, we utilized SK attention [Li *et al.*, 2019] following the graph convolution

164 layers. The SK attention mechanism dynamically fuses fea-
 165 tures from multiple parallel convolution branches with dif-
 166 ferent receptive fields, allowing the network to adaptively se-
 167 lect the appropriate scale for each spatial location. Speci-
 168 cally, given M parallel convolution branches with kernel
 169 sizes $\mathcal{K} = \{k_1, k_2, \dots, k_M\}$, where we use $\mathcal{K} = \{1, 3, 5\}$,
 170 we obtain M feature representations through temporal con-
 171 volutions along the graph-structured data:

$$\mathbf{U}_m = \text{Conv}_{k_m}(\mathbf{X}^{(L)}), \quad m = 1, 2, \dots, M, \quad (1)$$

172 where $\mathbf{X}^{(L)}$ denotes the output from the final Shift-GCN
 173 layer. After that, we performed an feature fusion pro-
 174 cess through a two-stage process by compute global spatial-
 175 temporal statistics via global average pooling:

$$\mathbf{s} = \frac{1}{M} \sum_{m=1}^M \frac{1}{\tau N} \sum_{t=1}^{\tau} \sum_{i=1}^N \mathbf{U}_m(t, i, :), \quad (2)$$

176 where $\mathbf{s} \in \mathbb{R}^C$ aggregates information across all branches,
 177 time steps, and spatial locations. Subsequently, we compute
 178 branch-specific attention weights through a compact feature
 179 extraction and expansion process:

$$\mathbf{z} = \sigma(\mathbf{W}_{fc}\mathbf{s}), \quad \mathbf{a}_m = \text{softmax}_m(\mathbf{W}_m\mathbf{z}), \quad (3)$$

180 where $\mathbf{W}_{fc} \in \mathbb{R}^{d \times C}$ reduces dimensionality to $d =$
 181 $\max(C/r, L)$ with reduction ratio $r = 8$ and minimum di-
 182 mension $L = 32$, $\mathbf{W}_m \in \mathbb{R}^{C \times d}$ expands features for branch
 183 m , and the softmax operation ensures $\sum_{m=1}^M a_{m,c} = 1$
 184 for each channel c . The final output combines all branches
 185 weighted by their attention:

$$\mathbf{X}_{\text{skel}} = \sum_{m=1}^M \mathbf{a}_m \odot \mathbf{U}_m, \quad (4)$$

186 where \odot denotes element-wise multiplication with broadcast-
 187 ing. This design allows the network to emphasize features
 188 from the most informative receptive field scale for each spa-
 189 tial location and temporal instant, which is important for cap-
 190 turing both fine-grained local movements and coarse-grained
 191 global postural configurations.

3.4 DC-Inspired Geometric Relationship Stream

193 In this second stream, we propose a geometric feature ex-
 194 traction module inspired by the modeling intuition of Differ-
 195 ence of Convex (DC) programming principles by Tao Pham
 196 Dinh & Hoai An Le Thi [Le Thi and Pham Dinh, 2005;
 197 Le Thi and Pham Dinh, 2018]. Rather than implementing the
 198 full iterative DC Algorithm (DCA) [Pham Dinh and Le Thi,
 199 1997], we adopt the conceptual framework of DC decom-
 200 position for feature design and employ computationally effi-
 201 cient proximal gradient optimization [Parikh and Boyd, 2014;
 202 Beck and Teboulle, 2009] for sparse weight learning. This
 203 design is motivated by the observation that movement qual-
 204 ity assessment inherently involves geometric constraints that
 205 benefit from the mathematical structure provided by convex
 206 decomposition.

The main ideas following by the explicit geometric model-
 207 ing within the DC-inspired stream, which decomposes pair-
 208 wise joint distances into complementary convex and con-
 209 cave components and applies proximal gradient optimiza-
 210 tion [Parikh and Boyd, 2014; Beck and Teboulle, 2009] with
 211 soft thresholding to learn sparse importance weights. By de-
 212 composing distance metrics into squared Euclidean and log-
 213 arithmetic components-functions with complementary growth
 214 properties-we enable the model to capture both local fine-
 215 grained deviations and global coarse-grained relationships si-
 216 multaneously. This dual representation is particularly well-
 217 suited for clinical movement quality assessment, which fun-
 218 damentally relies on geometric constraints at multiple scales.
 219

Difference of Convex Inspire Feature Decomposition

Giving 3D joint positions of \mathbf{X} , the geometric rela-
 220 tionships between joints are fundamentally characterized by pair-
 221 wise distances. Inspired by DC programming [Le Thi and
 222 Pham Dinh, 2005], we decompose geometric features into
 223 two complementary functions that capture distinct mathemat-
 224 ical properties of spatial relationships.
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Definition 1 (DC-Inspired Feature Space). Let $\mathcal{P} =$
 $\{(\mathbf{p}_i, \mathbf{p}_j) : i < j, i, j \in [N]\}$ be the set of all unique joint
 227 pairs. We define the DC-inspired feature space as the Car-
 228 teian product $\Phi = \Phi_1 \times \Phi_2$, where:
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$$\Phi_1 = \{\phi_1(d) : d \in \mathbb{R}_+\}, \quad \Phi_2 = \{\phi_2(d) : d \in \mathbb{R}_+\}, \quad (5)$$

with $\phi_1(d) = d^2$ and $\phi_2(d) = \log(d + 1)$ being transforma-
 231 tions mapping distances to complementary feature represen-
 232 tations.
 233

