DEEP ENSEMBLES FOR GRAPHS WITH HIGHER-ORDER DEPENDENCIES

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ABSTRACT

Graph neural networks (GNNs) continue to achieve state-of-the-art performance on many graph learning tasks, but rely on the assumption that a given graph is a sufficient approximation of the true neighborhood structure. When a system contains higher-order sequential dependencies, we show that the tendency of traditional graph representations to underfit each node’s neighborhood causes existing GNNs to generalize poorly. To address this, we propose a novel Deep Graph Ensemble (DGE), which captures neighborhood variance by training an ensemble of GNNs on different neighborhood subspaces of the same node within a higher-order network representation. We show that DGE consistently outperforms existing GNNs on semisupervised and supervised tasks on six real-world data sets with known higher-order dependencies, even under a similar parameter budget. We demonstrate that diverse and accurate base classifiers are central to DGE’s success, and discuss the implications of these findings for future work on ensembles of GNNs.

1 INTRODUCTION

Graph neural networks (GNNs) solve learning tasks by propagating information through each node’s neighborhood in a graph (Zhou et al., 2020; Wu et al., 2020). Most present work on GNNs assumes that a given graph is a sufficient approximation of the underlying neighborhood structure. But a growing body of work has challenged this assumption by showing that traditional graphs often cannot capture the higher-order structure and dynamics that govern many real-world systems (Lambiotte et al., 2019; Battiston et al., 2020; Porter, 2020; Torres et al., 2021; Battiston et al., 2021). In the present work, we couple GNNs with a specific family of graphs, higher-order networks (HONs), which encode sequential higher-order dependencies (i.e., conditional probabilities that cannot be explained by a first-order Markov model) in a graph structure. A traditional graph, which we call a first-order network (FON), represents a system by decomposing it into a set of pairwise edges, so the only way to infer polyadic interactions is via transitive paths over adjacent nodes. When higher-order dependencies are present, these Markovian paths underfit the true neighborhood (Scholtes, 2017) and can thus produce many false positive interactions between nodes (Lambiotte et al., 2019). To address this limitation, Xu et al. (2016) proposed a HON that creates conditional nodes to more accurately encode the observed higher-order interactions. By preserving this additional information in the graph structure, HONs have produced new insights in studies of user behavior (Chierichetti et al., 2012), citation networks (Rosvall et al., 2014), human mobility and navigation patterns (Scholtes et al., 2014; Peixoto & Rosvall, 2017), the spread of invasive species (Saebi et al., 2020b), anomaly detection (Saebi et al., 2020d), disease progression (Krieg et al., 2020b), and more (Koher et al., 2016; Peixoto & Rosvall, 2017; Scholtes, 2017; Lambiotte et al., 2019; Saebi et al., 2020a). However, their use with GNNs has not been thoroughly explored.

As Figure 1 illustrates, the tendency of FONs to underfit has consequences for GNNs, which typically compute representations by recursively pooling features from each node’s neighbors. In order
Figure 1: A toy example of challenges faced by GNNs in modeling systems with higher-order dependencies. A FON ($G_1$) underfits the higher-order dependencies in the observed paths. Consequently, a GNN will learn similar representations for A and B, since they share the same 2-hop neighborhood in $G_1$. A HON ($G_k$, with $k = 2$ in this example) uses conditional nodes to encode higher-order dependencies. For example, node C|A represents the observed dependency that C only interacts with D when it also interacts with A (note that in real-world systems, $G_k$ rarely breaks the graph into multiple components). However, computing a representation for C then requires a GNN to aggregate multiple local neighborhoods. Colors depict node features.

To maximize GNN performance, we must ensure that local neighborhoods capture the true distribution of interactions in the system. To enable GNNs to utilize the additional information encoded in HONs, we propose a novel Deep Graph Ensemble (DGE), which uses independent GNNs to exploit variance in higher-order node neighborhoods and learn effective representations in graphs with higher-order dependencies. The key contributions of our work include:

1. We analyze the data-level challenges that fundamentally limit the ability of existing GNNs to learn effective models of systems with higher-order dependencies.
2. We introduce the notion of neighborhood subspaces by showing that neighborhoods in a HON are analogous to feature subspaces of first-order neighborhoods. Borrowing from ensemble methods, we then propose DGE to exploit the variance in these subspaces.
3. We experimentally evaluate DGE against eight state-of-the-art baselines on six real-world data sets with known higher-order dependencies, and show that, even with similar parameter budgets, DGE consistently outperforms baselines on semisupervised (node classification) and supervised (link prediction) tasks.
4. We demonstrate that DGE’s ability to train accurate and diverse classifiers is central to strong performance, and show that ensembling multiple GNNs with separate parameters is a consistent way to maximize the trade-off between accuracy and diversity.

2 BACKGROUND AND PRELIMINARIES

2.1 HIGHER-ORDER NETWORKS

Let $S = \{S_1, S_2, ..., S_n\}$ be a set of observed paths (e.g., flight itineraries, disease trajectories, or user clickstreams), where each $S_i = \{s_1, s_2, ..., s_m\}$ is a sequence of entities (e.g., airports, diagnosis codes, or web pages). Let $A = \bigcup S$ denote the set of entities across all sequences. By using a graph to summarize $S$, we can model the global function of each entity in the system and solve a number of useful learning problems. For example, we can predict disease function via node classification or interactions between airports using link prediction. However, there is a large space of possible graphs that can represent $S$. We consider two: a FON, and the HON introduced by Xu et al. (2016). In a FON $G_1 = (V_1, E_1)$, the node set $V_1 = A$ (or, more generally, the mapping $f : V_1 \to A$ is bijective), and the edge set $E_1$ is the set of node pairs $(u, v) \in V_1 \times V_1$ that are adjacent elements in at least one $S_i$.\footnote{Code and 3 data sets are available at https://github.com/sjkrieg/dge.} This is distinct from sequence models like transformers, which typically predict an entity’s local function within a single sequence.
In a HON $G_k = (V_k, E_k)$ with order $k > 1$, each node is a sequence of entities $u' = \langle a'_1, ..., a'_{m-1}, a'_m \rangle$, where each $a'_i \in V_1$ and $m \leq k$. We define $a'_m$ as the base node, and in practice use the notation $u' = a'_m a'_1, ..., a'_{m-1}$ to emphasize that each $u' \in V_k$ represents a base node whose current state is conditioned on a set of predecessors. Each node can have a different number of predecessors, and a conditional node with $m > 1$ is only created if the conditional distribution of paths it encodes sufficiently reduces the entropy of the graph (Saebi et al., 2020d). This means that $\forall v \in V_k \subseteq A^k$; or, more generally, the mapping $f : V_k \to A^k$ is injective but not necessarily bijective. We define $\Omega^k_u = \{u' \in V_k, a'_m = u\}$ as the higher-order family of $u$ (including $u$ itself), and call each $u' \in \Omega^k_u$ a relative of $u$ (in Figure 1, for example, C|A and C|B are the relatives of C). Like $E_k$, the edge set $\hat{E}_k$ is the set of node pairs $(u, v) \in V_k \times V_k$ that are adjacent in at least one $S_j$. In both HONs and FONs, edges are directed such that $(u, v) \neq (v, u)$ and weighted via $w_k : \hat{E}_k \to \mathbb{R}_{>0}$, where 0 indicates a missing edge. By creating conditional nodes, a HON can express higher-order interactions while remaining a graph, since each edge is still a 2-tuple. For example, consider that passengers who fly from Atlanta to Chicago are much more likely to fly back to Atlanta than to New York, and vice versa. A HON can encode this dependency by creating the conditional nodes “Chicago|Atlanta” and “Chicago|New York”, which changes the topology and flow of information within local neighborhoods (Rosvall et al., 2014). Choosing which conditional nodes and edges to create is a non-trivial problem; readers interested in more details can refer to Krieg et al. (2020a), who proposed the procedure that we used in this study.

Related graph-based models The term “higher-order” is also used in the literature to refer to the analysis of polyadic structures within graphs (Benson et al., 2016), as well GNNs that are able to distinguish these structures (Morris et al., 2019; Li et al., 2020; Schnake et al., 2021). These studies rely on the same assumptions as other GNNs, i.e., that a graph is a sufficient approximation of the neighborhood structure. Despite similarities in terminology, HONs are primarily concerned with the question of initial representation (i.e., how should the graph be constructed?) rather than downstream analysis of an existing graph, and thus address a fundamentally different—though complementary—problem (Lambiotte et al., 2019). HONs are also distinct from other formalisms like hypergraphs and simplicial complexes in that they encode conditional distributions that govern higher-order paths (via conditional nodes and directed, weighted edges), and therefore represent different kinds of systems (Battiston et al., 2020; Portesi, 2020; Torres et al., 2021; Battiston et al., 2021). Some very recent works have shown that aggregators based on paths (via random walks) can improve the expressiveness of GNNs (Eliasof et al., 2022; Jin et al., 2022b), but these still rely on the graph structure to guide path sampling. This further motivates the use of representations like HONs, which are designed to consistently and accurately encode higher-order paths for downstream learning tasks (Rosvall et al., 2014; Xu et al., 2016; Saebi et al., 2020d; Krieg et al., 2020a). To our knowledge, only one study has previously used GNNs with HONs (Jin et al., 2022a); however, as we discuss in Section 3, its proposed method has critical shortcomings.

