Kolmogorov Regularization for Link Prediction

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Abstract

Link prediction in graphs is an important task in the fields of network science and machine learning. We propose a flexible means of regularization for link prediction based on an approximation of the Kolmogorov complexity of graphs. Informally, the Kolmogorov complexity of an object is the length of the shortest computer program that produces the object. Complex networks are often generated, in part, by simple mechanisms; for example, many citation networks and social networks are approximately scale-free and can be explained by preferential attachment. A preference for predicting graphs with simple generating mechanisms motivates our choice of Kolmogorov complexity as a regularization term. Our method is differentiable, fast and compatible with recent advances in link prediction algorithms based on graph neural networks. We demonstrate the effectiveness of our regularization technique on a set of diverse real-world networks.

1 Introduction

Network models have become an indispensable tool to study complex systems of discrete objects and their interactions. Network science has been applied with great success to a variety of scientific disciplines, often resulting in rich data sets that can be studied under the auspices of machine learning. A significant area of research that has emerged from the study of networks is link prediction between nodes based on incomplete instances of data [1]. The importance of link prediction techniques is underscored by its breadth of applications to topics such as protein function anticipation [2], friendship identification for social network users [3] and scientific collaboration inference [4].

Many complex networks have simple causal mechanisms that underlie their generation. For example, it has been theorized that scale-free networks such as the World Wide Web are often generated by a system of preferential attachment whereby new nodes are more likely to attach to nodes that are already well connected [5]. The small-world property, which states that any two nodes can reach each other via a short path, is present in many real-world phenomenon such as social networks and can be artificially generated by the simple Watts-Strogatz model [6]. The observation that complex networks can be characterized by such simple mechanisms raises the possibility of incorporating complexity biases for network modelling.

In recent years, graph neural networks (GNNs) have proven to be highly adept at solving link prediction problems [7,8,9]. By leveraging relational inductive biases to obtain high-level node representations, architectures such as graph auto-encoders can decode meaningful, unseen links from their latent representations [7,9]. Another benefit of GNNs is that they can be trained in the canonical way by defining a loss function and employing gradient-based learning methods. This flexible training framework affords an opportunity to incorporate information about the generating mechanisms of networks through a regularization term.

In this work, we introduce a differentiable regularization term for link prediction that penalizes graphs with large estimated Kolmogorov complexity to encourage the creation of graphs with sim-
The dotted and dashed lines represent two competing sets of inferred edges on a graph. The graph on the right has a much lower estimated Kolmogorov complexity than the graph on the left. Correspondingly, the graph on the right can be explained by a simple preferential attachment generating mechanism (where the red nodes are hubs).

The Kolmogorov complexity of an object is the length of the shortest computer program that outputs that object. Kolmogorov complexity and the related field of algorithmic probability rigorously define a notion of simplicity, and have been used to discover simple neural networks with a marked ability to generalize [10]. By proposing Kolmogorov complexity as a regularizer on graph outputs, we aim to augment the ability of models such as graph neural networks to generalize when predicting links. Although Kolmogorov complexity is uncomputable, we use recent approximation methods to develop a straightforward, fast and differentiable means of regularization that can be used with any model that learns by minimizing a loss function. In experiments on real-world networks our method proves to be highly effective.

Related Work In [11], Kolmogorov complexity was proposed by Hernández-Orozco et al. as a regularization term with a weighting parameter in conjunction with a general loss function. Similar to Schmidhuber’s approach in [10], Hernández-Orozco et al.’s form of regularization pushes a model’s parameters towards a low complexity, thus increasing the algorithmic probability of the model. In our methodology, we use Kolmogorov complexity to regulate the output, rather than the parameters, of a model. By taking this approach we are rewarding the model for learning to generate objects from simpler rule sets. Hernández-Orozco et al. also point out that a Kolmogorov penalty function, as they have defined it, cannot be optimized by gradient-based methods [11]. We overcome this problem by using a probabilistic interpretation of the regularization term.

In order to approximate the gradient of our Kolmogorov complexity based regularization term, we use a method based on perturbing the predicted adjacency matrix. The study of the algorithmic causality of an object by means of a perturbation calculus on the object’s estimated Kolmogorov complexity was pioneered by Zenil et al. in [12]. Their approach was recently used to define a highly effective unsupervised algorithm for identifying generating mechanisms in graphs [13].