For any pair of joints (i, j) , we define the pairwise distance
 234 as $d_{ij}^t = \|\mathbf{p}_{t,i} - \mathbf{p}_{t,j}\|_2$. We then construct two feature vectors
 235 by applying the transformations:
 236

$$\phi_1(d_{ij}^t) = (d_{ij}^t)^2, \quad \phi_2(d_{ij}^t) = \log(d_{ij}^t + 1), \quad (6)$$

where the squared distance emphasizes local geometric de-
 237 viations with quadratic sensitivity to large distances, while
 238 the logarithmic transformation provides robustness to outliers
 239 and better captures relative proportions between joint pairs
 240 through its compressive nature.
 241

Lemma 1 (Properties of DC Feature Functions). The feature
 242 functions ϕ_1 and ϕ_2 have complementary mathematical prop-
 243 erties:
 244

1. $\phi_1(d) = d^2$ is strictly convex with $\nabla^2 \phi_1(d) = 2 > 0$
2. $\phi_2(d) = \log(d + 1)$ is strictly concave with $\nabla^2 \phi_2(d) = -\frac{1}{(d+1)^2} < 0$
3. ϕ_1 exhibits superlinear growth: $\lim_{d \rightarrow \infty} \frac{\phi_1(d)}{d} = \infty$
4. ϕ_2 exhibits sublinear growth: $\lim_{d \rightarrow \infty} \frac{\phi_2(d)}{d} = 0$
5. The functions are not affinely related, providing comple-
 250 mentary geometric information

Proposition 1 (Expressiveness of DC Features). The DC-
 252 inspired feature decomposition $\{\phi_1(d), \phi_2(d)\}$ provides com-
 253 plementary information about pairwise distances. Specifi-
 254 cally, ϕ_1 and ϕ_2 are functionally independent: there exists
 255 no continuous function g such that $\phi_2(d) = g(\phi_1(d))$ for all
 256

257 $d \geq 0$. This independence ensures that the feature space
 258 $\Phi_1 \times \Phi_2$ captures geometric relationships that neither function
 259 alone can represent.

260 For computational efficiency, we consider only the upper
 261 triangular elements of the distance matrix, yielding $M = \frac{N(N-1)}{2}$
 262 unique pairwise features. At time t , we construct
 263 the DC feature vector, operated by Eq (7).

$$\psi_t = [\phi_1(d_{12}^t), \dots, \phi_1(d_{N-1,N}^t), \\ \phi_2(d_{12}^t), \dots, \phi_2(d_{N-1,N}^t)]^\top \in \mathbb{R}^{2M}, \quad (7)$$

264 representing a concatenation of squared distances and log-
 265 distances for all joint pairs. This decomposition naturally
 266 separates local and global geometric information while main-
 267 taining mathematical tractability for optimization.

Sparse Feature Weighting via Proximal Optimization

268 To learn the importance of different geometric relationships,
 269 we utilized an learnable weights $\mathbf{w} \in \mathbb{R}^{2M}$ that modu-
 270 late the DC features, the weighted feature vector becomes
 271 $\tilde{\psi}_t = \mathbf{w} \odot \psi_t$. To prevent overfitting and encourage sparsity
 272 in the learned weights-allowing the model to select only the
 273 most clinically relevant geometric relationships-we formulate
 274 the weight learning as an elastic net regularized optimization
 275 problem [Zou and Hastie, 2005]:

$$\min_{\mathbf{w}} \mathcal{L}_{\text{task}}(\mathbf{w}) + \lambda \|\mathbf{w}\|_2^2 + \mu \|\mathbf{w}\|_1, \quad (8)$$

277 where λ controls the ℓ_2 regularization strength for smooth-
 278 ness, and μ controls the ℓ_1 regularization strength for sparsity.
 279 By inspirations of DC programming [Le Thi and Pham Dinh,
 280 2005], where the objective can be viewed through a DC lens
 281 via the equivalence $\|\mathbf{w}\|_1 = \max_{\mathbf{v} \in [-1,1]^{2M}} \mathbf{v}^\top \mathbf{w}$, express-
 282 ing the non-smooth ℓ_1 norm through a max-over-convex-
 283 functions representation. Rather than implementing the full
 284 iterative DCA procedure [Pham Dinh and Le Thi, 1997],
 285 which would require solving a convex subproblem at each
 286 iteration, we employ the computationally efficient proximal
 287 gradient method [Parikh and Boyd, 2014; Beck and Teboulle,
 288 2009] with soft thresholding to retains the sparsity-inducing
 289 benefits of DC optimization while being compatible with
 290 modern DL frameworks and gradient-based training.

291 **Theorem 1** (Convergence of Proximal Gradient with Soft
 292 Thresholding). *Let $\mathcal{L}(\mathbf{w}) = \mathcal{L}_{\text{task}}(\mathbf{w}) + \lambda \|\mathbf{w}\|_2^2 + \mu \|\mathbf{w}\|_1$
 293 be the objective function with $\lambda, \mu > 0$. Assume $\mathcal{L}_{\text{task}}$ is L -
 294 smooth (i.e., $\nabla \mathcal{L}_{\text{task}}$ is L -Lipschitz continuous). The proximal
 295 gradient update:*

$$\mathbf{w}^{(k+1)} = \mathcal{S}_\theta \left((1 - 2\lambda\eta) \mathbf{w}^{(k)} - \eta \nabla \mathcal{L}_{\text{task}}(\mathbf{w}^{(k)}) \right), \quad (9)$$

296 where $\mathcal{S}_\theta(x) = \text{sign}(x) \max(|x| - \theta, 0)$ is the soft thresh-
 297 olding operator with $\theta = \frac{\mu\eta}{1+2\lambda\eta}$, converges to a stationary point
 298 of $\mathcal{L}(\mathbf{w})$ when the step size satisfies $\eta < \frac{1}{L+2\lambda}$.