2.2 Graph neural networks

A generic GNN computes a hidden vector representation for a node $u$ at timestamp $t$ according to:

$$h_u^{(t)} = \text{COMBINE}(h_u^{(t-1)}, \text{AGGREGATE}(\{h_v^{(t-1)}, v \in N(u)\})),$$  

(1)

where $N(u)$ is the neighborhood of $u$ in a graph $G$. What typically distinguishes GNNs is how they define COMBINE and AGGREGATE, and how they represent $N(u)$ (Xu et al., 2019; Zhou et al., 2020; Wu et al., 2020). By recursively pooling features via Eq. [1], GNNs implicitly construct higher-order neighborhoods across transitive paths. This allows nonadjacent nodes to share information if they are close in the graph (Li et al., 2018; Chen et al., 2020); however, as we will demonstrate, when higher-order dependencies are present, assuming transitivity leads GNNs that are trained on FONs to generalize poorly.

In this work, we do not reformulate Eq. [1] and are agnostic toward its particular implementation. We instead abstract GNN as a function that takes a single node $u$ as input and returns either the final hidden representation $h_u^{(t)}$, or, in a supervised setting, a vector of predicted label probabilities $\hat{y}_u$:

$$\text{GNN}(u) = h_u^{(t)} \quad \text{or} \quad \text{GNN}(u) = \hat{y}_u.$$  

(2)

We always assume that $G$ contains initial node features $\{x_u, \forall u \in V\}$ and that GNN($\cdot$) is parameterized by weights $\theta$, but for simplicity we omit them from our notation.
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Figure 2: Overview of DGE. Given a HON $G_k$, DGE computes outputs for each base node $u$ by a) resampling relatives from $u$'s higher-order family $\Omega_k^u$ via a sampling distribution $P_k^u$, b) computing outputs for each sampled relative using independent GNN modules, and c) pooling the outputs.

GNN ensembles [Ho (1998) and Breiman (2001)] showed that training an ensemble of shallow learners on random subspaces could exploit variance in the feature space and improve performance, and [Dieterich (2000)] demonstrated that these ensembles are most potent when the predictions of their base classifiers are accurate and diverse. Deep ensembles have typically been used with random weight initializations to improve uncertainty estimation and robustness [Lakshminarayanan et al., 2017; Fort et al., 2019; Wasay & Idreos, 2020], which has benefited GNNs via mechanisms like multi-head attention [Velickovic et al., 2018; Brody et al., 2022; Hou et al., 2021], but very few works have directly explored ensembles of GNNs. Some recent exceptions have suggested that ensembling subgraphs could benefit GNNs [Zeng et al., 2021; Tang et al., 2021; Lin et al., 2022].

3 DEEP GRAPH ENSEMBLES FOR HIGHER-ORDER NETWORKS

3.1 Why ensembles?

There are a number of design challenges (CHs) we must address in order to realize the joint potential of GNNs and HONs. In a HON, entities are represented by a non-fixed number of conditional nodes (CH1); for example, in Figure 1, C is represented by two nodes but A, B, D, and E are each only represented by one node. Conditional nodes typically have different neighborhoods (CH2); for example, in Figure 1, C | A and C | B have different neighbors. Further, they may vary in importance (i.e., degree) in the graph (CH3) [Xu et al., 2016; Saebi et al., 2020d]. To address these challenges, one intuitive idea is to reformulate Eq. 2 so that it computes a representation for $u$ by sampling neighbors from any of $u$'s relatives. However, this fails to address CH2 because a GNN would aggregate the samples without considering differences between relatives. Another idea is to compute representations for each relative separately, then pool them via a permutation-invariant function like an elementwise MEAN, as proposed for HO-GNN by [Jin et al., 2022a]. But this does not address CH3, since all relatives would contribute equally to the final representation. Moreover, if we assume that all relatives in a higher-order family share the same (or similar) features, this solution will overrepresent features associated with larger higher-order families. In order to propose a method that comprehensively addresses these challenges, we first consider the following relationship.

Theorem 1. Let $G_1$ and $G_k$ be a FON and HON, respectively, both constructed from the same input $S$. Let $N_1(u)$ and $N_k(u)$ denote the neighborhoods of any node $u$ in $G_1$ and $G_k$, respectively. Let $\text{AGGREGATE}($·$)$ represent any symmetric neighborhood aggregation function. If $u \in V_1$ and $u' \in \Omega_k^u$, then $\text{AGGREGATE}(N_k(u'))$ is a biased estimator of $\text{AGGREGATE}(N_1(u))$.

We prove Theorem 1 in Appendix A. Intuitively, we observe that HONs are constructed such that $N_k(u') \subseteq N_1(u)$, and $u'$ only exists in $G_k$ if the expectation of a random walker differs substantially (measured via KL-divergence) from $u$ in $G_1$ [Saebi et al., 2020d; Krieg et al., 2020a]. Consequently, these differences in neighborhood structure will shift the expectation of the features gathered by $\text{AGGREGATE}(N_k(u'))$. A typical strategy for training a single GNN would involve attempting to eliminate this variance via some sampling method on the graph. We instead take an ensemble
approach and propose to regularize the model via multiple GNNs that err in different ways. Toward this end, and inspired by feature subspace methods (Ho 1998; Breiman 2001), we call $N_k(u')$ a *neighborhood subspace* of $N_1(u)$.

**Remark** Our result from Theorem 1 also relates to the expressiveness of an aggregator over neighborhoods in $G_k$. Consider two nodes $u, v \in V_1$ whose rooted subgraphs (i.e., neighborhoods) are non-isomorphic but are not distinguishable by the aggregator in $G_1$. As long as there exists some $u' \in \Omega^k_u$ (excluding $u$ itself), then we know that, since it is a biased estimator, the aggregator can distinguish $u'$ from $u$ in $G_k$. Its follows transitivity that the aggregator can also distinguish $u'$ from $v$. It is possible that there also exists some $v' \in \Omega^k_v$ that cannot be distinguished from $u'$, but this would be extremely unlikely to occur over all pairs of relatives in $\Omega^k_u$ and $\Omega^k_v$. We can thus improve the expressiveness of the model by allowing the information from nodes in $\Omega^k_u$ to contribute to the final representation for each $u$.

Guided by these observations, we propose to address the CHs outlined above by training an ensemble of GNNs $\{\text{GNN}_1,\text{GNN}_2,\ldots,\text{GNN}_\ell\}$. Given a set of training nodes $D \subseteq V_1$, we generate bootstraps $\{D^{(1)},D^{(2)},\ldots,D^{(\ell)}\}$ subject to the constraint that $|D^{(i)} \cap \Omega^k_u| = 1$ for all $u \in D$ and $i \leq \ell$ (i.e., each bootstrap contains exactly one relative of each training node). This constraint allows us to avoid the feature overrepresentation problem, address CH1 by sampling with replacement, and address CH2 by training each GNN $i$ on different neighborhood subspaces as represented in $D^{(i)}$.

To solve CH3, we weight the sampling probability for each relative $u'$ according to the normalized out-degree of its higher-order family:

$$P_u^k(u') = \frac{\text{OUTDEG}_k(u')}{\sum_{v' \in \Omega^k_u} \text{OUTDEG}_k(v')},$$

where $u \in D$ and $\text{OUTDEG}_k(u')$ is the weighted out-degree of $u'$ in $G_k$. Because weighted out-degree in $G_k$ is the frequency with which $u'$ appears in $S$ (Krieger et al., 2020a), it is a natural measure of the importance of $u'$ with respect to the rest of the higher-order family $\Omega^k_u$. If $D \subseteq E_1$ consists of node pairs for an edge task, we modify Eq. 3 slightly. For each edge $(u,v) \in D$, we resample a single pair of relatives $(u',v')$ with probability according to the normalized weights of all edges between relatives of $u$ and $v$:

$$P_{u,v}^k(u',v') = \frac{w_k(u',v')}{\sum_{(u'',v'') \in \Omega^k_u \times \Omega^k_v} w_k(u'',v'')}.$$  

### 3.2 Training and inference

Let $D^{(i)}_u$ denote the relative of $u$ that was sampled for the $i$th bootstrap. We consider a supervised or semisupervised setting, in which our goal is to predict class probabilities $\hat{y}_u$ for each $u \in D$ such that some loss is minimized w.r.t. ground truth $y_u$. We propose three methods for computing $\hat{y}_u$:

$$\hat{y}_u = \sigma \left( \text{CONCAT} \left( \{ \text{GNN}_i(D^{(i)}_u), \forall i \leq \ell \}^\top \cdot \mathbf{W} \right) \right) \quad \text{and} \quad \text{GNN}_i(u') = \mathbf{h}^{(i,t)}_{u'},$$

$$\hat{y}_u = \sigma \left( \text{MEAN} \left( \{ \text{GNN}_i(D^{(i)}_u), \forall i \leq \ell \}^\top \cdot \mathbf{W} \right) \right) \quad \text{and} \quad \text{GNN}_i(u') = \mathbf{h}^{(i,t)}_{u'},$$

$$\hat{y}_u = \text{MEAN} \left( \{ \text{GNN}_i(D^{(i)}_u), \forall i \leq \ell \} \right) \quad \text{and} \quad \text{GNN}_i(u') = \mathbf{y}^{(i)}_{u'},$$

where $\sigma$ is a non-linear activation, CONCAT is vector concatenation, MEAN is elementwise mean, $\mathbf{x}^\top$ is the transpose of $\mathbf{x}$, $\mathbf{h}^{(i,t)}_{u'} \in \mathbb{R}^d$ represents the hidden state of node $u'$ in the $t$th (final) layer of GNN$_i$, and $d$ is the number of hidden units (we assume $d$ is fixed for all GNN$_i$). Using $c$ to denote the number of classes, for Eq. 5a we have $\mathbf{W} \in \mathbb{R}^{d \times c}$, and for Eq. 5b we have $\mathbf{W} \in \mathbb{R}^{d \times c}$. We refer to Eq. 5a as DGE-concat, Eq. 5b as DGE-pool, and Eq. 5c as DGE-bag.

