
GNNs Meet Sequence Models Along the Shortest-Path: an Expressive Method for Link Prediction

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Abstract

Graph Neural Networks (GNNs) often struggle to capture the link-specific structural patterns crucial for accurate link prediction, as their node-centric message-passing schemes overlook the subgraph structures connecting a pair of nodes. Existing methods to inject such structural context either incur high computational cost or rely on simplistic heuristics (e.g., common neighbor counts) that fail to model multi-hop dependencies. We introduce SP4LP (Shortest Path for Link Prediction), a novel framework that combines GNN-based node encodings with sequence modeling over shortest paths. Specifically, SP4LP first applies a GNN to compute representations for all nodes, then extracts the shortest path between each candidate node pair and processes the resulting sequence of node embeddings using a sequence model. This design enables SP4LP to capture expressive multi-hop relational patterns with computational efficiency. Empirically, SP4LP achieves state-of-the-art performance across link prediction benchmarks. Theoretically, we prove that SP4LP is strictly more expressive than standard message-passing GNNs and several state-of-the-art structural features methods, establishing it as a general and principled approach for link prediction in graphs.

1 Introduction

Graph Neural Networks (GNNs) are widely adopted for link-level tasks such as link prediction [33, 18, 35], link classification [21, 28, 3] and link regression [16, 6] with applications spanning recommender systems [31], knowledge graph completion [20], and biological interaction prediction [12].

Despite their popularity, standard GNNs struggle to accurately represent links, as they typically construct link embeddings by aggregating the representations of the two endpoint nodes. This node-centric strategy leads to a key limitation: structurally distinct links may be mapped to the same representation when their endpoints are automorphic [22, 2, 34]. For example, in the graph of Figure 1, links (v, u) and (v, u') yield identical representations under any standard GNN, even if one pair shares a common neighbor and the other does not. This issue, known as the automorphic node problem [2], highlights an expressivity bottleneck in message-passing schemes for link representation.

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To address this, several methods enhance GNNs with structural features (SFs), which can be broadly classified into three paradigms [27]: SF-then-GNN, which injects structural context into the graph before message passing (e.g., SEAL [34], NBFNet [36]); SF-and-GNN, which computes SFs and node embeddings in parallel (e.g., Neo-GNN [32], BUDDY [2]); and GNN-then-SF, which applies message passing once to compute node representations and then combines them using task-specific structural context (e.g., NCN and NCNC [27]).

While SF-then-GNN methods are expressive, they are computationally inefficient, often requiring subgraph extraction or retraining per link. SF-and-GNN models are efficient but rely on predefined heuristics, limiting their ability to capture rich relational patterns. GNN-then-SF approaches offer a compelling trade-off between expressivity and scalability, but current methods in this class, i.e., NCN and NCNC, depend on the presence of common neighbors between link endpoints. When such neighbors are absent, they revert to standard GNN behavior and lose expressive power.

In this paper we propose SP4LP, a novel method in the GNN-then-SF paradigm that combines high expressiveness with computational efficiency. SP4LP constructs a path-aware representation by incorporating the embeddings of all nodes along the shortest path connecting the two endpoints. These node embeddings, obtained via a base GNN, are then processed as a sequence using a dedicated sequence model, such as a Transformer [23], LSTM [10]. Unlike structural features such as common neighbors, which may not exist for many node pairs and can lead to degenerate cases in sparse graphs, the shortest path is always defined between any two nodes if the graph is connected. Shortest paths are more broadly defined, as common neighbors imply a path, but not vice versa. We formally prove that SP4LP is strictly more expressive than existing SF-and-GNN and GNN-then-SF approaches.

SP4LP achieves **state-of-the-art performance** across several benchmark datasets. Under the challenging HearT evaluation protocol [15], it consistently outperforms existing link prediction methods.

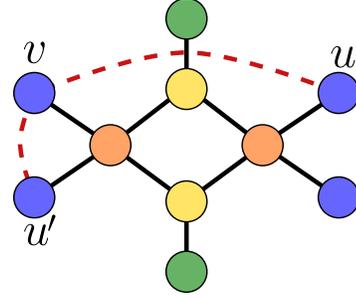


Figure 1: Links (v, u) and (v, u') have different structural roles within the graph, yet a GNN assigns them identical representations.

2 Preliminaries

Definition 2.1 (graph). A **graph** is a tuple $G = (V, E, \mathbf{X}^0)$ where $V = \{1, \dots, n\}$ is a set of nodes, $E \subseteq V \times V$ is a set of edges and $\mathbf{X}^0 \in \mathbb{R}^{n \times f}$ is the node features matrix. To each graph is associated an adjacency matrix $\mathbf{A} \in \{0, 1\}^{n \times n}$ with $\mathbf{A}_{i,j} = 1$ if and only if $(i, j) \in E$. In this work, we consider simple, finite and undirected graphs.

Definition 2.2 (message passing). Let $G = (V, E, \mathbf{X}^0)$ be a graph. In **message passing** scheme, representation of nodes $v \in V$ is iteratively updated as follows:

$$\mathbf{x}_v^0 = \mathbf{X}_{[v,:]}^0 \tag{1}$$

$$\mathbf{x}_v^l = \text{UPDATE}(\mathbf{x}_v^{l-1}, \text{AGGREGATE}(\{\mathbf{x}_u^{l-1} \mid u \in N(v)\})) \tag{2}$$

where $N(v)$ is the first-order neighborhood of node v .

GNNs are a class of neural architectures that operate on graphs by iteratively updating node representations through the message passing scheme. It has been proven that GNNs are at most as effective as the Weisfeiler–Lehman (WL) test in distinguishing between graphs [19, 29].

Definition 2.3 (GNN link representation model). A **GNN link representation model** M is a class of functions $F : ((u, v), G) \mapsto \mathbf{x}_{(u,v)} \in \mathbb{R}^d$ which maps node pairs in $(u, v) \in V \times V$ to vector representations using the message passing scheme defined in Definition 2.2.

Note that the pair (u, v) belongs to $V \times V$, meaning the model learns representations for all possible pairs (links), not just those connected by an edge. A widely adopted approach for learning such representations is what we refer to as a *pure GNN*, defined as follows:

Definition 2.4 (pure GNN). A **pure GNN** model calculates representation $\mathbf{x}_{(u,v)} \in \mathbb{R}^d$ for each pair of nodes (u, v) with $u, v \in V$ as follows:

$$\mathbf{x}_{(u,v)} = g(\mathbf{x}_u^L, \mathbf{x}_v^L) \quad (3)$$

where g is an aggregation function and $\mathbf{x}_u^L, \mathbf{x}_v^L$ are the node representation of u and v learned by L layers of message passing as defined in Definition 2.2.