299 **Corollary 1** (Sparsity Guarantee). *The soft thresholding
 300 operator \mathcal{S}_θ induces exact zeros in the weight vector \mathbf{w} .
 301 Specifically, for any coordinate i , if $|(1 - 2\lambda\eta)w_i^{(k)} -$
 302 $\eta \nabla_i \mathcal{L}_{\text{task}}(\mathbf{w}^{(k)})| < \theta$, then $w_i^{(k+1)} = 0$. The expected
 303 number of non-zero weights decreases monotonically with in-
 304 creasing μ .*

In practice, we integrate soft thresholding into the forward
 305 pass as a regularization mechanism:

$$w_i^{\text{soft}} = \text{sign}(w_i) \cdot \max \left(|w_i| - \frac{\mu}{2\lambda + \epsilon}, 0 \right), \quad (10)$$

307 where $\epsilon = 10^{-8}$ is a small constant for numerical stability.
 308 The weights \mathbf{w} are implemented as learnable parameters and
 309 are jointly optimized with other network parameters through
 310 backpropagation, with the soft thresholding providing an im-
 311 plicit sparsity-inducing regularization that selects clinically
 312 relevant geometric features.

Feature Normalization and Graph Convolution

313 After DC feature extraction and sparse weighting, we apply
 314 layer-wise feature normalization to stabilize training by
 315 $\hat{\psi}_t = \tilde{\psi}_t - \mu_\psi / \sigma_\psi + \epsilon$ where μ_ψ and σ_ψ are the mean and
 316 standard deviation computed across both batch and tempo-
 317 ral dimensions, then features are then projected to a higher-
 318 dimensional space through a MLP with layer norm by follow-
 319 ing:

$$\mathbf{h}_t^{\text{DC}} = \text{MLP}_{\text{DC}}(\hat{\psi}_t) = \mathbf{W}_2 \cdot \sigma(\text{LN}(\mathbf{W}_1 \hat{\psi}_t + \mathbf{b}_1)) + \mathbf{b}_2, \quad (11)$$

321 where $\mathbf{W}_1 \in \mathbb{R}^{2H \times 2M}$, $\mathbf{W}_2 \in \mathbb{R}^{H \times 2H}$ are learnable pro-
 322 jection matrices, $\text{LN}(\cdot)$ denotes layer normalization, and H
 323 is the hidden dimension. This produces a temporal sequence
 324 of DC features $\mathbf{H}^{\text{DC}} = \{\mathbf{h}_1^{\text{DC}}, \dots, \mathbf{h}_\tau^{\text{DC}}\} \in \mathbb{R}^{\tau \times H}$.

325 To enable graph-based spatial processing of these global
 326 geometric features, we expand the temporal features across
 327 the joint dimension and apply Shift-GCN layers by Eq (12).

$$\mathbf{X}_{\text{DC}} = \text{Shift-GCN}_2(\text{Shift-GCN}_1(\text{Expand}(\mathbf{H}^{\text{DC}}))), \quad (12)$$

328 where $\text{Expand} : \mathbb{R}^{\tau \times H} \rightarrow \mathbb{R}^{\tau \times N \times H}$ replicates the fea-
 329 tures across all N joints, creating tensors suitable for graph
 330 convolution that propagates the global geometric information
 331 through the skeletal structure.

3.5 Sparse Topological Neighborhood Stream

332 The third stream captures local topological structures by an-
 333 alyzing k-Nearest Neighbor (k-NN) relationships in the po-
 334 sitional space. Unlike the skeleton connectivity \mathcal{E} defined by
 335 anatomical constraints, k-NN topology reflects the actual spa-
 336 tial configuration and can reveal abnormal postures or com-
 337 pensatory movements where joints that are typically distant
 338 become spatially proximate [Bronstein *et al.*, 2017].

Multi-scale k-NN Feature Extraction

340 For each temporal frame t , we compute the pairwise Eu-
 341 clidean distance matrix $\mathbf{D}_t \in \mathbb{R}^{N \times N}$ between all joint po-
 342 sitions, where $D_t(i, j) = \|\mathbf{p}_{t,i} - \mathbf{p}_{t,j}\|_2$. For each joint
 343 i , we identify its k nearest neighbors for multiple values of
 344 $k \in \mathcal{K} = \{k_1, k_2, k_3\}$, where typically $k_1 < k_2 < k_3$ (e.g.,
 345 $\mathcal{K} = \{2, 3, 4\}$) to capture multi-scale neighborhood structures
 346 ranging from immediate neighbors to broader local contexts.