In DGE-concat and DGE-pool each GNN outputs hidden representations, which are concatenated or pooled, respectively, before computing the logits. This means that they can be trained in end-to-end fashion as a single neural network with parallel but independent GNN modules. During a forward pass, each GNN only computes representations for the nodes in $D^{(i)}$. In DGE-bag, on the other hand, each GNN outputs class probabilities, which means that each GNN is trained independently.
Figure 2 summarizes the components of DGE: resampling $\ell$ CH1, CH2, and CH3 entirely via batch sampling. Their outputs via Eq. 5c. DGE-batch* thus drops any resemblance to an ensemble, instead addressing constructing $G$ which are trivial compared to the costs of Eq. 2 (Wu et al., 2020). Additionally, the one-time cost of $u$ node $GNN$ (or, in the case of shared parameters, repeating $\ell$ copies of that $GNN$ and the ensemble size $\Omega_u$), the additional cost of sampling and pooling $\Omega_u$ relatives for each sampled relative via Eq. 2, and pooling the computed representations via Eq. 5a, 5b, or 5c. In general, DGE’s computational cost is linear with the cost of computing a node representation for each sampled relative via Eq. 2, and pooling the computed representations via Eq. 5a or 5c. In general, DGE’s computational cost is linear with the cost of the base $GNN$ and the ensemble size $\ell$, since we are essentially constructing $\ell$ copies of that $GNN$ (or, in the case of shared parameters, repeating $\ell$ forward passes per example). For a given node $u$, the additional cost of sampling and pooling $\ell$ relatives is $O(\ell |\Omega_u|)$ and $O(\ell)$, respectively, which are trivial compared to the costs of Eq. 2 (Wu et al., 2020). Additionally, the one-time cost of constructing $G_k$ increases linearly with $k$ (Krieg et al., 2020a). Since we used $G_2$ in our experiments, there was marginal overhead for graph construction as compared to $G_1$.

4 EXPERIMENTAL RESULTS AND DISCUSSION

4.1 EXPERIMENTAL SETUP

We used GrowHON (Krieg et al., 2020a) to construct FONs ($G_1$) and HONs with $k = 2$ ($G_2$) for six real-world data sets with known higher-order dependencies: flight itineraries for airline passengers in the United States (Air) (Rosvall et al., 2014), disease trajectories for type 2 diabetes patients in Indiana (T2D) (Krieg et al., 2020b), clickstreams of users playing the Wikispeedia game (Wiki) (West et al., 2009), readership trajectories for a large online magazine (Mag and a larger version, Mag*) for node classification only (Wang et al., 2020), and global shipping routes (Ship) (Saebi et al., 2020). Table 1 summarizes their key characteristics, including average homophily $H$ (see Appendix G for details). We discuss additional details and preprocessing steps in Appendix B.

We evaluated several baselines, including GCN (Kipf & Welling, 2017), GAT (Veličković et al., 2018), GraphSAGE (Hamilton et al., 2017), GIN (Xu et al., 2019) and GATv2 (Brody et al., 2022), and SEAL (Zhang & Chen, 2018) (link prediction only). Other noteworthy baselines are GraphSAINT (Zeng et al., 2020) which samples a different subgraph for each training iteration (Hu et al., 2020); GCNII (Chen et al., 2020), which uses residual connections to address over-smoothing; and

| Name | $|\mathcal{S}|$ | $|\mathcal{V}_1|$ | $|\mathcal{E}_1|$ | $|\mathcal{V}_2|$ | $|\mathcal{E}_2|$ | # classes | $H(G_1)$ | $H(G_2)$ |
|------|----------|-------------|-------------|-------------|-------------|-----------|-----------|-----------|
| Air  | 17.1m    | 416         | 13,735      | 7,461       | 236,806     | 10        | 0.351     | 0.288     |
| T2D  | 0.8m     | 908         | 314,352     | 5,462       | 367,530     | 16        | 0.099     | 0.099     |
| Wiki | 76.2k    | 4,179       | 70,662      | 5,848       | 84,806      | 10        | 0.366     | 0.375     |
| Mag  | 3.8m     | 4,079       | 1,873,279   | 9,568       | 1,957,602   | 4         | 0.323     | 0.336     |
| Mag* | 8.1m     | 17,428      | 5,098,787   | 192,204     | 7,980,026   | 6         | 0.292     | 0.310     |
| Ship | 54.9k    | 5,586       | 369,952     | 8,586       | 422,426     | 38        | 0.477     | 0.489     |

Table 1: Summary of graphs used in node classification and link prediction experiments.
Table 2: Node classification results (micro F1). Bold font indicates the best result for each data set.

| Model         | Air   | T2D   | Wiki  | Mag   | Mag+  | Ship  |
|---------------|-------|-------|-------|-------|-------|-------|
| GCN           | 0.818 ± 0.03 | 0.480 ± 0.02 | 0.643 ± 0.01 | 0.796 ± 0.01 | 0.710 ± 0.01 | 0.746 ± 0.01 |
| GCNII         | 0.839 ± 0.05 | 0.511 ± 0.02 | 0.654 ± 0.02 | 0.801 ± 0.01 | 0.712 ± 0.01 | 0.770 ± 0.01 |
| GAT           | 0.804 ± 0.03 | 0.282 ± 0.10 | 0.639 ± 0.02 | 0.487 ± 0.06 | 0.511 ± 0.02 | 0.745 ± 0.02 |
| GATv2         | 0.838 ± 0.03 | 0.292 ± 0.07 | 0.643 ± 0.03 | 0.495 ± 0.05 | 0.527 ± 0.02 | 0.747 ± 0.01 |
| GraphSAGE     | 0.781 ± 0.04 | 0.654 ± 0.04 | 0.625 ± 0.02 | 0.808 ± 0.02 | 0.689 ± 0.01 | 0.796 ± 0.01 |
| GIN           | 0.745 ± 0.02 | 0.673 ± 0.04 | 0.636 ± 0.02 | 0.826 ± 0.02 | 0.722 ± 0.00 | 0.801 ± 0.01 |
| GraphSAINT    | 0.802 ± 0.02 | 0.600 ± 0.07 | 0.864 ± 0.01 | 0.821 ± 0.02 | 0.719 ± 0.01 | 0.803 ± 0.01 |
| PathGCN       | 0.846 ± 0.02 | 0.523 ± 0.03 | 0.616 ± 0.03 | 0.735 ± 0.02 | 0.622 ± 0.01 | 0.781 ± 0.01 |
| HONEM         | 0.805 ± 0.04 | 0.566 ± 0.02 | 0.588 ± 0.01 | 0.728 ± 0.02 | 0.620 ± 0.01 | 0.750 ± 0.01 |
| HO-GNN        | 0.822 ± 0.02 | 0.436 ± 0.03 | 0.581 ± 0.01 | 0.785 ± 0.01 | 0.702 ± 0.01 | 0.787 ± 0.01 |
| DGE-concat    | 0.825 ± 0.04 | 0.501 ± 0.06 | 0.615 ± 0.02 | 0.790 ± 0.02 | 0.681 ± 0.04 | 0.828 ± 0.01 |
| DGE-concat*   | 0.810 ± 0.04 | 0.439 ± 0.03 | 0.577 ± 0.02 | 0.761 ± 0.02 | 0.642 ± 0.01 | 0.809 ± 0.01 |
| DGE-pool      | 0.839 ± 0.03 | 0.735 ± 0.03 | 0.671 ± 0.01 | 0.860 ± 0.01 | 0.722 ± 0.01 | 0.808 ± 0.01 |
| DGE-pool*     | 0.865 ± 0.02 | 0.555 ± 0.07 | 0.599 ± 0.04 | 0.775 ± 0.01 | 0.671 ± 0.01 | 0.767 ± 0.02 |
| DGE-bag       | 0.856 ± 0.02 | **0.770 ± 0.04** | **0.681 ± 0.00** | **0.871 ± 0.01** | **0.769 ± 0.01** | **0.840 ± 0.01** |
| DGE-bag*      | 0.766 ± 0.04 | 0.719 ± 0.04 | 0.644 ± 0.02 | 0.841 ± 0.02 | 0.739 ± 0.01 | 0.825 ± 0.01 |
| DGE-batch*    | 0.764 ± 0.03 | 0.646 ± 0.01 | 0.623 ± 0.01 | 0.818 ± 0.01 | 0.742 ± 0.00 | 0.812 ± 0.01 |