Pure GNNs are inherently limited in terms of expressiveness. In particular even when the base GNN is the most powerful, they may assign the same representation to structurally different links. We provide a formal definition of what it means for two links to be different.

Definition 2.5 (node permutation). A **node permutation** $\pi : \{1, \dots, n\} \rightarrow \{1, \dots, n\}$ is a bijective function that assigns a new index to each node of the graph. All the $n!$ possible node permutations constitute the permutation group Π_n . Given a subset of nodes $S \subseteq V$, we define the permutation π on S as $\pi(S) := \{\pi(i) | i \in S\}$. Additionally, we define $\pi(\mathbf{A})$ as the matrix \mathbf{A} with rows and columns permuted based on π , i.e., $\pi(\mathbf{A})_{\pi(i), \pi(j)} = \mathbf{A}_{i,j}$.

Definition 2.6 (automorphism). An **automorphism** on the graph $G = (V, E, \mathbf{X}^0)$ is a permutation $\sigma \in \Pi_n$ such that $\sigma(\mathbf{A}) = \mathbf{A}$. All the possible automorphisms on a graph constitute the automorphism group Σ_n^G .

Definition 2.7 (automorphic nodes). Let $G = (V, E, \mathbf{X}^0)$ be a graph and Σ_n^G its automorphism group. Two nodes $u, v \in V$ are said to be **automorphic nodes** ($u \simeq v$) if:

$$\exists \sigma \in \Sigma_n^G \quad \text{s.t.} \quad \sigma(\{u\}) = \{v\}. \quad (4)$$

Definition 2.8 (automorphic links). Let $G = (V, E, \mathbf{X}^0)$ be a graph and Σ_n^G its automorphism group. Two pairs of nodes $(u, v), (u', v') \in V \times V$ are said to be **automorphic links** ($(u, v) \simeq (u', v')$) if:

$$\exists \sigma \in \Sigma_n^G \quad \text{s.t.} \quad \sigma(\{u, v\}) = \{u', v'\}. \quad (5)$$

Proposition 2.9. *Pure GNN methods suffer from the automorphic node problem, i.e., for any graph $G = (V, E, \mathbf{X}^0)$, for pairs of links $(u, v), (u', v') \in V \times V$ such that there exist $\sigma_1 \in \Sigma_n^G$ and $\sigma_2 \in \Sigma_n^G$ with $\sigma_1(u) = u'$ and $\sigma_2(v) = v'$, $\mathbf{x}_{(u,v)} = \mathbf{x}_{(u',v')}$, independently whether (u, v) and (u', v') are isomorphic, i.e, whether exist $\sigma \in \Sigma_n^G$ with $\sigma(\{u, v\}) = \{u', v'\}$.*

This limitation does not arise from the expressiveness bounds of GNNs, which are constrained by the WL test. Even considering higher-order GNNs, i.e., k -GNN [19], automorphic nodes will be assigned to the same representation as the k -WL algorithm preserves graph automorphisms for every k [17, 5, 1]. To tackle this, several models have been proposed that enhance message passing by incorporating structural features [27, 36, 2, 26, 34], thereby increasing the expressive power. We provide a formal definition of what it means for one link representation model to be more expressive and strictly more expressive than another.

Definition 2.10 (more expressive). Let M_1 and M_2 be two link representation models (Def. 2.3). M_2 is **more expressive** than M_1 ($M_1 \preceq M_2$) if, for any graph $G = (V, E, \mathbf{X}^0)$ and any pair $(u, v), (u', v') \in V \times V$ with $(u, v) \not\sim (u', v')$:

$$\exists F_1 \in M_1 : F_1((u, v), G) \neq F_1((u', v'), G) \Rightarrow \exists F_2 \in M_2 : F_2((u, v), G) \neq F_2((u', v'), G). \quad (6)$$

Definition 2.11 (strictly more expressive). Let M_1 and M_2 be two link representation models (Def. 2.3). We say that M_2 is **strictly more expressive** than M_1 ($M_1 \prec M_2$) if:

- M_2 is more expressive than M_1 (Def. 2.10), and
- there exists a graph $G = (V, E, \mathbf{X}^0)$ and a pair of links $(u, v), (u', v') \in V \times V$ with $(u, v) \not\sim (u', v')$ such that:

$$\begin{aligned} \forall F_1 \in M_1 : F_1((u, v), G) &= F_1((u', v'), G) \\ \text{and } \exists F_2 \in M_2 : F_2((u, v), G) &\neq F_2((u', v'), G). \end{aligned}$$

In the following section, we introduce our model SP4LP and demonstrate its improved expressive power in distinguishing structurally different links.

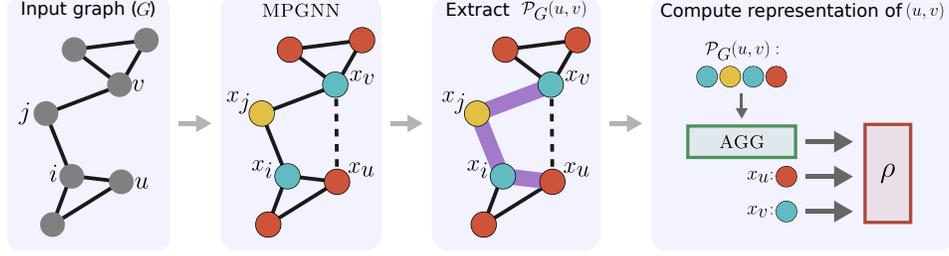


Figure 2: Overview of the SP4LP framework. First, a GNN is used to compute contextualized embeddings for all nodes in the graph. Then, for each target link, the shortest path connecting the two endpoints is extracted. The embeddings of the nodes along this path are passed to a sequence model (e.g., Transformer or LSTM) to compute a path-aware link representation.

3 SP4LP: An Expressive SF-then-GNN Model for link Representation

We propose a model, SP4LP, which adheres to the GNN-then-SF framework, but incorporates additional pairwise information by encoding the sequence of node embeddings along the shortest path connecting the endpoints. This design choice provides a crucial advantage in terms of expressiveness: unlike common neighbors, the shortest path is always defined for nodes in a connected graph and captures richer structural patterns, even in sparse or incomplete graphs. In the following, we introduce the necessary definitions, formally describe the model, and present theoretical results characterizing its expressive power.