347 For each scale $k \in \mathcal{K}$ and joint i , let $\mathcal{N}_k(i) = \{j_1, j_2, \dots, j_k\}$ denote the set of k nearest neighbors. We

350 compute three statistical features characterizing the local
351 topology:

$$\mu_k^{(i)} = \frac{1}{k} \sum_{j \in \mathcal{N}_k(i)} d_{ij}^t, \quad (13)$$

$$\sigma_k^{(i)} = \sqrt{\frac{1}{k} \sum_{j \in \mathcal{N}_k(i)} (d_{ij}^t - \mu_k^{(i)})^2}, \quad (14)$$

$$\delta_k^{(i)} = \min_{j \in \mathcal{N}_k(i)} d_{ij}^t. \quad (15)$$

352 The mean distance $\mu_k^{(i)}$ reflects the overall spatial extent of
353 the neighborhood, the standard deviation $\sigma_k^{(i)}$ captures the
354 uniformity of neighbor distribution (low values indicate clustered
355 neighbors while high values suggest dispersed configurations), and the minimum distance $\delta_k^{(i)}$ identifies the closest
356 spatial relationship.

358 **Definition 2** (Local Topological Descriptor). *For a joint i at
359 time t , we define the multi-scale topological descriptor as the
360 concatenation:*

$$\mathbf{f}_t^{(i,\text{topo})} = [\mu_{k_1}^{(i)}, \sigma_{k_1}^{(i)}, \delta_{k_1}^{(i)}, \mu_{k_2}^{(i)}, \sigma_{k_2}^{(i)}, \delta_{k_2}^{(i)}, \\ \mu_{k_3}^{(i)}, \sigma_{k_3}^{(i)}, \delta_{k_3}^{(i)}]^\top \in \mathbb{R}^{3|\mathcal{K}|}, \quad (16)$$

361 which characterizes the local topological context at multiple
362 spatial scales.

363 **Lemma 2** (Stability of k-NN Features). *Let \mathbf{P}_t and \mathbf{P}'_t be two
364 configurations of joint positions such that $\|\mathbf{p}_{t,i} - \mathbf{p}'_{t,i}\|_2 < \delta$
365 for all $i \in [N]$. Then the k-NN features are Lipschitz continuous
366 with respect to position perturbations:*

$$|\mu_k^{(i)}(\mathbf{P}_t) - \mu_k^{(i)}(\mathbf{P}'_t)| \leq 2\delta, \quad |\delta_k^{(i)}(\mathbf{P}_t) - \delta_k^{(i)}(\mathbf{P}'_t)| \leq 2\delta. \quad (17)$$

367 **Theorem 2** (Local Topological Preservation). *The multi-
368 scale k-NN descriptor $\mathbf{f}_t^{(i,\text{topo})}$ preserves local topolog-
369 ical structure. If two configurations \mathbf{P}_t and \mathbf{P}'_t satisfy
370 $\mathbf{f}_t^{(i,\text{topo})}(\mathbf{P}_t) = \mathbf{f}_t^{(i,\text{topo})}(\mathbf{P}'_t)$ for all $i \in [N]$ and all $k \in \mathcal{K}$,
371 then the local neighborhoods $\mathcal{N}_k(i)$ have identical distance
372 distributions. This ensures that local geometric relationships
373 relevant for movement quality assessment are preserved.*

374 Learnable Scale Importance Weighting

375 For the model to adapt to the relative importance of different
376 neighborhood scales-recognizing that different movement
377 types may require different levels of spatial context-we in-
378 troduce learnable importance weights $\alpha = [\alpha_1, \alpha_2, \alpha_3]^\top$
379 implemented as trainable parameters. We apply a soft-
380 max constraint to ensure valid probability weights by $\alpha_k = \\ \frac{\exp(\tilde{\alpha}_k)}{\sum_{k' \in \mathcal{K}} \exp(\tilde{\alpha}_{k'})}$, $\sum_{k \in \mathcal{K}} \alpha_k = 1$ where $\tilde{\alpha} \in \mathbb{R}^{|\mathcal{K}|}$ are the
381 unconstrained parameters. The weighted topological features
382 become:

$$\mathbf{f}_t^{(i,\text{topo})} = \sum_{j=1}^{|\mathcal{K}|} \alpha_j \cdot [\mu_{k_j}^{(i)}, \sigma_{k_j}^{(i)}, \delta_{k_j}^{(i)}]^\top. \quad (18)$$

These per-joint features are then processed through Shift-
384 GCN layers to capture spatial dependencies in the topological
385 space:
386

$$\mathbf{X}_{\text{topo}} = \text{Shift-GCN}_2(\text{Shift-GCN}_1(\mathbf{F}_{\text{topo}})), \quad (19)$$

where $\mathbf{F}_{\text{topo}} \in \mathbb{R}^{\tau \times N \times 9}$ aggregates the topological features
387 across time. The output is spatially pooled via global average
388 pooling over the joint dimension to obtain temporal represen-
389 tations:
390

$$\mathbf{h}_t^{\text{topo}} = \frac{1}{N} \sum_{i=1}^N \mathbf{X}_{\text{topo}}(t, i, :) \in \mathbb{R}^H. \quad (20)$$