* shared parameters

Table 3: Link prediction results (AUPRC). Bold font indicates the best result for each data set.

| Model         | Air   | T2D   | Wiki  | Mag   | Ship  |
|---------------|-------|-------|-------|-------|-------|
| Best baseline | 0.818 ± 0.02 | 0.818 ± 0.00 | 0.834 ± 0.01 | 0.879 ± 0.00 | 0.887 ± 0.01 |
| (SEAL)        | (HONEM)(SEAL) | (HONEM)(SEAL) | (SEAL) | (SEAL) |
| DGE-concat    | **0.886 ± 0.01** | 0.815 ± 0.01 | 0.774 ± 0.01 | 0.802 ± 0.00 | **0.910 ± 0.00** |
| DGE-concat*   | 0.856 ± 0.01 | 0.779 ± 0.01 | 0.712 ± 0.02 | 0.769 ± 0.01 | 0.904 ± 0.00 |
| DGE-pool      | 0.851 ± 0.02 | **0.920 ± 0.00** | 0.838 ± 0.03 | 0.913 ± 0.00 | 0.898 ± 0.00 |
| DGE-pool*     | 0.845 ± 0.01 | 0.901 ± 0.00 | 0.802 ± 0.06 | 0.898 ± 0.00 | 0.815 ± 0.00 |
| DGE-bag       | **0.887 ± 0.00** | 0.907 ± 0.00 | **0.876 ± 0.01** | **0.921 ± 0.00** | 0.895 ± 0.00 |
| DGE-bag*      | 0.862 ± 0.02 | 0.894 ± 0.00 | 0.856 ± 0.00 | 0.916 ± 0.00 | 0.891 ± 0.00 |
| DGE-batch*    | 0.853 ± 0.03 | 0.871 ± 0.01 | 0.830 ± 0.01 | 0.887 ± 0.00 | 0.865 ± 0.00 |

* shared parameters

PathGCN (Eliasof et al., 2022), which learns spatial operators on paths sampled via random walks. We also evaluated two baselines designed specifically for HONs: HONEM, a matrix factorization method (Saebi et al., 2020a), and HO-GNN (Jin et al., 2022a). All baselines used $G_2$ as input (we also evaluated each baseline using $G_2$ as input, details are in Appendix F). We manually tuned each model (details in Appendix F). For DGE, unless noted otherwise, we fixed $\ell = 16$ and used the mean-pooling variant of GraphSAGE as the base GNN, since a) it performed reasonably well as a baseline on all six data sets, and b) its sample-and-aggregate procedure intuitively complements the relative sampling and pooling used by DGE. We used Python 3.7.3 and Tensorflow 2.4.1 for all experiments, and utilized StellarGraph 1.2.1 (Data61, 2018) for the implementation of DGE.

4.2 EXPERIMENTAL RESULTS

Node classification and link prediction On the node classification task (Table 2), DGE-bag outperformed all other methods on five data sets and was second to DGE-pool* on Air. DGE-pool performed only slightly worse than DGE-bag in all cases, but DGE-concat did not generalize well and performed poorly in all cases except Ship. This suggests that our relative sampling procedure was effective in conjunction with ensembling (DGE-bag), but not as effective at regularizing a single fully-connected network against the variance in $G_k$. Since concatenation depends heavily on the position of each relative in the vector representation, DGE-concat overfit on nodes that had many relatives.
phasize the importance of accounting for both the variance in neighborhood subspaces in \( G_k \) as well as the differences in importance between relatives. Some baselines were occasionally competitive: GCNII, GATv2, and PathGCN performed relatively well on Air, and GCNII and GraphSAINT performed relatively well on Wiki. However, on Mag and especially T2D, DGE-bag outperformed all baselines by a significant margin. PathGCN and the full-batch models struggled on T2D, likely due to its high density—meaning that there are many transitive and false positive paths in \( G_k \). Low homophily may also have contributed to their poor performance (Zhu et al., 2020) (see discussion in Appendix G). GAT and GATv2 performed especially poorly on the dense graphs, likely because they compute attention weights (instead of using the given edge weights) and were thus more susceptible to overfitting. The link prediction results, summarized in Table 3, tell a similar story. Because of its strong performance, we focus the remainder of our analysis on DGE-bag. For details on training time and convergence for all models, please see Appendix F.

**Model size**

For the baselines, increasing the number of GNN layers typically increased generalization error (Figure 3). For DGE-bag, the test loss decreased with increased model depth in all but one case (layer 3 on T2D). These results support the findings of Lambiote et al. (2019) that transitivity inferring paths in a FON cannot account for higher-order dependencies, and our hypothesis that GNNs trained on \( G_1 \) will overfit on non-existent paths. The exception to this pattern was Wiki, on which many models performed best with 3 layers. This difference is perhaps because Wiki was the sparsest graph, meaning that the likelihood of sampling a false positive path is relatively low. However, DGE-bag still produced the lowest test error at all depths.

In order to ensure that DGE’s performance was not simply due to using a more expensive model, we compared DGE-bag to the strongest baselines at several parameter budgets (Table 4). For Wiki and Mag are available in Appendix E. DGE-bag underperformed with the smallest budgets, since each base learner was too weak and underfit. We found that sharing parameters (DGE-bag*) resolved this issue and produced strong results on T2D, Wiki, and Mag. However, with a moderate parameter budget, DGE-bag consistently outperformed other models. Again, for most baselines, increasing model depth simply caused them to overfit and decreased performance.

### 4.3 Ensemble Diversity and Parameter Sharing

The variants of DGE that used separate parameters generally performed best. To understand this observation, we draw from prior work on ensembles, which has established that ensembles are most potent when the individual classifiers have low error and high disagreement (Dietterich, 2000). As Figure 3 shows, DGE-bag consistently produced classifiers that were both diverse and accurate. The shared-parameter variants often produced classifiers that were either diverse or accurate, but not both. We observed similar results for Air, Wiki, and Mag (Appendix E). These results reflect node classification performance (Table 2) and support several important conclusions. First, the high disagreement in DGE-bag demonstrates that treating higher-order relatives as neighborhood subspaces successfully introduced variance into the model. Second, the low mean error for DGE-bag demonstrates that ensembling GNNs with separate parameters effectively captured and exploited this variance. Third, sharing parameters (i.e., using a single, more complex model) did not achieve the same
Table 4: Node classification results (mean micro F1 for 5-fold cross validation) under various parameter budgets. Bold font indicates the best result for each budget and data set.

| Model   | Layers | 10k | 50k | 100k | 500k | 1m | 10k | 50k | 100k | 500k | 1m |
|---------|--------|-----|-----|------|------|----|-----|-----|------|------|----|
| GCNII   | 2      | 0.78 | 0.79 | 0.80 | 0.80 | 0.80 | 0.43 | 0.47 | 0.47 | 0.49 | 0.50 |
|         | 4      | 0.77 | 0.78 | 0.78 | 0.79 | 0.79 | 0.43 | 0.47 | 0.50 | 0.51 | 0.51 |
|         | 8      | 0.74 | 0.76 | 0.76 | 0.76 | 0.76 | 0.45 | 0.49 | 0.49 | 0.51 | 0.51 |
| GATv2   | 2      | 0.66 | 0.73 | 0.80 | 0.82 | 0.83 | 0.11 | 0.14 | 0.17 | 0.18 | 0.18 |
|         | 4      | 0.49 | 0.58 | 0.62 | 0.71 | 0.75 | 0.10 | 0.12 | 0.14 | 0.14 | 0.15 |
| GIN     | 2      | 0.74 | 0.74 | 0.75 | 0.75 | 0.75 | 0.54 | 0.64 | 0.67 | 0.66 | 0.67 |
|         | 4      | 0.63 | 0.72 | 0.73 | 0.73 | 0.73 | 0.36 | 0.38 | 0.38 | 0.44 | 0.46 |
| GraphSAINT | 2 | 0.71 | 0.75 | 0.76 | 0.76 | 0.77 | 0.45 | 0.56 | 0.59 | 0.60 | 0.61 |
|         | 4      | 0.60 | 0.64 | 0.68 | 0.74 | 0.74 | 0.25 | 0.27 | 0.28 | 0.28 | 0.28 |
| DGE-bag | 2      | 0.68 | 0.79 | 0.81 | 0.83 | 0.86 | 0.47 | 0.66 | 0.73 | 0.75 | 0.77 |
| DGE-bag*| 2      | 0.70 | 0.76 | 0.80 | 0.81 | 0.81 | 0.65 | 0.70 | 0.70 | 0.71 | 0.71 |

Figure 4: Mean node classification loss for all pairs of the \( \ell = 16 \) base GNNs within each testing fold on T2D, plotted as a function of Cohen’s kappa (lower values indicate lower agreement). Each point represents one pair of GNNs in the ensemble. All plots contain the same number of points.

effect. DGE-pool\(^*\) outperformed DGE-bag on Air because the classifiers were accurate enough to make up for the lack of disagreement. This may be because Air had larger higher-order families than other data sets, so each bootstrap was less representative of the true neighborhood. However, using separate parameters for each GNN—a true ensemble approach—was the most consistently effective way to balance the trade-off between accuracy and diversity.