Definition 3.1 (path). Let $G = (V, E, \mathbf{X}^0)$ be a graph and $u, v \in V$ to nodes. A **path** in G from u to v is a sequence of nodes $P = (u_0, u_1, \dots, u_k)$ with (i) $u_i \in V$ for all $i = 0, \dots, k - 1$, (ii) $u_0 = u$ and $u_k = v$, (iii) $(u_i, u_{i+1}) \in E$ for all $i = 0, \dots, k - 1$, and (iv) all nodes in the sequence are distinct (i.e., $u_i \neq u_j$ for all $i \neq j$). The length of a path P , $\text{len}(P)$ is the number of edges it contains.

Definition 3.2 (shortest path length). Let $\mathcal{P}_G(u, v)$ denote the set of all paths from u to v in G . The **shortest path length** $d_G(u, v)$ is the minimum length among all paths i.e., $d_G(u, v) = \min_{P \in \mathcal{P}(u, v)} \text{len}(P)$.

Definition 3.3 (shortest path). A **shortest path** between u to v in G is any path $P^* \in \mathcal{P}_G(u, v)$ such that $\text{len}(P^*) = d_G(u, v)$. The set of all the shortest path from u to v in G is denoted as $\mathcal{P}_G^*(u, v)$.

Let $G = (V, E, \mathbf{X}^0)$ be a graph, $u, v \in V$. SP4LP is a GNN link representation model (see Definition 2.3) that computes link representation as follows:

$$\text{SP4LP}((u, v), G) = \rho \left(\text{GNN}(u, G), \text{GNN}(v, G), \text{AGG} \left(\left\{ \phi \left(\text{GNN}(u_i, G) \right)_{i=1}^k \mid (u_i)_{i=1}^k \in \mathcal{P}_G^*\{u, v\} \right\} \right) \right) \quad (7)$$

where $k = d_G(u, v)$, $\text{GNN}(u, G) \in \mathbb{R}^d$ is the representation of node $u \in V$ obtained at the final layer of message passing as in Definition 2.2, $\phi : \mathbb{R}^{k \times d} \rightarrow \mathbb{R}^d$ is a sequence model on the GNN representations of nodes in the shortest path from u to v , AGG is an aggregation function over multiset of shortest paths representations and $\rho : \mathbb{R}^d \times \mathbb{R}^d \times \mathbb{R}^d \rightarrow \mathbb{R}^d$ combine the endpoint nodes representations with the shortest paths representation to get a final link representation. Since we consider undirected graphs, we consider $\mathcal{P}_G^*\{u, v\} := \mathcal{P}_G^*(u, v) \cup \mathcal{P}_G^*(v, u)$.

For a graph with n nodes and m edges, the shortest paths from a single source node can be computed via a breadth-first search (BFS) [4] in $O(m)$ time. Consequently, computing shortest paths between all pairs of nodes requires $O(nm)$ time overall. In sparse graphs, where $m = O(n)$, this yields a quadratic cost $O(n^2)$, which remains tractable in practice. Notably, this computation is performed only once as a **preprocessing step** and can be amortized across multiple downstream predictions.

SP4LP is a general framework: the GNN component can be instantiated with architectures such as GCN [13], GAT [24] or GraphSAGE [9], while the sequence model can range from simple aggregation functions like injective summation as the one proposed in Xu et al. [29], to more complex architectures such as LSTMs [10] or Transformers [23]. An overview of SP4LP is illustrated in Figure 2.

Proposition 3.4. *SP4LP does not suffer from the automorphic node problem.*

Table 1: MRR and Hits@K (%) results across all datasets, following the HeaRT evaluation setting [15]. The top three results for each metric are highlighted using **first**, **second**, and **third**. *OOM* indicates that the model ran out of memory, while *>24h* denotes that the method did not complete within 24 hours. Standard deviations over 5 runs are reported in the appendix B.

Models	Cora		Citeseer		Pubmed		Ogbl-ddi		Ogbl-collab		Ogbl-ppa		Ogbl-Citation2		
	MRR	Hits@10	MRR	Hits@10	MRR	Hits@10	MRR	Hits@20	MRR	Hits@20	MRR	Hits@20	MRR	Hits@20	
GCN	16.61	36.26	21.09	47.23	7.13	15.22	13.46	64.76	6.09	22.48	26.94	68.38	19.98	51.72	
	13.84	32.89	19.58	45.30	4.95	9.99	12.92	66.83	4.18	18.30	<i>OOM</i>	<i>OOM</i>	<i>OOM</i>	<i>OOM</i>	
	14.74	34.65	21.09	48.75	9.40	20.54	12.60	67.19	5.53	21.26	27.27	69.49	22.05	53.13	
		18.32	37.95	25.25	49.65	5.27	10.50	3.49	17.81	<i>OOM</i>	<i>OOM</i>	<i>OOM</i>	<i>OOM</i>	<i>OOM</i>	
SF GNN	SEAL	10.67	24.27	13.16	27.37	5.88	12.47	9.99	49.74	6.43	21.57	29.71	76.77	20.60	48.62
	BUDDY	13.71	30.40	22.84	48.35	7.56	16.78	12.43	58.71	5.67	23.35	27.70	71.50	19.17	47.81
	Neo-GNN	13.95	31.27	17.34	41.74	7.74	17.88	10.86	51.94	5.23	21.03	21.68	64.81	16.12	43.17
	NCN	14.66	35.14	28.65	53.41	5.84	13.22	12.86	65.82	5.09	20.84	35.06	81.89	23.35	53.76
	NCNC	14.98	36.70	24.10	53.72	8.58	18.81	<i>>24h</i>	<i>>24h</i>	4.73	20.49	33.52	82.24	19.61	51.69
	NBFNet	13.56	31.12	14.29	31.39	<i>>24h</i>	<i>>24h</i>	<i>>24h</i>	<i>OOM</i>	<i>OOM</i>	<i>OOM</i>	<i>OOM</i>	<i>OOM</i>	<i>OOM</i>	<i>OOM</i>
	PEG	15.73	36.03	21.01	45.56	4.40	8.70	12.05	50.12	4.83	18.29	<i>OOM</i>	<i>OOM</i>	<i>OOM</i>	<i>OOM</i>
SP4LP (our)	17.27	38.52	41.08	66.28	10.87	23.01	15.00	47.96	9.46	20.00	36.45	76.90	24.91	55.45	

The proof can be found in Appendix A. As an example of non-automorphic links composed of automorphic nodes that SP4LP can successfully distinguish, consider the links (v, u) and (v, u') shown in Figure 1. While $u' \simeq u$ via the identity, and $v \simeq u$ via an automorphism induced by a vertical axis of symmetry (i.e., a mirror reflection), the links (v, u) and (v, u') are not automorphic. This asymmetry is captured by the distinct shortest paths between the endpoints: the shortest path from v to u' consists of v , and orange node, and u' , whereas the shortest path from v to u includes v , and orange node, a yellow node, another orange node, and finally u . In addition to overcoming this limitation, SP4LP is strictly more expressive than several state-of-the-art message passing methods for link representation learning.