Paper Outline This paper is structured as follows. Section 2 briefly presents several fundamental concepts in algorithmic information theory and discusses a method for approximating Kolmogorov complexity that we later use in our methodology. Section 3 describes our Kolmogorov complexity regularization term for link prediction that is fully compatible with standard differentiable approaches to training neural networks, such as backpropagation. Section 4 presents experiments on five diverse real-world networks, demonstrating the effectiveness of our method on powerful graph neural network approaches to link prediction. Section 5 concludes the paper and considers challenges and future work.

2 Background Information

Algorithmic information theory (AIT) primarily studies the irreducible information content of objects and was charmingly summarized by one of its founders, Gregory Chaitin, as “the result of putting Shannon’s information theory and Turing’s computability theory into a cocktail shaker and
shaking vigorously [14]. The information content or complexity of an object is measured by the size, in bits, of the shortest computer program that can compute the object. For example, consider the following two binary strings:

10101010101010101010101010101010
000110110110011001010001000110110

The first string has a simple pattern which can be described concisely as:

```
print '10' 16 times
```

The second string does not have a clear pattern and likely has no simpler description than merely printing the string itself. In more formal terms, the Kolmogorov complexity of a string $s$ is:

$$
K(s) = \min \{|p| : U(p) = s \}
$$

where $p$ is a program of length $|p|$ bits that, when run on a universal Turing machine $U$, outputs $s$. Kolmogorov complexity is invariant to the choice of $U$ up to an additive constant independent of the choice of $s$.

Closely related to Kolmogorov complexity is the concept of algorithmic probability, a method for assigning a universal prior probability to objects. Consider a program $p$ that produces a binary string $s$ when run on a universal prefix-free Turing machine $U$. The universal prior probability for each string $s$ is defined as:

$$
m(s) = \sum_{p: U(p) = s} 2^{-|p|}
$$

As $T$ is a universal prefix-free Turing machine, the group of valid programs on $U$ are a prefix-free set and thus, by Kraft’s inequality, the sum is bounded by one. Kolmogorov complexity and algorithmic probability are beautifully linked through Levin’s Coding Theorem which gives the following result:

$$
-\log_2 m(s) = K(s) + O(1)
$$

Algorithmic probability is, in part, guided by Epicurus’ principle of multiple explanations (if several theories are consistent with the data, retain them all) and Occam’s razor (among theories consistent with the data, choose the simplest) [15]. The prior probability $m(s)$ satisfies these principles by assigning a non-zero probability to every string and giving a higher probability to strings with shorter generating programs. For more information on the field of algorithmic information theory in general, we refer the reader to the following references [15, 16, 17, 18, 19, 20, 21].

2.1 Approximating Kolmogorov Complexity

Both Kolmogorov complexity and algorithmic probability are uncomputable for reasons related to the halting problem, therefore approximations are required. Statistical lossless compression algorithms are a popular approach to estimate Kolmogorov complexity [22]. Lossless compression techniques like the Lempel-Ziv-Welch (LZW) algorithm clearly provide intuitive estimates of the Kolmogorov complexity of an object, but suffer from an inability to capture meaning beyond classical Shannon information theory [22, 23].

**Coding Theorem Method**  The Coding Theorem Method (CTM) provides a straightforward approximation to the Kolmogorov complexity of an object that captures algorithmic features rather than merely statistical features [24]. The CTM directly approximates the algorithmic probability of small strings by exploring the large space of Turing machines with a fixed number of symbols and states. Let $(n, 2)$ be the class of all $n$-state 2-symbol Turing machines $T$ using the Turing machine formalism outlined in the busy beaver game [25]. The CTM defines the following function for a binary string $s$:

$$
D_{(n,2)}(s) = \frac{|\{T \in (n, 2) : T \text{ produces } s\}|}{|\{T \in (n, 2) : T \text{ halts}\}|}
$$