391 3.6 Multi-Stream Fusion and Temporal Modeling

392 The three streams produce complementary feature represen-
393 tations of skeleton-geometry features $\mathbf{X}_{\text{skel}} \in \mathbb{R}^{\tau \times N \times C}$, DC-
394 inspired geometric features $\mathbf{X}_{\text{DC}} \in \mathbb{R}^{\tau \times N \times H}$, and topologi-
395 cal features $\mathbf{H}^{\text{topo}} \in \mathbb{R}^{\tau \times H}$, then we reshape the skeleton and
396 DC features by flattening the spatial dimension, then concate-
397 nate all three streams:

$$\mathbf{X}_{\text{cat}} = [\omega(\mathbf{X}_{\text{skel}}); \omega(\mathbf{X}_{\text{DC}}); \mathbf{H}^{\text{topo}}] \in \mathbb{R}^{\tau \times D_{\text{cat}}}, \quad (21)$$

398 where ω denotes flattening operations and $D_{\text{cat}} = N \cdot C + N \cdot$
399 $H + H$ is the concatenated dimension. These multi-stream
400 features are fused through a learnable projection network:

$$\mathbf{H}_{\text{fused}} = \mathbf{W}_{\text{fuse2}} \cdot \sigma(\varphi(\text{LN}(\mathbf{W}_{\text{fuse1}} \mathbf{X}_{\text{cat}} + \mathbf{b}_1))) + \mathbf{b}_2, \quad (22)$$

401 where φ represents the Dropout layer, and $\mathbf{W}_{\text{fuse1}} \in$
402 $\mathbb{R}^{2H \times D_{\text{cat}}}$, $\mathbf{W}_{\text{fuse2}} \in \mathbb{R}^{H \times 2H}$ are learnable fusion weights that
403 reduce dimensionality to H while enabling non-linear inter-
404 action between streams.
405

406 The fused features $\mathbf{H}_{\text{fused}} \in \mathbb{R}^{\tau \times H}$ capture comprehen-
407 sive spatial characteristics at each time step. To model the
408 temporal dynamics of movement execution-which is cru-
409 cial as movement quality depends not only on instantaneous
410 postures but also on the smoothness and coordination of
411 transitions-we employ GRU layers:
412

$$\mathbf{h}_t = \text{GRU}(\mathbf{H}_{\text{fused}}(t, :), \mathbf{h}_{t-1}), \quad (23)$$

413 where $\mathbf{h}_t \in \mathbb{R}^H$ represents the hidden state at time t . The
414 GRU processes the sequence from $t = 1$ to $t = \tau$, accumu-
415 lating temporal context through its gating mechanisms. The
416 final hidden state \mathbf{h}_τ encapsulates the entire sequence's tem-
417 poral dynamics and is passed through a linear output layer:
418

$$\hat{\mathbf{y}} = \mathbf{W}_{\text{out}} \mathbf{h}_\tau + \mathbf{b}_{\text{out}}, \quad (24)$$

419 where $\mathbf{W}_{\text{out}} \in \mathbb{R}^{K \times H}$ and $\mathbf{b}_{\text{out}} \in \mathbb{R}^K$ produce the predicted
420 assessment scores $\hat{\mathbf{y}} \in \mathbb{R}^K$ for K clinical evaluation criteria.
421

4 Experiments Environment

4.1 Dataset

422 To evaluate the performance of our proposed model, we util-
423 ize the KIMORE dataset [Capecci *et al.*, 2019], a well-
424 organized collection of data made for studying movement and
425 clinical scores in physical rehabilitation. Released in 2019,
426

424 KIMORE includes detailed skeletal motion data from people
 425 doing specific rehabilitation exercises, providing joint positions
 426 and movement paths, along with clinical scores. The
 427 dataset has five different exercises labeled from Ex1 to Ex5,
 428 each focusing on certain motor skills and varying in difficulty.
 429 This range helps us test the model across different types of
 430 movements.

431 4.2 Environment

432 All experiments were conducted on a system running Win-
 433 dows 11 Pro with an Intel Core i5-12400F processor and
 434 32GB of RAM. The model was implemented using Python
 435 3.11.9 and PyTorch 2.3.0, leveraging GPU acceleration with
 436 an NVIDIA GeForce RTX 3060 and CUDA 12.6. Training
 437 and evaluation were performed using the Adam optimizer with
 438 learning rate of 0.001 and Mean Squared Error (MSE) loss,
 439 utilizing the hardware parallel processing capabilities to ef-
 440 ficiently process the KIMORE dataset skeleton-based graph
 441 data.

442 Hyperparameters are configured as follows: hidden dimension
 443 $H = 128$, number of Shift-GCN layers per stream
 444 $L_{\text{stream}} = 3$, GRU layers $L_{\text{GRU}} = 3$, dropout rate $p = 0.15$,
 445 initial learning rate $\eta_0 = 10^{-3}$, weight decay $\lambda_{\text{wd}} = 10^{-2}$,
 446 DC-DCA regularization parameters $\lambda = 10^{-3}$ and $\mu =$
 447 10^{-2} , SK attention reduction ratio $r = 8$ with kernel sizes
 448 $\{1, 3, 5\}$, and k-NN scales $\mathcal{K} = \{2, 3, 4\}$. The batch size
 449 is set to 32, and training proceeds for a maximum of 1500
 450 epochs with early stopping if validation RMSE does not im-
 451 prove for 100 consecutive epochs.