Theorem 3.5. *SP4LP is strictly more expressive than Pure GNNs, NCN, BUDDY, NBFnet and Neo-GNN.*

The proof can be found in Appendix A. In the following sections, we complement the theoretical analysis with an extensive experimental evaluation, showing that SP4LP also achieves state-of-the-art performance on standard link prediction benchmarks.

4 Experiments

We evaluate the performance of SP4LP on real-world link prediction benchmarks against several baselines. In particular, we use Cora, Citeseer, and Pubmed [30] as well as datasets from Open Graph Benchmark [11]. For Cora, Citeseer, and Pubmed, we use a single fixed data split in all experiments. Table 6 in appendix C provides a summary of dataset statistics.

As baseline methods we consider two class of models: 1) **Pure GNN methods**: Graph Convolutional Network (GCN) [13], Graph Attention Network (GAT) [25], GraphSAGE [8], and Graph Autoencoder (GAE) [14]; 2) **Structural Features GNN methods**: SEAL [33], BUDDY [2], Neo-GNN [32], NBFNet [36], NCN [27], NCNC [27] and PEG [26].

Importantly, as described in Section 3, SP4LP is a general framework that allows for different choices of both the underlying GNN architecture and the sequence model (ϕ Equation 7). In our experimental setting, we treat the choice of GNN and the choice of ϕ as hyperparameters, and perform hyperparameter tuning based on validation set performance. Specifically, we explore GCN, GAT, and GraphSAGE as GNN backbones, and LSTM, Transformer, and an injective sum Xu et al. [29] aggregator as sequence models. Moreover, we choose an MLP for ρ and as AGG we choose to select the first shortest path retrieved by the BFS procedure for computational efficiency. Appendix D provides implementation details. Code to reproduce all experiments is available at².

Evaluation Setting We evaluate model performance under the more challenging and realistic HeaRT evaluation setting [15]. We adopt two standard ranking metrics: Hits@K and Mean Reciprocal Rank (MRR). Following the HeaRT protocol, we report Hits@10 and MRR for Cora, Citeseer, and Pubmed, and Hits@20 along with MRR for ogbl-collab and ogbl-ddi.

²<https://anonymous.4open.science/r/sp4lp-3875/README.md>

Results on Real-World Benchmarks As shown in Table 1, SP4LP ranks first in terms of MRR on four out of five datasets and second on the remaining one. The improvements in MRR are often substantial: on Citeseer, for instance, SP4LP achieves a 43% gain over the second-best method, NCN. SP4LP also achieves the best Hits@K score on three out of five datasets. On the Ogb1-Collab dataset, SP4LP is comparable on Hits@20 to the third-best model (SEAL), when accounting for standard deviations (Appendix B). On Ogb1-ddi, where SP4LP performs worse, the lower score can be explained by the lack of node features. Our model benefits from the availability of node features, as it leverages nodes representations obtained via message passing. In settings where such features are absent, like in Ogb1-ddi, the discriminative power of the learned representations is reduced. In addition to achieving the best performance in several datasets, SP4LP is also the most consistent model across all benchmarks.

Ablation Study We perform an ablation study to evaluate the contribution of the main components of our model. In particular, we investigate two simplified variants: **(1) Sequence Model Only:** in this variant, the sequential model operates directly on the raw input features of the nodes along the shortest path, without incorporating node representations learned by the GNN. **(2) GNN + Shortest Path Length:** in this variant, the sequential model is completely removed and link prediction is performed using only the learned node representations from the GNN, combined with the length of the shortest path between the target nodes.

Table 2: Ablation study results (%). Standard deviations over 5 runs are reported in Appendix B

Models	Cora		Citeseer		Pubmed	
	MRR	Hits@10	MRR	Hits@10	MRR	Hits@10
<i>GNN + SP len.</i>	14.21	33.43	20.90	47.82	7.12	5.63
<i>Sequence Model</i>	16.86	36.03	27.45	54.20	8.58	12.87
SP4LP	17.27	38.52	41.08	66.28	10.87	23.01

Table 2 reports the results, conducted on Cora, Citeseer, and Pubmed. Both variants show a performance drop compared to the full model, demonstrating the importance of jointly leveraging node representations and sequential modeling of the shortest paths.

Scalability Analysis We assess the scalability of SP4LP by examining how its GPU memory consumption and inference time evolve as the batch size increases, in comparison to several baseline methods. The results, presented in Figure 3, highlight the superior resource efficiency of SP4LP across a wide range of batch sizes. In terms of GPU memory usage, SP4LP exhibits remarkable efficiency: memory consumption remains nearly constant across small to medium batch sizes, and increases moderately only for the largest batches. PEG also maintains low memory usage; however, this advantage is undermined by its impractically slow inference. SEAL, while competitive with SP4LP in terms of inference speed, suffers from excessive memory consumption.

Considering inference time, SP4LP matches the efficiency of SEAL. Although NCN and Buddy consistently achieve low inference times, this comes at the cost of substantially lower predictive performance, as shown in Table 1.

In summary, SP4LP achieves an excellent balance between low memory consumption, fast inference, high predictive accuracy, and strong model expressiveness. It effectively scales to large batch sizes where alternative approaches either become prohibitively slow, fail due to memory constraints, or cannot deliver competitive results.

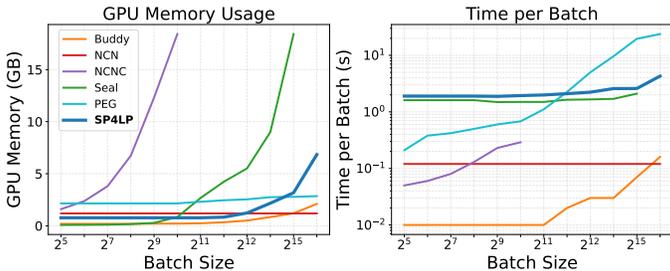


Figure 3: Inference time and GPU memory usage on ogbl-collab, measured during the prediction of a single batch of test links.