Of course, in general it is impossible to know if a machine will halt; however, for the 2 symbol case the largest number of steps taken before halting are known up to $n = 4$ (and theorized for $n = 5$) [24]. Therefore, $D_{(n,2)}(s)$ can be computed for small $n$ using brute-force. Using an approximate form of Levin’s Coding Theorem, the CTM estimate of Kolmogorov complexity, denoted by $CTM_{(n,2)}(s)$, is given as:

$$
CTM_{(n,2)}(s) = -\log_2 D_{(n,2)}(s)
$$
Block Decomposition Method  Unfortunately, it is ultimately uncomputable to use the CTM approximation on large objects due to the rapid growth of the busy beaver function. To address this limitation, the Block Decomposition Method (BDM) \cite{23} extends the CTM via an aggregation rule designed to reconstruct the Kolmogorov complexity of a large object from its smaller components. The BDM has been applied with great success to problems in machine learning and causality\cite{11,13} and is described as follows. For a given binary string \( s \), decompose the string into the multiset \( S_s = \{s_1, s_2, \ldots, s_r\} \) where \( s_i \) are consecutive slices from \( s \) of size \( r \). The value of \( r \) is chosen to be small enough so that the CTM can compute an approximation to the Kolmogorov complexity of the slice (we assume the length of \( s \) is divisible by \( r \)). Let \( U_s \) be the set of unique values in \( S_s \) and let \( c_u \) be the number of times slice \( u \in U_s \) appears in \( S_s \). The BDM is based on the following intuition: if the CTM approximates the Kolmogorov complexity for each slice \( u \), then a program with an estimated complexity of \( \sum_{u \in U_s} \text{CTM}(n, 2)(u) \) can be used to generate all of the unique building blocks of \( s \). The number of times each slice \( u \) appears in \( s \) can be specified in \( \log_2(c_u) \) bits, thus the BDM approximation for the Kolmogorov complexity of \( s \) is:

\[
BDM(n, 2)(s) = \sum_{u \in U_s} \text{CTM}(n, 2)(u) + \log_2(c_u)
\]

Approximating Graph Complexity  The BDM can be extended to approximate the Kolmogorov complexity of a graph by applying a two-dimensional variant of the BDM to the graph’s binary adjacency matrix \( A \). \cite{23,26}. The CTM component of the BDM approximation is computed using Turing machines that run on a 2-dimensional tape and produce arrays rather than strings. The BDM approximation of the graph’s Kolmogorov complexity \( BDM(n, 2)(A) \) is computed over a partition of \( A \) into block matrices small enough to have a CTM value (for a more explicit formulation please see Section 3.2). Using the adjacency matrix as the descriptor of a graph introduces a potential challenge as adjacency matrices corresponding to isomorphic graphs can have different Kolmogorov complexity. However, this discrepancy is bounded by a constant independent of the choice of graph \cite{27} and in practice the BDM works very well as a Kolmogorov complexity estimator for networks \cite{13,28,29}.

Larger values of \( n \) allow for CTM estimates of larger arrays, and in turn lead to better BDM approximations \cite{23}. Therefore, for the remainder of this paper we will replace the notation \( BDM(n, 2) \) with \( K_{BDM} \), where \( n \) is assumed to be the largest number of states for which there are CTM values available. We will also exclusively use the function \( K_{BDM} \) in reference to the two-dimensional variant of the BDM.

3 Methodology

Notation  We consider an unweighted graph \( G = (V, E) \) with \( N = |V| \) nodes. The \( N \times N \) adjacency matrix \( A \) of \( G \) has elements \( a_{ij} \in \{0, 1\} \). Given a learning algorithm \( \mathcal{M} \) that predicts links (in the form of an adjacency matrix), we denote the output of \( \mathcal{M} \) as an \( N \times N \) matrix \( \hat{A} \). The elements of \( \hat{A} \) have been mapped to the open interval \((0, 1)\) by the output activation function of \( \mathcal{M} \). We will treat \( \hat{A} \) as a matrix of Bernoulli parameters where \( \hat{a}_{ij} \) represents the independent probability that there is an edge from node \( i \) to node \( j \). The reasoning for treating \( \hat{A} \) as a matrix of probabilities will be discussed in Section 3.2. When referring to the Bernoulli random variable parameterized by \( \hat{a}_{ij} \) we will write \( \tilde{a}_{ij} \). Additionally, when referring to the matrix of independent Bernoulli random variables parameterized by the values in \( \hat{A} \), we will write \( \tilde{A} \).

3.1 Regularized Loss Function

Let \( L \) denote a general loss function used to train \( \mathcal{M} \) over \( A_{\text{Train}} \), a noisy or restricted view of \( A \). For example, a reasonable choice of \( L \) is the binary cross entropy loss function with weighting to account for a sparsity of edges.