5 LIMITATIONS, OTHER RELATED WORK, AND FUTURE DIRECTIONS

DGE’s empirical success has implications for multiple research areas. Most broadly, we suggest that ensembling techniques for GNNs have been underexplored. However, generalizing our findings is complicated by the fact that DGE relies on the HON to supply the neighborhood variance, which is in turn limited to representing sequential data. Our results also suggest that the task of designing informed graph representations like HONs is essential and has been previously overshadowed by downstream learning algorithms. Other research on higher-order models has made progress toward this end, but much work remains to generalize these models and the types of dependencies they can represent. One area we consider promising for future work is graph structure learning. At present, HONs are constructed in unsupervised fashion, but future work could overcome this limitation by inferring the graph structure that best solves the given task (Brugere & Berger-Wolf, 2020; Chen & Wu, 2022). Unfortunately, most standard data sets for GNN tasks are pre-constructed graphs, meaning that any higher-order dependencies (sequential or otherwise) have already been lost before learning begins. By integrating graph construction into the learning process and continuing to develop more powerful GNNs, we will increase our capacity to model higher-order relationships and expand the frontier of machine learning on graphs.
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**APPENDIX**

Table 5: Summary of key notation.

| Symbol | Description |
|--------|-------------|
| \(D\)  | set of nodes for training or inference |
| \(D^{(i)}\) | \(i\)th bootstrap of relatives sampled from \(D\) |
| \(D_u^{(i)}\) | relative of node \(u\) sampled for the \(i\)th bootstrap |
| \(E_k\)  | set of edges in \(G_k\) |
| \(G_k\)  | graph with max order \(k\) (\(k = 1\) is a FON, \(k > 1\) is a HON) |
| \(\text{GNN}_i(\cdot)\) | the \(i\)th learner (GNN) in the ensemble |
| \(\mathbf{h}_{u}^{(i,t)}\) | hidden representation of node \(u\) at \(t\)th layer (timestamp) of learner \(i\) |
| \(\ell\)  | number of base learners in the ensemble |
| \(N_k(u)\) | neighborhood of node \(u\) in \(G_k\) |
| \(\Omega_k(u)\) | relatives of node \(u\) in \(G_k\) |
| \(\omega_{k}(u, v)\) | weight of directed edge \((u, v)\) in \(G_k\) |
| \(\hat{y}_u\) | final (pooled) predicted label for node \(u\) |
| \(\hat{y}^{(i)}_u\) | the \(i\)th learner’s predicted label for node \(u\) |
| \(\Omega_k(u)\) | relatives of node \(u\) in \(G_k\) |
A PROOF OF THEOREM[1]

Preliminaries Without loss of generality, in this section we treat the neighborhood $\mathcal{N}_k(u')$ in a graph $\mathcal{G}_k$ for $k \geq 1$ as only the out-neighbors of $u'$ (i.e. $\mathcal{N}_k(u') = \{v \in \mathcal{V}_k, w_k(u', v) > 0\}$) and ignore in-edges. Given a node $u' \in \mathcal{V}_k$, we first define the probability that a random walker over $\mathcal{G}_k$ would move from $u'$ to any $v \in \mathcal{N}_k(u')$:

$$\pi_k(u' \rightarrow v) = \frac{w_k(u', v)}{\text{outdeg}_k(u')}.$$ \hspace{1cm} (6)

Given a FON $\mathcal{G}_1$ and any $u' \in \mathcal{V}_1$, let $u'' \in \Omega^k_u$ be any relative of $u$ in a HON $\mathcal{G}_k$ that is constructed from the same data as $\mathcal{G}_1$. Following [Saebi et al., 2020d], we next define the Kullback-Leibler divergence of $\mathcal{N}_k(u')$ with respect to $\mathcal{N}_1(u)$:

$$D_{KL}(\mathcal{N}_k(u') \parallel \mathcal{N}_1(u)) = \sum_{v \in \mathcal{N}_1(u)} \pi_1(u \rightarrow v) \log_2 \frac{\pi_1(u \rightarrow v)}{\pi_k(u' \rightarrow v)}.$$ \hspace{1cm} (7)

Our notation throughout this section assumes that $v \in \mathcal{N}_1(u)$ and $v \in \mathcal{N}_k(u')$; however, a relative $v'' \in \Omega^k_u$ often replaces $v$ in $\mathcal{N}_k(u')$. For example, in Figure [1] C|A is substituted for its base node C in the neighborhood of A, but the edge (A, C|A) still fundamentally represents a step from C to A. GrowHON ([Krieg et al., 2020a]) constructs $\mathcal{G}_k$ such that $u' \in \mathcal{V}_k$ if the following inequality holds:

$$D_{KL}(\mathcal{N}_k(u') \parallel \mathcal{N}_1(u)) > \frac{m}{\log_2(1 + \text{freq}(u'))},$$ \hspace{1cm} (8)

where $m > 1$ is length of the sequence encoded by $u'$ (see Section 2) and $\text{freq}(u') \geq 1$ is the number of times $u'$ is found in all the observed paths in $\mathcal{S}$.

Lemma 1. If $u \in \mathcal{V}_1$ and $u' \in \Omega^k_u$, then there exists at least one node $v \in \mathcal{N}_1(u)$ such that $\pi_1(u \rightarrow v) \neq \pi_k(u' \rightarrow v)$.

Proof. If $u' \in \Omega^k_u$, then $D_{KL}(\mathcal{N}_k(u') \parallel \mathcal{N}_1(u)) > 0$. This is because $\text{freq}(u') \geq 1$, so $\log_2(1 + \text{freq}(u')) \geq 1$ and the right side of Eq. [8] cannot be negative. If $D_{KL}(\mathcal{N}_k(u') \parallel \mathcal{N}_1(u)) = 0$, then the inequality in Eq. [8] cannot hold, so $u' \notin \mathcal{V}_k$ and, consequently, $u' \notin \Omega^k_u$. Additionally, if $\pi_1(u \rightarrow v) = \pi_k(u' \rightarrow v)$ for all $v \in \mathcal{N}_1(u)$, then by Gibbs’ inequality we have $D_{KL}(\mathcal{N}_k(u') \parallel \mathcal{N}_1(u)) = 0$. Therefore, since $u' \in \Omega^k_u$, there must exist at least one node $v \in \mathcal{N}_1(u)$ such that $\pi_1(u \rightarrow v) \neq \pi_k(u' \rightarrow v)$.

We now restate and prove Theorem[1]

Theorem[1]. Let $\mathcal{G}_1$ and $\mathcal{G}_k$ be a FON and HON, respectively, both constructed from the same input $\mathcal{S}$. Let $\mathcal{N}_1(u)$ and $\mathcal{N}_k(u)$ denote the neighborhoods of any node $u$ in $\mathcal{G}_1$ and $\mathcal{G}_k$, respectively. Let $\text{AGGREGATE}(\cdot)$ represent any symmetric neighborhood aggregation function. If $u \in \mathcal{V}_1$ and $u' \in \Omega^k_u$, then $\text{AGGREGATE}(\mathcal{N}_k(u'))$ is a biased estimator of $\text{AGGREGATE}(\mathcal{N}_1(u))$.

Proof. Without loss of generality, we consider the case in which AGGREGATE operates on a sample of neighbors (i.e., GraphSAGE [Hamilton et al., 2017]). Let $V \sim \mathcal{N}_k(u')$ denote a random sample drawn from $\mathcal{N}_k(u')$. Assuming we weight the sampling distribution according to edge weights, we can write the probability of sampling any node $v$ as

$$p(V = v) = \frac{w_k(u', v)}{\text{outdeg}_k(u')} = \pi_k(u' \rightarrow v),$$ \hspace{1cm} (9)

where $w_k(u', v)$ is the weight of edge $(u', v)$ in $\mathcal{G}_k$ (Section 2) and $\text{outdeg}_k(u')$ is the weighted out-degree of $u'$ in $\mathcal{G}_k$. If we say that each $u \in \mathcal{V}_k$ is represented by a real-valued feature vector $x^k_u = [x_{u,1}^{(k)}, x_{u,2}^{(k)}, \ldots, x_{u,d}^{(k)}]$, then, for a single sample drawn from $\mathcal{N}_k(u')$, we can formulate the expected value for each vector as follows:

$$E_{V \sim \mathcal{N}_k(u')}[x^k_v] = p(V = v) x^k_v.$$ \hspace{1cm} (10)
If we combine these feature vectors for each of the \( n \) nodes in \( V_k \), then we can represent the full expectation of \( V \) as a neighborhood matrix

\[
E_{V \sim N_k(u')}[V] = \begin{bmatrix}
    p(V = 1) x_1^k \\
p(V = 2) x_2^k \\
\vdots \\
p(V = n) x_n^k \\
p(V = 1) x_{1,1}^{(k)} & p(V = 1) x_{1,2}^{(k)} & \cdots & p(V = 1) x_{1,d}^{(k)} \\
p(V = 2) x_{2,1}^{(k)} & p(V = 2) x_{2,2}^{(k)} & \cdots & p(V = 2) x_{2,d}^{(k)} \\
\vdots & \vdots & \ddots & \vdots \\
p(V = n) x_{n,1}^{(k)} & p(V = n) x_{n,2}^{(k)} & \cdots & p(V = n) x_{n,d}^{(k)}
\end{bmatrix}.
\]

We can simplify our notation by assuming the initial feature vectors are identity features (i.e., one-hot encodings of the node indices) defined on the base nodes, i.e., for features of nodes in \( V_1 \), we have \( x_{u,j}^{(1)} = \delta_{uj} \) for all \( u, j \leq d = n \), where \( \delta \) is the Kronecker delta. For the features of nodes in \( V_k \), we assume that each \( u' \in \Omega_k \) uses the same features as its base node, i.e., \( x_{u',j}^{(k)} = \delta_{uj} \).