5 Conclusion

We introduced SP4LP, a novel message-passing based framework for link representation that enhances the expressiveness of standard GNNs by incorporating sequential modeling over the shortest path between target nodes. We formally proved that SP4LP is strictly more expressive than several state-of-the-art link representation models. Extensive experiments under the HeaRT evaluation protocol confirm that SP4LP achieves state-of-the-art performance across diverse datasets.

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A Proofs

Proposition 3.4 SP4LP does not suffer from the automorphic node problem.

Proof. According to Proposition 2.9, a model M suffers from the *automorphic node problem* if, for any graph $G = (V, E, \mathbf{X}^0)$, for any pairs of links $(u, v), (u', v') \in V \times V$, for any $F \in M$ it holds that:

$$(u, v) \not\cong (u', v'), \quad u \simeq u', \quad v \simeq v', \quad \text{and} \quad F((u, v), G) \neq F((u', v'), G).$$

In order to prove that SP4LP does not suffer from the automorphic node problem, it suffices to provide an example of a graph $G = (V, E, X^0)$ and node pairs $(u, v), (u', v') \in V \times V$ such that:

$$(u, v) \not\cong (u', v'), \quad u \simeq u', \quad v \simeq v', \quad \text{and} \quad \text{SP4LP}((u, v), G) \neq \text{SP4LP}((u', v'), G).$$

Such an example is provided in Figure 1: the shortest path from v to u' consists of v , an orange node, and u' , whereas the shortest path from v to u includes v , an orange node, a yellow node, another orange node, and finally u . Thus, simply using a summation as function ϕ , leads to distinct representations for links (v, u) and (v, u') . □

Theorem 3.5 SP4LP is strictly more expressive than Pure GNN, NCN, BUDDY, NBFnet and Neo-GNN.

Proof. We proceed by prove each comparison separately.

- SP4LP is strictly more expressive than Pure GNN.

We prove this by noting that SP4LP architecture (Equation 7) generalizes that of pure GNNs, meaning that SP4LP can simulate any pure GNN by simply ignoring the shortest path information. Thus, for any pair of non-automorphic links that a specific pure GNN can distinguish, there exists a configuration of SP4LP that distinguishes them as well. We can conclude that SP4LP is strictly more expressive than pure GNNs considering as example of pair of links indistinguishable by GNNs but distinguishable by SP4LP the one provided is provided in the proof of Proposition 3.4.

- SP4LP is strictly more expressive than NCN.

We prove this in two steps: (1) When two links share no common neighbors, NCN on them reduces to a pure GNN. As we have already proved that SP4LP is strictly more expressive than pure GNNs, it follows that SP4LP is also strictly more expressive than NCN in this case. (2) When the links have common neighbors, setting AGG as summation, ρ as the Hadamard product between the endpoint representations and the concatenation with the result of the aggregation, and ϕ as the identity function, SP4LP reduces exactly to NCN. Therefore, if NCN can distinguish two links under some configuration, SP4LP can as well. By definition 2.10), this implies that SP4LP is more expressive than NCN. We can conclude that SP4LP is strictly more expressive than NCN considering the example in Figure 4. The graph is regular and thus all nodes receive the same embedding from a GNN. Consider nodes u and v : $N(u) \cap N(v) = N(u) \cap N(v') = \emptyset$. In this case, NCN reduces to a pure GNN and is thus unable to distinguish the links (u, v) and (u', v') . Now, let $\mathbf{x} \in \mathbb{R}^d$ be the representation assigned to every node by a GNN. Then, the representation of the shortest path $\mathcal{P}_G^*(u, v)$ is simply $(\mathbf{x}, \mathbf{x}, \mathbf{x})$, while $\mathcal{P}_G^*(u', v') = (\mathbf{x}, \mathbf{x}, \mathbf{x}, \mathbf{x})$. Even using a simple sum as aggregation function, SP4LP successfully distinguishes between the two links.

- SP4LP is strictly more expressive than Neo-GNN and BUDDY.

We have already shown that SP4LP is strictly more expressive than NCN. In Theorem 2 of the NCN paper [27], it has been proved that NCN is more expressive than both Neo-GNN and BUDDY. It follows that SP4LP is strictly more expressive than Neo-GNN and BUDDY as well.

- SP4LP is strictly more expressive than NBFNet.

We prove that NBFNet is as expressive as a pure GNN. Therefore, since we have already proven that SP4LP is strictly more expressive than pure GNNs, it immediately follows that SP4LP is also strictly more expressive than NBFNet.

To complete the argument, we prove that NBFNet is as expressive as a pure GNN. First of all we report the formulation of NBFNet for simple undirected graph. Given a graph $G = (V, E, \mathbf{X}^0)$, NBFNet assigns a representation $\mathbf{x}(u, v)$ to each edge $(u, v) \in E$. The iterative update rule follows a message-passing scheme:

$$\begin{aligned} \mathbf{x}_{(u,v)}^{(0)} &= \text{INDICATOR}(\mathbf{x}_u^0, \mathbf{x}_v^0), \\ \mathbf{x}_{(u,v)}^{(l)} &= \text{AGGREGATE} \left(\left\{ \text{MESSAGE}(h_{(i,j)}^{(l-1)}) \mid (i, j) \in N(u, v) \right\} \cup \{h_{(u,v)}^{(0)}\} \right) \end{aligned} \quad (8)$$

where INDICATOR assigns an initial representation based on the nodes $u, v \in V$ and $N(u, v)$ is the set of edges incident to (u, v) .

We prove that, at any layer l , the representations of the two links produced by a pure GNN are equal if and only if also the ones produced by NBFNet are equal, i.e.,

$$\mathbf{x}_{(u,v)}^{\text{GNN}^l} = \mathbf{x}_{(i,j)}^{\text{GNN}^l} \Leftrightarrow \mathbf{x}_{(u,v)}^{\text{NBF}^l} = \mathbf{x}_{(i,j)}^{\text{NBF}^l} \quad \forall l \quad (9)$$

where $\mathbf{x}_{(u,v)}^{\text{NBF}^l}$ and $\mathbf{x}_{(i,j)}^{\text{NBF}^l}$ are calculated via Equation 8, while $\mathbf{x}_{(u,v)}^{\text{GNN}^l}$ and $\mathbf{x}_{(i,j)}^{\text{GNN}^l}$ are calculated following the standard message passing scheme reported below:

$$\begin{aligned} \mathbf{x}_v^{\text{GNN}^0} &= \mathbf{x}_v^0, \\ \mathbf{x}_v^{\text{GNN}^l} &= \text{COMB} \left(\mathbf{x}_v^{\text{GNN}^{l-1}}, \text{AGG} \left(\{ \mathbf{x}_u^{\text{GNN}^{l-1}} \mid u \in N(v) \} \right) \right) \\ \mathbf{x}_{(u,v)}^{\text{GNN}^l} &= g(\mathbf{x}_u^{\text{GNN}^l}, \mathbf{x}_v^{\text{GNN}^l}) \end{aligned} \quad (10)$$

Let the functions INDICATOR, AGGREGATE and MESSAGE of Equation 8, as well as the functions COMB, AGG and g be injective. We prove Equation 9 by induction on the number of layer l .