Kolmogorov Regularization  Given a learning algorithm that predicts links in a graph, we define Kolmogorov-regularized functions as the class of loss functions with the form:

\[
\hat{L} = L + \lambda \cdot \mathbb{E}[K(\tilde{A})]
\]  

(1)
where $\lambda \in \mathbb{R}^+$ is a weighting hyperparameter and $\mathbb{E}[K(\mathbf{A})]$ is the expected Kolmogorov complexity of $\mathbf{A}$. Because the Kolmogorov complexity of an object is uncomputable, we rely on the BDM to produce an approximation $\mathbb{E}[K_{BDM}(\mathbf{A})]$ of the expected Kolmogorov complexity of $\mathbf{A}$. We denote the class of loss functions that use this approximation to Kolmogorov regularization as:

$$\mathcal{L}_{BDM} = \mathcal{L} + \lambda \cdot \mathbb{E}[K_{BDM}(\mathbf{A})]$$  \hspace{1cm} (2)

### 3.2 Practical Differentiable Formulation

Let us partition the binary adjacency matrix $\mathbf{A}$ into blocks of size $R \times R$ as follows:

$$\mathbf{A} = \begin{bmatrix}
A_{11} & A_{12} & \cdots & A_{1N'} \\
A_{21} & A_{22} & \cdots & A_{2N'} \\
\vdots & \vdots & \ddots & \vdots \\
A_{N'1} & A_{N'2} & \cdots & A_{N'R'}
\end{bmatrix}$$

where $N' = N/R$. We have made the mild assumption that $N$ is divisible by $R$; if this is not the case we pad $\mathbf{A}$ with zeros (see Section 4.1). We refer to the multiset of all blocks in $\mathbf{A}$ as $\mathcal{A}_A = \{A_{11}, A_{21}, A_{12}, \ldots, A_{N'R'}\}$ and the set of unique elements in $\mathcal{A}_A$ as $\mathcal{U}_A$. Recall that the BDM approximation of the Kolmogorov complexity of a binary adjacency matrix is:

$$K_{BDM}(\mathbf{A}) = \sum_{\mathbf{U} \in \mathcal{U}_A} CTM(\mathbf{U}) + \log_2(c_\mathbf{U})$$

where $c_\mathbf{U}$ is the number of times a block $\mathbf{U} \in \mathcal{U}_A$ appears in $\mathcal{A}_A$. Because we would like our regularization term to be used with gradient-based training algorithms such as backpropagation, we must alter the BDM approximation to be differentiable. This requirement motivates our designation of the model output $\mathbf{A}$ as an adjacency matrix of edge probabilities. By treating $\mathbf{A}$ as a collection of $N^2$ independent Bernoulli random variables parameterized by $\mathbf{A}$, we have a regularization term $\mathbb{E}[K_{BDM}(\mathbf{A})]$ that is clearly differentiable with respect to the elements of $\mathbf{A}$. However, this decision also introduces a computational complexity problem. To appreciate this problem, note that each of the $N^2$ blocks in $\mathbf{A}$ has a unique probability mass function over the $2^{R^2}$ possible binary matrices of size $R \times R$. Therefore, there are $(2^{R^2})^{N^2} = 2^{N^2}$ unique probabilities to be computed in order to directly determine $\mathbb{E}[K_{BDM}(\mathbf{A})]$ (this is also apparent from the fact that there are $2^{N^2}$ possible realizations of $\mathbf{A}$).

**Monte Carlo Perturbation** To mitigate the computational complexity problem, we adopt a Monte Carlo approach where we sample $m$ times from $\mathbf{A}$. However, instead of approximating $\mathbb{E}[K_{BDM}(\mathbf{A})]$ we use the samples to directly approximate the gradient $\nabla \mathbb{E}[K_{BDM}(\mathbf{A})]$. Consider the partial derivative of $\mathbb{E}[K_{BDM}(\mathbf{A})]$ with respect to $\tilde{a}_{ij}:

$$\frac{\partial \mathbb{E}[K_{BDM}(\mathbf{A})]}{\partial \tilde{a}_{ij}} = \frac{\partial}{\partial \tilde{a}_{ij}} \cdot \mathbb{P}(\tilde{a}_{ij} = 1) \cdot \mathbb{E}[K_{BDM}(\mathbf{A})|\tilde{a}_{ij} = 1]$$