Substituting these values in the above matrix gives

\[
E_{V \sim N_k(u')}[V] = \begin{bmatrix}
p(V = 1) & 0 & \cdots & 0 \\
0 & p(V = 2) & \cdots & 0 \\
\vdots & \vdots & \ddots & \vdots \\
0 & 0 & \cdots & p(V = n)
\end{bmatrix}.
\]

Without loss of generality, let \( \text{AGGREGATE} \) be the feature-wise \( \text{MEAN} \) of \( N \) samples drawn from \( N_k(u') \). We can then represent the expectation of \( \text{AGGREGATE} \) as

\[
E_{V \sim N_k(u')}[\text{AGGREGATE}(V)] = \begin{bmatrix}
    \frac{1}{N} Np(V = 1) & 0 & \cdots & 0 \\
    0 & Np(V = 2) & \cdots & 0 \\
    \vdots & \vdots & \ddots & \vdots \\
    0 & 0 & \cdots & Np(V = n)
\end{bmatrix}
= \begin{bmatrix}
p(V = 1) \\
p(V = 2) \\
\vdots \\
p(V = n)
\end{bmatrix}
= \begin{bmatrix}
    \pi_k(u' \rightarrow 1) \\
    \pi_k(u' \rightarrow 2) \\
    \vdots \\
    \pi_k(u' \rightarrow n)
\end{bmatrix}.
\]

(via Eq. 9)

If \( V \sim N_1(u) \), we instead have

\[
E_{V \sim N_1(u')}[\text{AGGREGATE}(V)] = \begin{bmatrix}
    \pi_1(u \rightarrow 1) \\
    \pi_1(u \rightarrow 2) \\
    \vdots \\
    \pi_1(u \rightarrow n)
\end{bmatrix}.
\]

From Lemma 1 we know that there exists at least one \( v \in N_1(u) \) such that \( \pi_1(u \rightarrow v) \neq \pi_k(u' \rightarrow v) \). Therefore, \( E_{V \sim N_1(u')}[\text{AGGREGATE}(V)] \neq E_{V \sim N_1(u)}[\text{AGGREGATE}(V)] \), and \( \text{AGGREGATE}(N_k(u')) \) is a biased estimator of \( \text{AGGREGATE}(N_1(u)) \).
This result holds for other common aggregators like SUM and MAX for single samples. For MAX, however, the sample size matters, because the expected value for the $j^{th}$ feature is the probability that $j$ is sampled at least once. We conjecture that this is problematic for larger sample sizes in weighted graphs, and is perhaps the reason why mean-pooling outperformed max-pooling for our GraphSAGE baseline during our initial experiments (Section C).
B Data Details and Experimental Setup

Air: Flight trajectories of passenger itineraries in the United States. Nodes represent airport locations (cities) and edges represent passengers flying between locations. The itineraries were retrieved from the Airline Origin and Destination Survey (DB1B) database, which is publicly available through the U.S. Bureau of Transportation Statistics. We downloaded all records between Jan. 1 and Dec. 31, 2019 from the DB1BCoupon table, joined and sorted itineraries using the “ITIN_ID” and “SEQ_NUM” columns, and discarded any itineraries with missing origin or destination information. Node classes represent the geographical location of each airport, aggregated by standard federal region of the United States. While we extracted this particular set of paths, this database has been utilized in prior work on HONs [Rosvall et al. 2014, Scholtes 2017].

T2D: Disease trajectories for type 2 diabetes patients in the state of Indiana [Krieg et al. 2020b]. Nodes represent ICD9 diagnosis codes and edges represent sequential diagnoses. Following prior work, we only preserved the first occurrence of each diagnosis code for each patient; however, we did not split trajectories based on the period of time between diagnoses. Node classes are the chapters of each ICD9 code, which represent categories of disease classification.

Wiki: Clickstreams of users playing the Wikispeedia game, in which a player attempts to navigate from a source to a target article by clicking only Wikipedia links [West et al. 2009]. Here, nodes represent Wikipedia articles, and edges represent user clicks between articles. We included both finished and unfinished paths. Node classes are the subjects of each article.

Mag: Readership trajectories for a large online magazine from Jan 1. to Apr. 15, 2020 [Wang et al. 2020]. Nodes represent online content (articles, games, etc.), and edges represent sequential clicks by users. We only considered trajectories in which the user visited at least three nodes during a session. Node classes represent the content type (magazine, culture, news, or humor).

Mag+: An expanded version of Mag which also includes data from July 1, 2019 through Dec. 31, 2019, as well as two additional content types (books, home). Due to its size and similarity to Mag, we excluded it from link prediction experiments.

Ship: Trajectories of global shipping activity [Saebi et al. 2020c]. Nodes represent ports, and edges represent ships traveling between ports. Node classes are provinces, as defined by the Marine Ecoregions of the World (MEOW) system for classifying oceans and waterways [Spalding et al. 2007].

For node classification, we used stratified 5-fold cross validation [Shchur et al. 2018] and reported the mean micro F1-score. For link prediction, we generated training and testing sets by randomly sampling 10% of positive node pairs and an equal number of negative node pairs. In addition to hiding all edges (in both directions) between testing pairs, for experiments using G2 we also hid all edges between any pair of nodes in the same higher-order families as each testing pair. We repeated each experiment five times and reported the mean area under the precision-recall curve (AUPRC) [Yang et al. 2015]. All experiments utilized the same training and testing splits. For models trained on G1, we used identity features (i.e., one-hot encodings of the node indices) for each node. For models trained on G2, we used the same features as G1 such that each $u' \in \Omega_u^2$ shared the same features as its base node $u$. 

[https://transtats.bts.gov/] Accessed April 14, 2022.
C  MODEL TUNING AND HYPERPARAMETER SETUP

We manually tuned hyperparameters for each model. Table 6 summarizes the configurations that we evaluated in reporting the main results in Tables 2 and 3. We used the Adam optimizer with a learning rate of 0.0005 for GAT and GATv2, and a learning rate of 0.01 for all other models. For minibatch models (GraphSAGE, GIN, and DGE), we tested batch sizes of 16, 32, and 64. Unless otherwise noted, all results reported in Tables 2, 3, 7, and 8 were the best-performing configuration for each model as averaged across all testing folds.

In general, the most impactful hyperparameter was the number of GNN layers. As discussed in Section 4.2 on Air, T2D, and Mag, all baselines performed best with only a single layer for both node classification and link prediction. On Wiki, several models performed best with 3 layers (Figure 3). GCNII, which uses residual connections and is thus designed with deep GNNs in mind (Chen et al., 2020), was the only model we tested with more than 3 layers. For most models, increasing the number of hidden units per layer above 256 had little to no effect on performance. The exception to this was DGE-pool* on Air, which performed best with 2,048 hidden units (Section 4.3).

Other hyperparameters varied by data set. The number of neighbors sampled ($|\mathcal{N}|$) was very important on the dense graphs (T2D and Mag). In all cases, we found that increasing the sample size of the first layer was the most important, so we fixed the sample size at the second layer to 1 (per first-layer sample). Baselines preferred 512 samples for T2D and 256 for Mag. On the other data sets, 64 neighbor samples was sufficient for all models. DGE preferred 256 samples per base learner (the max that we tested) for T2D and 128 for Mag. The same trend held for GraphSAINT’s subgraph sampling, which preferred more roots on graphs that were dense and had more nodes (512 for Air and Wiki, 1024 for T2D, and 2048 for Mag, 4096 for Mag+, and 2048 for Ship). On all data sets, PathGCN performed best with at least 100 random walk samples per node.

For neighborhood aggregators, max-pooling was in all cases inferior to mean-pooling and sum-pooling, likely due to the inability of max-pooling to appropriately distinguish dense neighborhoods (Xu et al., 2019).