Base Case: $l = 0$

$$\begin{aligned} \mathbf{x}_{(u,v)}^{\text{GNN}^0} = \mathbf{x}_{(i,j)}^{\text{GNN}^0} &\stackrel{(10)}{\Leftrightarrow} g(\mathbf{x}_u^0, \mathbf{x}_v^0) = g(\mathbf{x}_i^0, \mathbf{x}_j^0) \stackrel{\text{inj}}{\Leftrightarrow} (\mathbf{x}_u^0, \mathbf{x}_v^0) = (\mathbf{x}_i^0, \mathbf{x}_j^0) \stackrel{\text{inj}}{\Leftrightarrow} \\ &\stackrel{\text{inj}}{\Leftrightarrow} \text{INDICATOR}(\mathbf{x}_u^0, \mathbf{x}_v^0) = \text{INDICATOR}(\mathbf{x}_i^0, \mathbf{x}_j^0) \stackrel{(8)}{\Leftrightarrow} \mathbf{x}_{(u,v)}^{\text{NBF}^0} = \mathbf{x}_{(i,j)}^{\text{NBF}^0} \end{aligned}$$

Inductive Step We assume Equation 9 holds for $l - 1$ and prove it holds for l .

In particular, we want to prove

$$\mathbf{x}_{(u,v)}^{\text{GNN}^l} = \mathbf{x}_{(i,j)}^{\text{GNN}^l} \Leftrightarrow \mathbf{x}_{(u,v)}^{\text{NBF}^l} = \mathbf{x}_{(i,j)}^{\text{NBF}^l} \quad (11)$$

using the inductive hypothesis

$$\mathbf{x}_{(u,v)}^{\text{GNN}^{l-1}} = \mathbf{x}_{(i,j)}^{\text{GNN}^{l-1}} \Leftrightarrow \mathbf{x}_{(u,v)}^{\text{NBF}^{l-1}} = \mathbf{x}_{(i,j)}^{\text{NBF}^{l-1}} \quad (12)$$

Applying Equation 10 to the left-hand side of Equation 11 we get

$$\begin{aligned} \mathbf{x}_{(u,v)}^{\text{GNN}^l} &= \mathbf{x}_{(i,j)}^{\text{GNN}^l} \\ &\stackrel{(10)}{\Leftrightarrow} \\ g(\text{COMB}(\mathbf{x}_u^{\text{GNN}^{l-1}}, \text{AGG}(\{ \mathbf{x}_x^{\text{GNN}^{l-1}} \mid x \in N(u) \})), \text{COMB}(\mathbf{x}_v^{\text{GNN}^{l-1}}, \text{AGG}(\{ \mathbf{x}_y^{\text{GNN}^{l-1}} \mid y \in N(u) \}))) & \\ = & \\ g(\text{COMB}(\mathbf{x}_i^{\text{GNN}^{l-1}}, \text{AGG}(\{ \mathbf{x}_m^{\text{GNN}^{l-1}} \mid m \in N(i) \})), \text{COMB}(\mathbf{x}_j^{\text{GNN}^{l-1}}, \text{AGG}(\{ \mathbf{x}_n^{\text{GNN}^{l-1}} \mid n \in N(j) \}))) & \end{aligned}$$

Given the injectivity of g , COMB and AGG, this is equivalent to

$$\begin{aligned}
& \mathbf{x}_u^{\text{GNN}^{l-1}} = \mathbf{x}_i^{\text{GNN}^{l-1}} \wedge \{\mathbf{x}_x^{\text{GNN}^{l-1}} \mid x \in N(u)\} = \{\mathbf{x}_m^{\text{GNN}^{l-1}} \mid m \in N(i)\} \wedge \\
& \wedge \mathbf{x}_v^{\text{GNN}^{l-1}} = \mathbf{x}_j^{\text{GNN}^{l-1}} \wedge \{\mathbf{x}_y^{\text{GNN}^{l-1}} \mid y \in N(v)\} = \{\mathbf{x}_n^{\text{GNN}^{l-1}} \mid n \in N(j)\} \\
& \iff \\
& \{(\mathbf{x}_u^{\text{GNN}^{l-1}}, \mathbf{x}_x^{\text{GNN}^{l-1}}) \mid x \in N(u)\} = \{(\mathbf{x}_i^{\text{GNN}^{l-1}}, \mathbf{x}_m^{\text{GNN}^{l-1}}) \mid m \in N(i)\} \\
& \quad \wedge \\
& \{(\mathbf{x}_v^{\text{GNN}^{l-1}}, \mathbf{x}_y^{\text{GNN}^{l-1}}) \mid y \in N(v)\} = \{(\mathbf{x}_j^{\text{GNN}^{l-1}}, \mathbf{x}_n^{\text{GNN}^{l-1}}) \mid n \in N(j)\} \\
& \iff \\
& \{(\mathbf{x}_u^{\text{GNN}^{l-1}}, \mathbf{x}_x^{\text{GNN}^{l-1}}) \mid x \in N(u)\} \cup \{(\mathbf{x}_v^{\text{GNN}^{l-1}}, \mathbf{x}_y^{\text{GNN}^{l-1}}) \mid y \in N(v)\} \\
& \quad = \\
& \{(\mathbf{x}_i^{\text{GNN}^{l-1}}, \mathbf{x}_m^{\text{GNN}^{l-1}}) \mid m \in N(i)\} \cup \{(\mathbf{x}_j^{\text{GNN}^{l-1}}, \mathbf{x}_n^{\text{GNN}^{l-1}}) \mid n \in N(j)\}.
\end{aligned}$$