$$+ \frac{\partial}{\partial \tilde{a}_{ij}} \cdot \mathbb{P}(\tilde{a}_{ij} = 0) \cdot \mathbb{E}[K_{BDM}(\mathbf{A})|\tilde{a}_{ij} = 0]$$

$$= \frac{\partial}{\partial \tilde{a}_{ij}} \cdot \mathbb{E}[K_{BDM}(\mathbf{A})|\tilde{a}_{ij} = 1]$$

$$+ \frac{\partial}{\partial \tilde{a}_{ij}} \cdot (1 - \tilde{a}_{ij}) \cdot \mathbb{E}[K_{BDM}(\mathbf{A})|\tilde{a}_{ij} = 0]$$

$$= \mathbb{E}[K_{BDM}(\mathbf{A})|\tilde{a}_{ij} = 1] - \mathbb{E}[K_{BDM}(\mathbf{A})|\tilde{a}_{ij} = 0]$$

Let $\mathbf{A}^{(1)}, \mathbf{A}^{(2)}, \ldots, \mathbf{A}^{(m)}$ denote $m$ binary matrices sampled from $\mathbf{A}$. For each sample $\mathbf{A}^{(k)}$ we can partition the matrix into $R \times R$ blocks and compute a frequency table of all the different blocks in $\mathcal{O}(N^2)$ time. We will use the notations $\tilde{A}_{ij=1}^{(k)}$ and $\tilde{A}_{ij=0}^{(k)}$ to denote a sample $\tilde{A}_{ij}^{(k)}$ that has
We perform our experimentation on five different real-world networks: a network of links between VGAE models require a node feature matrix, we use an appropriately sized identity matrix as a with weighting proportional to the ratio of negative to positive labels. The VGAE network uses the we test the regularization term on a graph auto-encoder (GAE) and a variational graph auto-encoder (VGAE) [9]. For both models, we follow the designs used in [9] where the encoders are two-layer ⊙ where ∇ the average value of and then simply incrementing and decrementing BDM values based on the existing frequencies. The it will be treated as a constant. Using both the lookup table and the frequency table, the difference R binary string of the values in the R × R block matrix containing aij(k) as the index key for both tables, and then simply incrementing and decrementing BDM values based on the existing frequencies. The average value of K_BDM(˜Aij=1) − K_BDM(˜Aij=0) approaches ∂E[K_BDM(˜A)]/∂hij as more samples are taken. Therefore, if we compute the value of K_BDM(˜Aij=1) − K_BDM(˜Aij=0) for each element aij(k) we have effectively sampled from the gradient ∇E[K_BDM(˜A)] in O(N^2) time.

**Loss Function Incorporation** Let ∇E[K_BDM(˜A)](k) denote a sample from ∇E[K_BDM(˜A)] and for simplicity of notation we write the sample mean of the gradient as:

\[ \mathbf{G}_{A,m} = \frac{1}{mn} \sum_{k=1}^{m} \nabla E[K_BDM(\tilde{A})]^{(k)} \]

In order to incorporate the sample mean of the gradient into the gradient of the loss function we simply multiply each element in ˜A with its corresponding element in ˜G_{A,m} and the weighting parameter λ, then sum these products back into the loss function. Note that despite the notation, the elements of ˜G_{A,m} are treated as constants. Our loss function is summarized below as:

\[ \hat{\mathcal{L}}_{BDM} = \mathcal{L} + \lambda \cdot \mathbf{1}^T (\tilde{A} \odot \tilde{G}_{A,m}) \mathbf{1} \quad (3) \]

where \( \mathcal{L} \) denotes element-wise multiplication and \( \mathbf{1} \) is the column vector of \( N \) ones.

# 4 Experiments

In order to assess the performance of our method, we measure the impact of the regularization term \( \lambda \cdot \mathbf{1}^T (\tilde{A} \odot \tilde{G}_{A,m}) \mathbf{1} \) on the ability of standard GNN frameworks to predict links. More specifically, we test the regularization term on a graph auto-encoder (GAE) and a variational graph auto-encoder (VGAE) [9]. For both models, we follow the designs used in [9] where the encoders are two-layer graph convolutional network (GCN) [30] with 32 and 16 hidden units, respectively. The decoders produce the edge probabilities by computing the sigmoid of the inner product of the latent node embeddings. As for the loss functions, the GAE network is trained on the binary cross-entropy loss with weighting proportional to the ratio of negative to positive labels. The VGAE network uses the same loss as a reconstruction term, but includes an additional Kullback-Leibler divergence term to measure the discrepancy between the approximation of the posterior and the latent prior, which we define as an isotropic Gaussian distribution with unit variance.