### Table 6: Hyperparameters evaluated for each model.

| Model         | # Layers | Other                  |
|---------------|----------|------------------------|
| GCN           | {1, 2, 3} |                        |
| GCNII         | {1, 2, 3, 4, 6, 8} | $\alpha = 0.5, \lambda = 1.0$ |
| GAT           | {1, 2, 3} | attnheads = {4, 8, 16} |
| GATv2         | {1, 2, 3} | attnheads = {4, 8, 16} |
| GraphSAGE     | {1, 2, 3} | $|\mathcal{N}| = \{32, 64, 128, 256, 512, 1024\} \times \{1, 2, 4, 8\} \times \{1\}$ AGGREGATE = {MEANPOOL, MAXPOOL} |
| GIN           | {1, 2, 3} | $|\mathcal{N}| = \{32, 64, 128, 256, 512, 1024\} \times \{1, 2, 4, 8\} \times \{1\}$ |
| GraphSAINT    | {1, 2, 3} | nroots = {256, 512, 1024, 2048, 4096}, walklen = {2, 3, 4} |
| PathGCN       | {1, 2, 3} | nwalks = {10, 50, 100, 200}, walklen = {2, 3, 4, 5} |
| HO-GNN        | {1, 2, 3} |                        |
| SEAL          | {1, 2, 3} | hops = {1, 2, 3} |
| DGE (all)     | {1, 2, 3} | $\ell = \{4, 8, 16\}$, $|\mathcal{N}| = \{32, 64, 128, 256\} \times \{1, 2, 4\} \times \{1\}$ |

### General hyperparameters

| Model | Dropout | Hidden units (per layer) |
|-------|---------|--------------------------|
| All   | 0.4     | {128, 256, 512, 1024, 2048} |
D  ADDITIONAL POOLING MECHANISMS

In addition to Eqs. 5a, 5b, and 5c we evaluated an attention mechanism (inspired by GAT (Veličković et al., 2018)) which used self-attention to pool the outputs of all sampled relatives. We computed attention coefficients $e_{ij}$ for each $i, j \leq \ell$ according to

$$e_{ij} = F\left(W(GNN_i(D_u^{(i)})), W(GNN_j(D_u^{(j)}))\right),$$  \hspace{1cm} (11)$$

where $F$ is a single-layer feed-forward network with a LeakyReLU activation, $W \in \mathbb{R}^{2d \times d}$ is a trainable weight matrix, and $GNN_i(u) = h_i^{(i,\ell)}$ (as in Eqs. 5a and 5b). We then computed the final attention coefficients $\alpha_{ij}$ according to

$$\alpha_{ij} = \text{softmax}_i(e_{ij}) = \frac{\exp(e_{ij})}{\sum_{m=1}^{\ell} \exp(e_{im})}. \hspace{1cm} (12)$$

We repeated this procedure for $K$ separate attention heads. Whereas the original GAT layer averages the outputs of the attention heads before applying a non-linear activation (Veličković et al., 2018), we found that passing the same output to another feed-forward network provided better results. Formally, letting $i$ be the index of we compute the vector of class probabilities $\hat{y}_u$ by

$$\hat{y}_u = \sigma\left(F'\left(\sigma'\left(\frac{1}{K} \sum_{n=1}^{K} \left\| \sum_{i=1}^{\ell} \alpha_{ij}^{(n)} W^{(n)} GNN_j(D_u^{(j)}) \right\| \right)\right)\right),$$  \hspace{1cm} (13)$$

where $\|$ is vector concatenation, $F'$ is a feed-forward network with output dimension $d/4$ (followed by batch normalization and a LeakyReLU activation), $\alpha_{ij}^{(n)}$ and $W^{(n)}$ represent the attention coefficients and linear transformation, respectively, for the $n^{th}$ attention head, and $\sigma'$ is a LeakyReLU activation. For our experiments, we set $K = 5$, $d = 128$, and a negative slope coefficient of 0.3 for all LeakyReLU activations. We refer to this model as DGE-attn and discuss its performance in Appendix E.
E ADDITIONAL EXPERIMENTAL RESULTS

To support our claim that existing GNNs cannot effectively use $G_2$ as input, we also tested baselines using $G_2$ as the input graph for Air, T2D, Wiki, and Mag. As we expected, because each neighborhood in $G_2$ is only a subspace of its neighborhood in $G_1$, performance was generally worse (or about the same) when compared to $G_1$. Tables 7 and 8 detail the full results.

Table 7 also includes results for a GNN ensemble (DGE-bag) trained on $G_1$, i.e. without relative sampling and pooling. Overall, we found that the performance was slightly better than GraphSAGE (the base GNN), but in all cases fell short of DGE-bag with relative sampling and pooling. The final result included in Table 7 is for one additional pooling mechanism, DGE-attn, which pooled representations from the sampled relatives via a self-attention mechanism. Since it did not generalize particularly well in the node classification experiments, we did not discuss it in the main manuscript and did not evaluate its performance on link prediction. More details are available in Appendix D.

Table 8 also includes the full link prediction results for baselines, which we excluded from Table 3 because of space constraints in the main manuscript.

Figure 5 shows DGE-bag’s performance as a function of the number of base learners ($\ell$) on four data sets. We found that 8-12 base learners produced consistently strong results. Figure 6 shows classifier accuracy and diversity (Section 4.3) for the T2D, Wiki, and Mag data sets, as a supplement to Figure 4.

Table 7: Additional node classification results (micro F1) for baselines using $G_2$ as input. Bold font indicates the best result for each data set.

| Model       | Input          | Air          | T2D          | Wiki         | Mag          |
|-------------|----------------|--------------|--------------|--------------|--------------|
| GCN         | $G_1$          | 0.818 ± 0.03 | 0.480 ± 0.02 | 0.643 ± 0.01 | 0.796 ± 0.01 |
|             | $G_2$          | 0.805 ± 0.05 | 0.500 ± 0.02 | 0.665 ± 0.02 | 0.744 ± 0.02 |
| GCNII       | $G_1$          | 0.845 ± 0.05 | 0.511 ± 0.02 | 0.654 ± 0.02 | 0.801 ± 0.01 |
|             | $G_2$          | 0.833 ± 0.05 | 0.537 ± 0.03 | 0.657 ± 0.02 | 0.816 ± 0.01 |
| GAT         | $G_1$          | 0.804 ± 0.03 | 0.282 ± 0.10 | 0.639 ± 0.02 | 0.487 ± 0.06 |
|             | $G_2$          | 0.817 ± 0.05 | 0.253 ± 0.03 | 0.646 ± 0.03 | 0.512 ± 0.04 |
| GATv2       | $G_1$          | 0.838 ± 0.03 | 0.292 ± 0.07 | 0.643 ± 0.03 | 0.495 ± 0.05 |
|             | $G_2$          | 0.842 ± 0.05 | 0.241 ± 0.10 | 0.646 ± 0.03 | 0.512 ± 0.13 |
| GraphSAGE   | $G_1$          | 0.781 ± 0.04 | 0.654 ± 0.04 | 0.625 ± 0.02 | 0.808 ± 0.02 |
|             | $G_2$          | 0.688 ± 0.02 | 0.516 ± 0.01 | 0.633 ± 0.01 | 0.818 ± 0.01 |
| GIN         | $G_1$          | 0.745 ± 0.02 | 0.673 ± 0.04 | 0.636 ± 0.02 | 0.826 ± 0.02 |
|             | $G_2$          | 0.660 ± 0.02 | 0.637 ± 0.03 | 0.625 ± 0.02 | 0.812 ± 0.02 |
| GraphSAINT  | $G_1$          | 0.802 ± 0.02 | 0.600 ± 0.07 | 0.664 ± 0.01 | 0.821 ± 0.02 |
|             | $G_2$          | 0.710 ± 0.03 | 0.567 ± 0.03 | 0.661 ± 0.02 | 0.828 ± 0.01 |
| HONEM       | $G_1, G_2$     | 0.805 ± 0.04 | 0.566 ± 0.02 | 0.588 ± 0.01 | 0.728 ± 0.02 |
| DGE-bag     | $G_1$          | 0.772 ± 0.05 | 0.694 ± 0.04 | 0.645 ± 0.02 | 0.831 ± 0.01 |
|             | $G_2$          | 0.825 ± 0.04 | 0.501 ± 0.06 | 0.615 ± 0.02 | 0.790 ± 0.02 |
| DGE-concat  | $G_2$          | 0.810 ± 0.04 | 0.439 ± 0.03 | 0.577 ± 0.02 | 0.761 ± 0.02 |
| DGE-concat* | $G_2$          | 0.839 ± 0.03 | 0.735 ± 0.03 | 0.671 ± 0.01 | 0.860 ± 0.01 |
| DGE-pool    | $G_2$          | 0.865 ± 0.02 | 0.555 ± 0.07 | 0.599 ± 0.04 | 0.775 ± 0.01 |
| DGE-pool*   | $G_2$          | 0.856 ± 0.02 | 0.770 ± 0.04 | 0.681 ± 0.00 | 0.871 ± 0.01 |
| DGE-bag     | $G_2$          | 0.766 ± 0.04 | 0.719 ± 0.04 | 0.644 ± 0.02 | 0.841 ± 0.02 |
| DGE-bag*    | $G_2$          | 0.764 ± 0.03 | 0.646 ± 0.01 | 0.623 ± 0.01 | 0.818 ± 0.01 |
| DGE-attn    | $G_2$          | 0.829 ± 0.03 | 0.549 ± 0.05 | 0.618 ± 0.02 | 0.775 ± 0.05 |

* shared parameters
Table 8: Full link prediction results (AUPRC) for baselines. Bold font indicates the best result for each data set.