By Definition of $N(u, v)$ (Equation 8), this is equivalent to

$$\begin{aligned}
& \{(\mathbf{x}_w^{\text{GNN}^{l-1}}, \mathbf{x}_t^{\text{GNN}^{l-1}}) \mid (w, t) \in N(u, v)\} = \{(\mathbf{x}_a^{\text{GNN}^{l-1}}, \mathbf{x}_b^{\text{GNN}^{l-1}}) \mid (a, b) \in N(i, j)\} \\
& \quad \xrightarrow{\text{inj}} \\
& \{g(\mathbf{x}_w^{\text{GNN}^{l-1}}, \mathbf{x}_t^{\text{GNN}^{l-1}}) \mid (w, t) \in N(u, v)\} = \{g(\mathbf{x}_a^{\text{GNN}^{l-1}}, \mathbf{x}_b^{\text{GNN}^{l-1}}) \mid (a, b) \in N(i, j)\} \\
& \quad \xrightarrow{(10)} \\
& \{\mathbf{x}_{(w,t)}^{\text{GNN}^{l-1}} \mid (w, t) \in N(u, v)\} = \{\mathbf{x}_{(a,b)}^{\text{GNN}^{l-1}} \mid (a, b) \in N(i, j)\} \\
& \quad \xrightarrow{\text{IND. HYP. (12)}} \\
& \{\mathbf{x}_{(w,t)}^{\text{NBF}^{l-1}} \mid (w, t) \in N(u, v)\} = \{\mathbf{x}_{(a,b)}^{\text{NBF}^{l-1}} \mid (a, b) \in N(i, j)\}.
\end{aligned}$$

Using the hypotheses of injective AGG and MESSAGE, this is equivalent to:

$$\begin{aligned}
& \text{AGG}(\{\text{MESSAGE}(\mathbf{x}_{(w,t)}^{\text{NBF}^{l-1}}) \mid (w, t) \in N(u, v)\}) = \text{AGG}(\{\text{MESSAGE}(\mathbf{x}_{(a,b)}^{\text{NBF}^{l-1}}) \mid (a, b) \in N(i, j)\}) \\
& \quad \xrightarrow{(8)} \\
& \mathbf{x}_{(u,v)}^{\text{NBF}^l} = \mathbf{x}_{(i,j)}^{\text{NBF}^l} \tag{13}
\end{aligned}$$

which complete the proof. □

B Additional results

We complement the main results of Section 4 with additional tables reporting the standard deviation computed over five different random seeds, to better assess the stability of each method.

Real-world Datasets: Results with Standard Deviations Table 3 and Table 4 expand the main results in Table 1 by including both the mean and standard deviation of MRR and Hits@ K across runs.

Ablation Study: Results with Standard Deviations Similarly, Table 5 complements the ablation results in Table 2 by reporting mean and standard deviation for Cora, Citeseer, and Pubmed.

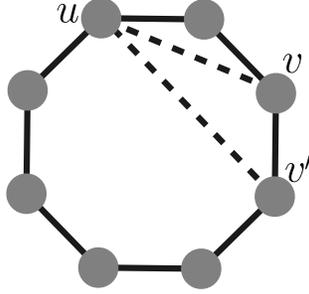


Figure 4: Links (u, v) , (u, v') are not distinguished by NCN while are distinguished by SP4LP.

Models	Cora MRR	Citeseer MRR	Pubmed MRR	Ogbl-ddi MRR	Ogbl-collab MRR
CN	9.78	8.42	2.28	7.11	4.20
AA	11.91	10.82	2.63	7.37	5.07
RA	11.81	10.84	2.47	9.10	6.29
Shortest Path	5.04	5.83	0.86	0	3.06
Katz	11.41	11.19	3.01	7.11	6.31
Node2Vec	14.47 (± 0.60)	21.17 (± 1.01)	3.94 (± 0.24)	11.14 (± 0.95)	4.68 (± 0.08)
MF	6.20 (± 1.42)	7.80 (± 0.79)	4.46 (± 0.32)	13.99 (± 0.47)	4.89 (± 0.25)
MLP	13.52 (± 0.65)	22.62 (± 0.55)	6.41 (± 0.25)	N/A	5.37 (± 0.14)
GCN	16.61 (± 0.30)	21.09 (± 0.88)	7.13 (± 0.27)	13.46 (± 0.34)	6.09 (± 0.38)
GAT	13.84 (± 0.68)	19.58 (± 0.84)	4.95 (± 0.14)	12.92 (± 0.39)	4.18 (± 0.33)
SAGE	14.74 (± 0.69)	21.09 (± 1.15)	9.40 (± 0.70)	12.60 (± 0.72)	5.53 (± 0.50)
GAE	18.32 (± 0.41)	25.25 (± 0.82)	5.27 (± 0.25)	3.49 (± 1.73)	OOM
SEAL	10.67 (± 3.46)	13.16 (± 1.66)	5.88 (± 0.53)	9.99 (± 0.90)	6.43 (± 0.32)
BUDDY	13.71 (± 0.59)	22.84 (± 0.36)	7.56 (± 0.18)	12.43 (± 0.50)	5.67 (± 0.36)
Neo-GNN	13.95 (± 0.39)	17.34 (± 0.84)	7.74 (± 0.30)	10.86 (± 2.16)	5.23 (± 0.90)
NCN	14.66 (± 0.95)	28.65 (± 1.21)	5.84 (± 0.22)	12.86 (± 0.78)	5.09 (± 0.38)
NCNC	14.98 (± 1.00)	24.10 (± 0.65)	8.58 (± 0.59)	>24h	4.73 (± 0.86)
NBFNet	13.56 (± 0.58)	14.29 (± 0.80)	>24h	>24h	OOM
PEG	15.73 (± 0.39)	21.01 (± 0.77)	4.40 (± 0.41)	12.05 (± 1.14)	4.83 (± 0.21)
SP4LP	17.27 (± 0.57)	41.08 (± 1.84)	10.87 (± 0.31)	15.00 (± 0.57)	9.46 (± 0.55)

Table 3: MRR results across all datasets, following the HearT evaluation setting [15]. The top three results for each metric are highlighted using **first**, **second**, and **third**. *OOM* indicates that the model ran out of memory, while *>24h* denotes that the method did not complete within 24 hours.

C Datasets statistics

Table 6 summarizes the main datasets used in our link prediction experiments. Cora, Citeseer, and Pubmed are well-known citation networks frequently used as benchmarks for graph-based learning methods. These datasets are relatively small, both in the number of nodes and edges. In contrast, the datasets from the Open Graph Benchmark (OGB), namely ogbl-collab and ogb-ddi, are substantially larger and more complex, offering challenging scenarios for evaluating model scalability and performance on large-scale graphs.