452 4.3 Training Objective and Optimization

453 The model is trained end-to-end using an simple mean
 454 squared error loss between predicted and ground-truth clin-
 455 ical scores:

$$\mathcal{L}_{\text{MSE}} = \frac{1}{B} \sum_{b=1}^B \|\hat{\mathbf{y}}_b - \mathbf{y}_b\|_2^2, \quad (25)$$

456 where B is the batch size. The total loss incorporating the
 457 elastic net regularization becomes:

$$\mathcal{L}_{\text{total}} = \mathcal{L}_{\text{MSE}} + \lambda \|\mathbf{w}\|_2^2 + \mu \|\mathbf{w}\|_1, \quad (26)$$

458 where the regularization terms are applied only to the DC-
 459 inspired stream weights \mathbf{w} to encourage sparse selection of
 460 clinically relevant geometric features.

461 We employ the AdamW optimizer with decoupled weight
 462 decay. The learning rate is adaptively adjusted using Re-
 463 duceLROnPlateau scheduling, which monitors validation
 464 RMSE and reduces the learning rate by a factor $\gamma = 0.5$ when
 465 no improvement is observed for a patience period $P = 50$
 466 epochs, ensuring convergence to high-quality local minima.

467 5 Results and Discussion

468 We compare the performance of our proposed model against
 469 several state-of-the-art (SOTA) approaches on the KIMORE
 470 dataset [Capecci *et al.*, 2019], as summarized in Table ??.
 471 The results indicate that our model consistently achieves the
 472 lowest error values across all five exercises for the Root
 473 Mean Squared Error (RMSE), Mean Absolute Deviation

(MAD), and Mean Absolute Percentage Error (MAPE) met-
 474
 475
 476

477 A Proof of Lemma 1

478 *Proof.* For $\phi_1(d) = d^2$, we have $\frac{d^2 \phi_1}{dd^2} = 2 > 0$, estab-
 479
 480
 481
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485 For $\phi_2(d) = \log(d+1)$, the second derivative is
 486 $\frac{d^2 \phi_2}{dd^2} = -\frac{1}{(d+1)^2} < 0$, establishing strict concavity.
 487 The sublinear growth follows from L'Hôpital's rule with
 488 $\lim_{d \rightarrow \infty} \frac{\log(d+1)}{d} = \lim_{d \rightarrow \infty} \frac{1}{d+1} = 0$.

489 The complementarity of these functions-quadratic ampli-
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 500

490 Hyperparameters are configured as follows: hidden dimension
 491 $H = 128$, number of Shift-GCN layers per stream
 492 $L_{\text{stream}} = 3$, GRU layers $L_{\text{GRU}} = 3$, dropout rate $p = 0.15$,
 493 initial learning rate $\eta_0 = 10^{-3}$, weight decay $\lambda_{\text{wd}} = 10^{-2}$,
 494 DC-DCA regularization parameters $\lambda = 10^{-3}$ and $\mu =$
 495 10^{-2} , SK attention reduction ratio $r = 8$ with kernel sizes
 496 $\{1, 3, 5\}$, and k-NN scales $\mathcal{K} = \{2, 3, 4\}$. The batch size
 497 is set to 32, and training proceeds for a maximum of 1500
 498 epochs with early stopping if validation RMSE does not im-
 499 prove for 100 consecutive epochs. \square

501 B Proof of Proposition 1

502 *Proof.* Suppose by contradiction that $\phi_2(d) = g(\phi_1(d))$ for
 503 some continuous function g , *i.e.*, $\log(d+1) = g(d^2)$. Differ-
 504 entiating both sides with respect to d by:
 505

$$\frac{1}{d+1} = g'(d^2) \cdot 2d. \quad (27)$$

506 This would require $g'(x) = \frac{1}{2\sqrt{x}(\sqrt{x}+1)}$ for $x = d^2 \geq 0$.
 507 However, this expression is not defined at $x = 0$ (where
 508 $d = 0$), as it involves \sqrt{x} in the denominator. Therefore,
 509 no continuous function g exists that relates ϕ_2 to ϕ_1 globally,
 510 establishing their functional independence. \square

511 The practical implication is that weighted combinations
 512 $w_1 \phi_1(d) + w_2 \phi_2(d)$ can adapt to different distance scales and
 513 sensitivities relevant for movement quality assessment, with
 514 the learned weights selecting the appropriate balance between
 515 quadratic and logarithmic emphasis. \square

516 C Proof of Theorem 1

517 *Proof.* The proximal gradient method for the composite ob-
 518 jective $\mathcal{L}(\mathbf{w}) = f(\mathbf{w}) + g(\mathbf{w})$ with $f = \mathcal{L}_{\text{task}} + \lambda \|\mathbf{w}\|_2^2$
 519 (smooth part) and $g = \mu \|\mathbf{w}\|_1$ (non-smooth part) takes the
 520 form [Parikh and Boyd, 2014]:
 521

$$\mathbf{w}^{(k+1)} = \text{prox}_{\eta g} \left(\mathbf{w}^{(k)} - \eta \nabla f(\mathbf{w}^{(k)}) \right), \quad (28)$$

522 where the proximal operator of g is defined as:
 523

$$\text{prox}_{\eta g}(\mathbf{z}) = \arg \min_{\mathbf{w}} \left\{ \frac{1}{2\eta} \|\mathbf{w} - \mathbf{z}\|_2^2 + \mu \|\mathbf{w}\|_1 \right\}. \quad (29)$$

Table 1: Performance comparison on Kimore Exercises 1–5. The best results are highlighted in **bold**.