## 4.1 Link Prediction on Real-World Networks

We perform our experimentation on five different real-world networks: a network of links between Wikipedia pages on chameleons [31], a road transportation network from Chicago [32, 33, 34], the Cora citation network of scientific publications [35], a protein-protein interaction network from PDZBase [36, 37], and a network of co-purchases of US political books [38]. Table I contains a brief overview of key details for each of the five networks. These data sets were chosen to represent a broad range of applications with highly different generating mechanisms. As both the GAE and VGAE models require a node feature matrix, we use an appropriately sized identity matrix as a dummy input for each of the five networks. Additionally, each network is processed to be undirected and contain no self-loops.

**Experiment Design** Our experiment design is largely based on that of [9, 39]. We begin by dividing each of the five networks into training, validation, and testing data sets. The training input...
Table 1: Overview of the networks used in our experiments.

| Network     | Node Count | Edge Count | Category           |
|-------------|------------|------------|--------------------|
| Chameleon   | 2277       | 31421      | Internet           |
| Chicago     | 1467       | 1298       | Transportation     |
| Cora        | 2708       | 5429       | Citation           |
| PDZBase     | 212        | 244        | Protein Interaction|
| Political Books | 105 | 441        | Purchasing         |

Table 2: Summary of \( \lambda \) values used for each data set.

| Network     | Chameleon | Chicago | Cora | PDZBase | Political Books |
|-------------|-----------|---------|------|---------|-----------------|
| \( \lambda \) Value | \( 5 \times 10^{-7} \) | \( 1 \times 10^{-5} \) | \( 4 \times 10^{-7} \) | \( 1 \times 10^{-4} \) | \( 3 \times 10^{-5} \) |

is the original adjacency matrix with 80% of the edges randomly retained. Because the graph convolutional operator requires the adjacency matrix to be updated with self-loops along the diagonal, our training label is simply the training input summed with the identity matrix. The validation set consists of half of the 20% of original edges not selected for training along with an equal number of false edges that do not exist in the original graph. All of the true edges and false edges are randomly selected. The test set consists of the remaining original edges along with an equal number of random false edges that do not exist in either the original graph or the set of false validation edges. Note that different random splits will, of course, give slightly different results.

We randomly initialize both the GAE and the VGAE models using Glorot initialization \[40\] and perform multiple trials to account for different initializations. We employ two standard metrics for binary classification: area under the ROC curve (AUC) and average precision (AP). After splitting the data sets, we establish preliminary results for both models without any regularization over 10 trials on each of the five validation sets. All trials described in this paper are run for 1000 epochs. During each trial, we save the model weights for both the maximum validation AUC and AP scores.

Using these preliminary results we search for an appropriate value of \( \lambda \) on the validation sets in a simple manner, with care taken to make sure this process is not overly tedious. The starting point for the search is the inverse of the square of the node count as the BDM can potentially grow quadratically with the node count (until the CTM dictionary is exhausted). We then proceed to search in proportional increments until neither increasing nor decreasing \( \lambda \) leads to a significant increase in validation performance. Table \[2\] contains the values of \( \lambda \) in both models for each of the five networks. Throughout our experiments we use \( m = 1 \) sample to approximate the gradient \[ \nabla \mathbb{E}[K_{BDM}(\mathbf{A})] \]. Increasing the value of \( m \) does not seem to improve results significantly, but does slow down training as the regularization term takes \( O(mN^2) \) time to compute. \( R \) is set to 4, the largest value for which there are binary CTM array estimates available \[41\]. If \( N \) is not divisible by \( R \) we can simply pad \( \mathbf{A} \) with zeros; this has a negligible effect on the BDM as \( R = 4 \ll N \).

After the \( \lambda \) values have been determined, we train all five data sets using Kolmogorov regularization. For both models, we repeat this process for 10 trials per data set, saving the model weights that yield the maximum validation AUC and AP scores for each trial and data set. Finally, we run all the saved validation model weights (with and without regularization) on the corresponding network test sets. We report the means and standard errors on the test sets for both AUC and AP scores in Tables \[3\] and \[4\] respectively.