For Cora, Citeseer, and Pubmed, we adopt a fixed train/validation/test split of 85/5/10%. For the OGB datasets, we use the official data splits provided by the OGB benchmark.

Models	Cora Hits@10	Citeseer Hits@10	Pubmed Hits@10	Ogbl-ddi Hits@20	Ogbl-collab Hits@20
CN	20.11	18.68	4.78	39.09	16.46
AA	24.10	22.20	5.51	40.15	19.59
RA	24.48	22.86	4.90	44.01	24.29
Shortest Path	15.37	16.26	0.38	0	16.38
Katz	22.77	24.84	5.98	39.09	24.34
<hr/>					
Node2Vec	32.77 (± 1.29)	45.82 (± 2.01)	8.51 (± 0.77)	63.63 (± 2.05)	16.84 (± 0.17)
MF	15.26 (± 3.39)	16.72 (± 1.99)	9.42 (± 0.80)	59.50 (± 1.68)	18.86 (± 0.40)
MLP	31.01 (± 1.71)	48.02 (± 1.79)	15.04 (± 0.67)	N/A	16.15 (± 0.27)
<hr/>					
GCN	36.26 (± 1.14)	47.23 (± 1.88)	15.22 (± 0.57)	64.76 (± 1.45)	22.48 (± 0.81)
GAT	32.89 (± 1.27)	45.30 (± 1.30)	9.99 (± 0.64)	66.83 (± 2.23)	18.30 (± 1.42)
SAGE	34.65 (± 1.47)	48.75 (± 1.85)	20.54 (± 1.40)	67.19 (± 1.18)	21.26 (± 1.32)
GAE	37.95 (± 1.24)	49.65 (± 1.48)	10.50 (± 0.46)	17.81 (± 9.80)	OOM
<hr/>					
SEAL	24.27 (± 6.74)	27.37 (± 3.20)	12.47 (± 1.23)	49.74 (± 2.39)	21.57 (± 0.38)
BUDDY	30.40 (± 1.18)	48.35 (± 1.18)	16.78 (± 0.53)	58.71 (± 1.63)	23.35 (± 0.73)
Neo-GNN	31.27 (± 0.72)	41.74 (± 1.18)	17.88 (± 0.71)	51.94 (± 10.33)	21.03 (± 3.39)
NCN	35.14 (± 1.04)	53.41 (± 1.46)	13.22 (± 0.56)	65.82 (± 2.66)	20.84 (± 1.31)
NCNC	36.70 (± 1.57)	53.72 (± 0.97)	18.81 (± 1.16)	>24h	20.49 (± 3.97)
NBFNet	31.12 (± 0.75)	31.39 (± 1.34)	>24h	>24h	OOM
PEG	36.03 (± 0.75)	45.56 (± 1.38)	8.70 (± 1.26)	50.12 (± 6.55)	18.29 (± 1.06)
<hr/>					
SP4LP	38.52 (± 1.19)	66.28 (± 0.63)	23.01 (± 0.39)	47.96 (± 3.82)	20.00 (± 1.20)

Table 4: Hits@K (%) results across all datasets, following the HeaRT evaluation setting [15]. The top three results for each metric are highlighted using **first**, **second**, and **third**. *OOM* indicates that the model ran out of memory, while *>24h* denotes that the method did not complete within 24 hours.

Models	Cora		Citeseer		Pubmed	
	MRR	Hits@10	MRR	Hits@10	MRR	Hits@10
<i>GNN + SP len.</i>	14.21 (± 1.44)	33.43 (± 2.69)	20.90 (± 0.79)	47.82 (± 1.11)	7.12 (± 0.41)	5.63 (± 0.52)
<i>Sequence Model</i>	16.86 (± 1.26)	36.03 (± 1.75)	27.45 (± 1.55)	54.20 (± 2.35)	8.58 (± 0.75)	12.87 (± 0.85)
SP4LP	17.27 (± 0.57)	38.52 (± 1.19)	41.08 (± 1.84)	66.28 (± 0.63)	10.87 (± 0.31)	23.01 (± 0.39)

Table 5: Ablation study results (%). MRR and Hits@K with mean and std. deviations over 5 runs with different seeds.

D Experimental Settings

This section outlines the experimental setup used to evaluate all models. We describe the computational resources and the hyperparameter search space. Moreover for SP4LP we include details regarding how the calculation of the shortest path is performed. Details are reported below.

Computational Resources All experiments were conducted on a workstation running Ubuntu 22.04 with an AMD Ryzen 9 7950X CPU (32 threads), 124GB of RAM, and two NVIDIA GeForce RTX 4090 GPUs (24GB each).

Hyperparameters All models are tuned using a grid search over learning rate $\in [1 \times 10^{-2}, 1 \times 10^{-3}]$, dropout $\in [0, 0.7]$, weight decay $\in [0, 10^{-4}, 10^{-7}]$, number of GNN layers $\in \{1, 2, 3\}$, hidden dimensions $\in \{32, 64, 128, 256\}$ and prediction layers $\in \{1, 2, 3\}$. For large-scale datasets, we follow the reduced search space adopted in Li et al. [15] to avoid excessive compute. For SP4LP, we additionally explore the choice of GNN component $\in \{GCN, GraphSAGE, GAT\}$ and sequence model $\in \{LSTM, Transformer\}$, the best models are shown in Table 7. The best hyperparameters are selected based on validation performance. All reported metrics are averaged over 5 different seeds.

Dataset	Cora	Citeseer	Pubmed	ogbl-collab	ogbl-ddi
#Nodes	2,708	3,327	18,717	235,868	4,267
#Edges	5,278	4,676	44,327	1,285,465	1,334,889
Mean Degree	3.90	2.81	4.74	10.90	625.68
Split Ratio	85/5/10	85/5/10	85/5/10	92/4/4	80/10/10

Table 6: Dataset statistics. The split ratio indicates the percentages used for train/validation/test.

Dataset	GNN model	Sequence model
Cora	GCN	Transformer
Citeseer	GCN	LSTM
Pubmed	SAGE	Transformer
ogbl-collab	SAGE	Transformer
ogbl-ddi	GCN	Transformer

Table 7: Best GNN and sequence models selected via hyperparameter tuning.

Shortest path calculation for SP4LP For the computation of shortest paths between node pairs u and v , we used the `shortest_path` function from the `networkx` library [7]. This function returns a single shortest path between two nodes to ensure computational efficiency, even when multiple shortest paths exist. If no path was found between u and v (i.e., they belonged to different connected components), we assigned a synthetic path of length one directly connecting u and v .