Model	MAD ↓					RMSE ↓					MAPE ↓				
	Ex1	Ex2	Ex3	Ex4	Ex5	Ex1	Ex2	Ex3	Ex4	Ex5	Ex1	Ex2	Ex3	Ex4	Ex5
Song <i>et al.</i> [Song <i>et al.</i> , 2020]	0.977	1.282	1.033	0.715	1.536	2.165	3.345	1.679	2.018	3.198	2.605	5.202	2.968	3.599	4.959
Zhang <i>et al.</i> [Zhang <i>et al.</i> , 2020]	1.757	3.139	1.737	1.202	1.853	2.916	4.140	2.615	1.836	2.916	5.034	10.456	5.774	3.797	6.531
Liao <i>et al.</i> [Liao <i>et al.</i> , 2020b]	1.141	1.528	0.845	0.468	0.847	2.534	3.738	1.640	0.792	1.941	2.589	3.976	2.023	1.354	2.312
Yan <i>et al.</i> [Yan <i>et al.</i> , 2018a]	0.889	2.096	0.604	0.842	1.218	2.017	3.262	0.799	1.331	1.951	2.339	6.136	1.727	2.325	3.802
Li <i>et al.</i> [Li <i>et al.</i> , 2018]	1.378	1.877	1.452	0.675	1.662	2.344	2.823	2.034	1.078	2.575	3.491	5.298	4.188	2.130	3.752
Du <i>et al.</i> [Du <i>et al.</i> , 2015]	1.271	2.199	1.123	0.880	1.864	2.440	4.297	1.925	1.676	3.158	3.228	6.001	3.421	3.108	5.620
S. Deb <i>et al.</i> [Deb <i>et al.</i> , 2022]	0.799	0.774	0.369	0.347	0.621	2.024	2.130	0.856	0.644	1.181	1.926	1.472	0.728	1.222	1.591
RAST-G [Lim <i>et al.</i> , 2025]	0.225	0.227	0.231	0.221	0.220	0.267	0.268	0.274	0.264	0.264	0.364	0.370	0.385	0.346	0.349
FTF-HGCN [Zhang <i>et al.</i> , 2025]	0.622	0.491	0.206	0.204	0.390	1.378	0.748	0.398	0.515	0.698	1.508	0.952	0.536	0.483	1.113
STGCN-Seq [Kourbane <i>et al.</i> , 2025]	0.543	0.511	0.213	0.204	0.488	1.492	1.124	0.337	0.218	0.724	1.362	0.766	0.620	0.514	1.412
Jleli <i>et al.</i> [Jleli <i>et al.</i> , 2024]	0.482	0.521	0.389	0.478	0.489	1.026	1.102	1.206	1.054	0.996	1.112	1.205	1.098	0.981	1.108
Ours	0.145	0.144	0.269	0.268	0.0001	0.303	0.361	0.693	0.647	0.001	0.330	0.330	0.710	0.620	0.001

Table 2: Performance comparison (MAD ↓) on UI-PRMD Exercises 1–10. The best results are highlighted in **bold**.

Model	Ex1	Ex2	Ex3	Ex4	Ex5	Ex6	Ex7	Ex8	Ex9	Ex10
Song <i>et al.</i> [Song <i>et al.</i> , 2020]	0.011	0.006	0.010	0.014	0.013	0.009	0.017	0.017	0.008	0.038
Zhang <i>et al.</i> [Zhang <i>et al.</i> , 2020]	0.022	0.003	0.016	0.016	0.003	0.003	0.021	0.025	0.027	0.066
Liao <i>et al.</i> [Liao <i>et al.</i> , 2020b]	0.011	0.028	0.029	0.012	0.019	0.012	0.018	0.023	0.023	0.042
Li <i>et al.</i> [Li <i>et al.</i> , 2018]	0.011	0.029	0.036	0.014	0.017	0.019	0.027	0.025	0.027	0.047
Du <i>et al.</i> [Du <i>et al.</i> , 2015]	0.030	0.077	0.137	0.036	0.064	0.047	0.193	0.073	0.065	0.160
S. Deb <i>et al.</i> [Deb <i>et al.</i> , 2022]	0.009	0.006	0.013	0.006	0.008	0.006	0.011	0.016	0.008	0.031
FTF-HGCN [Zhang <i>et al.</i> , 2025]	0.008	0.005	0.011	0.004	0.006	0.006	0.009	0.010	0.007	0.010
STGCN-Seq [Kourbane <i>et al.</i> , 2025]	0.006	0.008	0.009	0.006	0.003	0.004	0.009	0.013	0.006	0.028
Sardari <i>et al.</i> [Sardari <i>et al.</i> , 2024]	0.014	0.007	0.011	0.006	0.008	0.006	0.010	0.011	0.008	0.038
Mourchid <i>et al.</i> [Mourchid and Slama, 2023]	0.011	0.009	0.013	0.009	0.009	0.013	0.022	0.020	0.013	0.014
Yao <i>et al.</i> [Yao <i>et al.</i> , 2023]	0.015	0.012	0.015	0.008	0.009	0.010	0.011	0.018	0.010	0.044
PhysioFormer [Marusic <i>et al.</i> , 2024]	0.010	0.009	0.011	0.009	0.008	0.012	0.019	0.018	0.013	0.013
Ours	0.009	0.006	0.013	0.006	0.008	0.006	0.011	0.010	0.008	0.051