Discussion In Tables \[3\] and \[4\] we showed that Kolmogorov regularization is highly effective for link prediction tasks on a broad variety of data sets. In particular, the performance gains on the Chameleon, Chicago and Political Books networks were impressive when the graph neural networks were trained with Kolmogorov regularization. Note that despite already being very high, the results on the Chameleon data set were significantly superior with the Kolmogorov regularizer as the standard error ranges for this network were very small. The Cora data set displayed a marginal increase on the AUC metric for the GAE and the AP metric for the VGAE, but in comparison to other networks the improvements were not as drastic. Regularization did have a notable positive
Table 3: Link prediction results for AUC metric.

| Network      | GAE No Reg. | GAE Kol. Reg. | VGAE No Reg. | VGAE Kol. Reg. |
|--------------|-------------|---------------|--------------|----------------|
| Chameleon    | 98.22 ± 0.01| 98.90 ± 0.02  | 98.17 ± 0.02 | 98.82 ± 0.02  |
| Chicago      | 76.38 ± 1.21| 88.00 ± 0.13  | 82.40 ± 0.79 | 88.11 ± 0.08  |
| Cora         | 81.85 ± 0.48| 82.60 ± 0.24  | 82.56 ± 0.28 | 82.94 ± 0.30  |
| PDZBase      | 71.51 ± 2.48| 83.14 ± 0.44  | 75.28 ± 1.64 | 83.89 ± 0.63  |
| Political Books | 83.70 ± 0.41| 88.94 ± 0.43  | 86.61 ± 0.43 | 88.96 ± 0.33  |

Table 4: Link prediction results for AP metric.

| Network      | GAE No Reg. | GAE Kol. Reg. | VGAE No Reg. | VGAE Kol. Reg. |
|--------------|-------------|---------------|--------------|----------------|
| Chameleon    | 98.48 ± 0.02| 98.96 ± 0.02  | 98.52 ± 0.02 | 98.88 ± 0.02  |
| Chicago      | 78.35 ± 1.03| 86.17 ± 0.31  | 82.59 ± 1.10 | 86.22 ± 0.15  |
| Cora         | 86.09 ± 0.23| 86.21 ± 0.19  | 86.26 ± 0.25 | 86.76 ± 0.21  |
| PDZBase      | 75.04 ± 1.42| 73.80 ± 2.25  | 76.02 ± 1.20 | 77.16 ± 2.52  |
| Political Books | 80.75 ± 0.45| 89.28 ± 0.40  | 85.04 ± 0.33 | 91.08 ± 0.25  |

effect on the AUC score of the PDZBase network, but the AP score did not improve. The volatility shown on the PDZBase network was likely due to its small edge count.

Training was performed on an Intel i7-4790k CPU with 8GB of RAM and an Nvidia GTX 970 GPU. Computation of the gradient sample for the Kolmogorov regularization term was done entirely on the CPU, although this process should scale well on a GPU. Each trial of 1000 epochs took from approximately 30 seconds (Political Books) to approximately 35 minutes (Cora) depending on the size of the network.

5 Conclusion

In this paper we have presented Kolmogorov complexity as an effective regularizer for link prediction and have introduced a fast, differentiable regularization term based on approximations of Kolmogorov complexity for graphs. Our method is flexible and can be used with any model that learns by minimizing a loss function through gradient-based methods. In experiments performed with recent approaches to link prediction using graph neural networks, our regularization term achieves impressive improvements for a variety of real-world networks.

An ambitious task for future research is the development of more powerful approximation methods of Kolmogorov complexity than the BDM. As the size of the network grows beyond the CTM, the BDM loses its ability to capture algorithmic complexity [23]. Naturally, better approximations would allow for applications to extremely large networks as well as an increase in the quality of regularization bias towards networks with simpler generating mechanisms.

Broader Impact

When discussing the broader impact of our work, it is important to consider existing applications of link prediction algorithms. As mentioned in the introduction, link prediction has been used by both the scientific community (protein interactions, potential collaborations, etc.) as well as by the commercial sector (recommendation systems, social networking tools, etc.) for positive means. It is easy to see that such applications may only improve in predictive power when applying the methods described in this paper. If an organization or government were to use link prediction for unethical purposes, our algorithm is general enough that they could make use of it. We do not think our form of regularization will contribute to the failure of a larger system. Additionally, the algorithm pushes towards networks with simple generating mechanisms, but it seems unlikely that this would exploit any unwanted bias in the data.
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