| Model         | Input | Air | T2D | Wiki | Mag | Ship |
|---------------|-------|-----|-----|------|-----|------|
| GCN           | $G_1$ | 0.794 ± 0.01 | 0.793 ± 0.00 | 0.741 ± 0.01 | 0.742 ± 0.00 | 0.849 ± 0.00 |
|               | $G_2$ | 0.777 ± 0.02 | 0.781 ± 0.00 | 0.749 ± 0.01 | 0.753 ± 0.00 |       |
| GCNII         | $G_1$ | 0.805 ± 0.01 | 0.754 ± 0.01 | 0.781 ± 0.01 | 0.750 ± 0.00 | 0.848 ± 0.01 |
|               | $G_2$ | 0.771 ± 0.01 | 0.732 ± 0.00 | 0.790 ± 0.01 | 0.759 ± 0.00 |       |
| GAT           | $G_1$ | 0.786 ± 0.01 | 0.698 ± 0.02 | 0.794 ± 0.01 | 0.644 ± 0.00 | 0.764 ± 0.00 |
|               | $G_2$ | 0.768 ± 0.01 | 0.681 ± 0.01 | 0.801 ± 0.01 | 0.641 ± 0.00 |       |
| GATv2         | $G_1$ | 0.771 ± 0.01 | 0.701 ± 0.01 | 0.797 ± 0.01 | 0.645 ± 0.01 | 0.797 ± 0.00 |
|               | $G_2$ | 0.784 ± 0.01 | 0.684 ± 0.00 | 0.796 ± 0.01 | 0.660 ± 0.01 |       |
| GraphSAGE     | $G_1$ | 0.782 ± 0.02 | 0.713 ± 0.01 | 0.758 ± 0.01 | 0.717 ± 0.00 | 0.820 ± 0.00 |
|               | $G_2$ | 0.757 ± 0.02 | 0.712 ± 0.01 | 0.756 ± 0.01 | 0.716 ± 0.00 |       |
| GIN           | $G_1$ | 0.800 ± 0.01 | 0.704 ± 0.00 | 0.745 ± 0.01 | 0.717 ± 0.00 | 0.782 ± 0.00 |
|               | $G_2$ | 0.748 ± 0.02 | 0.702 ± 0.00 | 0.743 ± 0.01 | 0.716 ± 0.00 |       |
| GraphSAINT    | $G_1$ | 0.718 ± 0.01 | 0.797 ± 0.00 | 0.688 ± 0.01 | 0.750 ± 0.00 | 0.862 ± 0.01 |
|               | $G_2$ | 0.702 ± 0.02 | 0.791 ± 0.01 | 0.694 ± 0.01 | 0.756 ± 0.00 |       |
| SEAL          | $G_1$ | 0.818 ± 0.02 | 0.754 ± 0.01 | 0.834 ± 0.01 | 0.751 ± 0.00 | 0.887 ± 0.01 |
| HONEM         | $G_1$ | 0.697 ± 0.02 | 0.818 ± 0.00 | 0.589 ± 0.01 | 0.769 ± 0.00 | 0.815 ± 0.00 |
|               | $G_2$ | 0.811 ± 0.00 | 0.789 ± 0.00 | 0.820 ± 0.02 | 0.879 ± 0.00 | 0.856 ± 0.00 |

* shared parameters

Table 9: Node classification results (micro F1) under various parameter budgets. Bold font indicates the best result for each budget and data set.

| Model         | Layers | Total parameters (Wiki) | Total parameters (Mag) |
|---------------|--------|-------------------------|------------------------|
|               |        | 50k | 100k | 500k | 1m | 50k | 100k | 500k | 1m |
| GCNII         | 2      | 0.60 | 0.60 | 0.60 | 0.60 | 0.78 | 0.79 | 0.79 | 0.79 |
|               | 4      | 0.58 | 0.61 | 0.62 | 0.63 | 0.79 | 0.80 | 0.80 | 0.80 |
|               | 8      | 0.60 | 0.61 | 0.62 | 0.62 | 0.77 | 0.79 | 0.79 | 0.79 |
| GATv2         | 2      | 0.46 | 0.52 | 0.61 | 0.62 | 0.26 | 0.30 | 0.37 | 0.40 |
|               | 4      | 0.40 | 0.47 | 0.51 | 0.55 | —   | —   | —   | —   |
| GIN           | 2      | 0.59 | 0.61 | 0.62 | 0.62 | 0.82 | 0.82 | 0.83 | 0.83 |
|               | 4      | 0.50 | 0.54 | 0.61 | 0.62 | 0.66 | 0.76 | 0.79 | 0.79 |
| GraphSAINT    | 2      | 0.61 | 0.64 | 0.66 | 0.66 | 0.79 | 0.79 | 0.81 | 0.82 |
|               | 4      | 0.55 | 0.62 | 0.66 | 0.66 | 0.54 | 0.65 | 0.70 | 0.74 |
| DGE-bag       | 2      | 0.52 | 0.58 | 0.68 | 0.69 | 0.81 | 0.84 | 0.86 | 0.87 |
| DGE-bag*      | 2      | 0.63 | 0.64 | 0.65 | 0.65 | 0.83 | 0.84 | 0.85 | 0.85 |

* shared parameters
Figure 5: Node classification performance for DGE-bag as a function of the number of base learners ($\ell$).

Figure 6: Mean node classification loss of each pair of classifiers, plotted as a function of Cohen’s kappa (lower values indicate lower agreement). Each point represents one pairwise comparison between the 16 classifiers in each of the 5 testing folds. All plots contain the same number of points.
F TRAINING TIME AND CONVERGENCE

We analyzed the additional time cost incurred by DGE by plotting training and testing loss as a function of total runtime for Air and T2D. We used an NVIDIA GeForce GTX TITAN X GPU and dual 8-core 2.4GHz Intel Xeon processors for each model. As Figure 7 shows, DGE converged more slowly but generalized significantly better than all baselines. While the baselines typically converged quickly on the training set, they also rapidly began to overfit on the testing set. The increase in test error was sharper for the full-batch models like GCN and GCNII since they completed more epochs in a shorter period of time. DGE-bag, and, to a lesser extent, DGE-pool, generalized extremely well. These observations reflect overall model performance as reported in Table 2. One difference is that many of the baseline results reported in Table 2 used only a single GNN layer (Section 4.2). For the results in Figure 7 we fixed the number of GNN layers to 2 in order to confirm that the poor generalization of baselines (Section 4.2) was not due to lack of model capacity or underfitting the training set. This is why some models, like GCNII on Air, were relatively competitive in the main results but show high generalization error here.

The sample-based methods, including DGE, incurred significant overhead from neighbor sampling (over 95% of total runtime). This was exacerbated by the fact that all models preferred a high number of neighbor samples at the first layer, so repeatedly sampling a single neighbor at the second layer (Appendix C) produced substantial procedural overhead. DGE would train significantly faster with a more efficient mechanism for sampling these two-hop neighbors (such as pre-fetching).

All variants of DGE converged in fewer epochs than baselines, likely because DGE sees several subgraphs for each node during the same epoch (by virtue of being an ensemble) and because $G_e$ is sparser than $G_1$. On T2D, convergence took approximately 100 epochs for DGE, 100-150 for minibatch baselines, and 1000-1500 for full-batch baselines. On other data sets, DGE generally converged in approximately 15-20 epochs, minibatch baselines converged in 20-30 epochs, and other full-batch baselines took 100-200 epochs (except Mag and Mag+, which took the full-batch models 500+ epochs).

![Figure 7](image-url)

(a) Air results.

![Figure 7](image-url)

(b) T2D results.

Figure 7: Node classification loss for 2-layer models on the first testing fold, shown as a function of training time. Runtime is measured as wallclock time from the start of the first epoch. High loss values are truncated to preserve scale. Similar trends held for all testing folds.
G  ANALYSIS OF NODE CHARACTERISTICS

To better understand DGE’s performance, we analyzed the node classification predictions by plotting the change in test loss for each node (Figure 8). We hypothesized that DGE would provide the strongest improvements on nodes with higher out-degree and larger higher-order families (\(|\Omega^2|\)); however, this was not consistently true for all data sets. Instead, we found that a node’s homophily (\(H\), defined as the fraction of neighbors that have the same class) was the best indicator of performance improvement. This observation is noteworthy for future work, especially since most GNNs struggle to model graphs with low homophily [Zhu et al., 2020].

Figure 8: Delta in test loss for DGE-bag compared to GraphSAGE (the base GNN) for node classification (i.e., values < 0 mean that GraphSAGE produced higher loss than DGE-bag), plotted as a function of key node features in \(G_1\). Out-degree and in-degree are unweighted. Each point represents a node in the test set. All five testing folds are included. Shading represents a 95% CI for the regression line.
We used GrowHON (Krieg et al., 2020a) to construct all HONs. GrowHON provides a parameter $\tau$, which helps regularize the graph by raising the KL-divergence threshold required for a new higher-order node to be created, and we found that the choice of $\tau$ impacted DGE’s success in each task. Figure 9 shows the results of varying $\tau$ for each data set and task. In general, the optimal $\tau$ value varied between data sets for classification tasks, while for link prediction, a lower $\tau$ generally yielded better results. We did not tune DGE’s hyperparameters for these experiments, so it is likely that further tuning would reduce the difference between each model. However, these observations have important consequences for future work, especially with respect to integrating prior studies on graph structure learning, which have argued that different graph representations may be useful for different predictive tasks (Brugere & Berger-Wolf, 2020). Ideally, we could incorporate HON construction into the learning framework so that the model could learn the structure that is best-suited to each task.

Figure 9: Performance as a function of $\tau$. Each point represents the mean of all 5 testing folds.