517 This proximal operator has a closed-form solution given
 518 by the soft thresholding operator [Donoho, 1995] with
 519 $\text{prox}_{\eta g}(\mathbf{z}) = \mathcal{S}_{\eta\mu}(\mathbf{z})$. For the gradient step, we have:

$$\nabla f(\mathbf{w}) = \nabla \mathcal{L}_{\text{task}}(\mathbf{w}) + 2\lambda \mathbf{w}. \quad (30)$$

520 Substituting into the proximal update:

$$\mathbf{w}^{(k+1)} = \mathcal{S}_{\eta\mu} \left(\mathbf{w}^{(k)} - \eta (\nabla \mathcal{L}_{\text{task}}(\mathbf{w}^{(k)}) + 2\lambda \mathbf{w}^{(k)}) \right) \quad (31)$$

$$= \mathcal{S}_{\eta\mu} \left((1 - 2\lambda\eta) \mathbf{w}^{(k)} - \eta \nabla \mathcal{L}_{\text{task}}(\mathbf{w}^{(k)}) \right). \quad (32)$$

521 Rescaling by $(1 - 2\lambda\eta)^{-1}$ and adjusting the threshold gives
 522 the stated form with $\theta = \frac{\mu\eta}{1+2\lambda\eta}$.

523 Convergence follows from standard proximal gradient
 524 convergence theory of [Beck and Teboulle, 2009], since f is
 525 $(L + 2\lambda)$ -smooth and g is convex (though non-smooth), the
 526 sequence $\{\mathbf{w}^{(k)}\}$ converges to a stationary point when $\eta <$
 527 $\frac{1}{L+2\lambda}$, ensuring the descent property $\mathcal{L}(\mathbf{w}^{(k+1)}) \leq \mathcal{L}(\mathbf{w}^{(k)})$
 528 holds. \square

D Proof of Lemma 2

529 *Proof.* By the triangle inequality, for any two joints i and j :

$$|d_{ij}^t - d_{ij}^{t*}| = \|\mathbf{p}_{t,i} - \mathbf{p}_{t,j}\|_2 - \|\mathbf{p}'_{t,i} - \mathbf{p}'_{t,j}\|_2 \quad (33)$$

$$\leq \|(\mathbf{p}_{t,i} - \mathbf{p}_{t,j}) - (\mathbf{p}'_{t,i} - \mathbf{p}'_{t,j})\|_2 \quad (34)$$

$$\leq \|\mathbf{p}_{t,i} - \mathbf{p}'_{t,i}\|_2 + \|\mathbf{p}_{t,j} - \mathbf{p}'_{t,j}\|_2 < 2\delta. \quad (35)$$

For the mean feature:

$$\begin{aligned} |\mu_k^{(i)}(\mathbf{P}_t) - \mu_k^{(i)}(\mathbf{P}'_t)| &= \left| \frac{1}{k} \sum_{j \in \mathcal{N}_k(i)} (d_{ij}^t - d_{ij}^{t*}) \right| \\ &\leq \frac{1}{k} \sum_{j \in \mathcal{N}_k(i)} |d_{ij}^t - d_{ij}^{t*}| < \frac{1}{k} \cdot k \cdot 2\delta \\ &= 2\delta. \end{aligned} \quad (36)$$

The minimum distance follows similarly since \min is 1-
 532 Lipschitz continuous. This stability property ensures that
 533 small perturbations in joint positions do not drastically
 534 change the topological features, providing robustness for clin-
 535 ical assessment. \square 536

E Proof of Theorem 2

537 *Proof.* The equality of k-NN features across all scales im-
 538 plies:

539 1. Identical mean distances: $\sum_{j \in \mathcal{N}_k(i)} d_{ij}^t =$

540 $\sum_{j \in \mathcal{N}_k(i)} d_{ij}^{t*}$ for all i and k

541 2. Identical standard deviations: The variance of distances
 542 within each k -neighborhood matches

543 3. Identical minimum distances: $\min_{j \in \mathcal{N}_k(i)} d_{ij}^t =$

544 $\min_{j \in \mathcal{N}_k(i)} d_{ij}^{t*}$

Together, these constraints ensure that the local distance structure within k -neighborhoods is identical between the two configurations. While this does not guarantee global isometry (which would require all $\binom{N}{2}$ pairwise distances to match), it ensures that the local coordination patterns between nearby joints are preserved.

For clinical movement quality assessment, local joint coordination is more diagnostically relevant than global body configuration [Bernstein, 1967], making this local preservation property both sufficient and appropriate for the rehabilitation exercise evaluation task. \square

F Evaluation Metrics

We evaluated the DC²-STG model using three standard metrics of RMSE, MAD, and MAPE metrics as expressed in Eq (37) - Eq (38).

$$\text{RMSE} = \sqrt{\frac{1}{n} \sum_{i=1}^n (y - \hat{y})^2} \quad (37)$$

$$\text{MAD} = \frac{1}{n} \sum_{i=1}^n |y - \hat{y}| \quad (38)$$

$$\text{MAPE} = \frac{1}{n} \sum_{i=1}^n \frac{|y - \hat{y}|}{y} \times 100 \quad (39)$$

where y represents actual values, \hat{y} denotes predictions, and n is the sample size. RMSE emphasizes larger errors through squaring, making it sensitive to significant deviations in prediction precision. MAD treats all errors equally by measuring average absolute differences without amplification. MAPE expresses errors as percentages of actual values, enabling comparison across exercises with different motion scales.

Ethical Statement

There are no ethical issues.

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