A Geometry-Aware Algorithm to Learn Hierarchical Embeddings in Hyperbolic Space

Abstract

Hyperbolic embeddings are a class of representation learning methods that offer competitive performances when data can be abstracted as a tree-like graph. However, in practice, learning hyperbolic embeddings of hierarchical data is difficult due to the different geometry between hyperbolic space and the Euclidean space. To address such difficulties, we first categorize three kinds of illness that harm the performance of the embeddings. Then, we develop a geometry-aware algorithm to tackle the above illnesses. Specifically, we introduce the dilation operation, the transitive closure regularization, and an improved negative sampling strategy to build our algorithm. We empirically validate these techniques and present a theoretical analysis of the mechanism behind the dilation operation. Experiments on synthetic and real-world datasets reveal superior performances of our algorithm.

1 Introduction

Learning data representation has gained popularity in modern machine learning as it provides a metric space that reveals or preserves the inherent data structure Mikolov et al. (2013); Pennington et al. (2014); Bojanowski et al. (2017); Hoff et al. (2002); Grover & Leskovec (2016); Perozzi et al. (2014); Nickel et al. (2011); Bordes et al. (2013); Riedel et al. (2013). Hyperbolic embeddings, a class of hierarchy representation methods, have shown competitive performances when data can be abstracted as a graph Chamberlain et al. (2017); Davidson et al. (2018); Ganea et al. (2018a); Gu et al. (2018); Tifrea et al. (2018).

In this paper, we focus on the following embedding task. Let $\mathcal{D}$ be a dataset incorporated with a set of hierarchical relations represented as edges in a tree-like graph $\mathcal{G}$. The goal is to learn an embedding Θ in the *hyperbolic* space by drawing positive and negative samples of edges from the graph such that Θ preserves the edge relationships, which are reflected by the order of similarity between data pairs. The formal problem statement is presented in Section A.1.

Theoretically, hyperbolic space, such as the Poincaré Ball model, benefit from high representational power due to their negative curvatures Nickel & Kiela (2017); Sala et al. (2018). This observation has motivated research on solving real-world problems in hyperbolic space. For datasets with an *observed* structure, hyperbolic space can embed the data and preserve the structure with arbitrarily low distortion (Nickel & Kiela, 2017; Chamberlain et al., 2017; Nickel & Kiela, 2018; Ganea et al., 2018b; Chami et al., 2019c). For datasets with a *latent* structure, especially those obeying the power-law, hyperbolic space can provide a natural metric such that finer concepts are embedded into areas allowing more subtlety (Tifrea et al., 2018; Leimeister & Wilson, 2018; Le et al., 2019).

Despite the theoretical advantages of hyperbolic embeddings, learning such representation in practice is difficult. Specifically, the following fundamental difficulties have not been well-studied in the literature. (1) Many properties of the Euclidean space do not transfer to hyperbolic space. For example, the latter generally do not have the scale or shift-invariance in the sense of preserving similarity orders. (2) Many nice properties exclusive to hyperbolic space may improve learning. However, it is unclear how to design algorithms to effectively incorporate these properties. (3) Optimization in hyperbolic space is (i) expensive due to a more sophisticated distance measure and (ii) unstable because gradient descent is performed on hyperbolic manifolds.

In this paper, we present a theoretical and empirical analysis of these difficulties and a set of solutions to them.

(a) The baseline Poincaré embedding algorithm (b) Our algorithm

Figure 1: Visualizations of two-dimensional embeddings of a synthetic balance tree (156 nodes, 155 edges) learned by the baseline Poincaré embedding algorithm in Nickel & Kiela (2017) and our geometry-aware algorithm, respectively. Both algorithms are trained for 3000 epochs to ensure convergence.

To begin with, in Figure 1a, we visualize the learned 2-dimensional hyperbolic embeddings of a synthetic tree according to Nickel & Kiela (2017). The lines refer to ground-truth edges and the points refer to the learned hyperbolic embeddings. The red lines indicate bad cases, i.e., the embeddings fail to reconstruct these ground-truth edges. Comparing it with the perfect reconstruction in Figure 1b, we find bad cases are of intrinsic different types. We categorize such bad cases into *capacity illness*, *inter-subtree illness*, and *intra-subtree illness*. Intuitively, the above categorization discriminates whether an incorrect edge involves nodes from different subtrees. Formal definitions and intuitive visualizations are presented in Section A.2. We also present a theoretical analysis of local capacity, capacity illness, and their relationship in Section A.3.

We then develop an algorithm that aims at reducing the above three kinds of illness in Section 3. To reduce capacity illness, we introduce the *dilation* operation during the learning process (see Section 3.1). Geometrically, it pushes points to locations with sufficient local capacity in hyperbolic space, where the sufficiency is guaranteed by equation 11. To reduce inter-subtree illness, we add transitive closure edges (i.e., edges between nodes and their non-parent ancestors) of data to positive samples in the objective equation 5. To reduce intra-subtree illness, we adopt the re-weighting strategy, which modifies the weights of transitive closure edges. We call the combination of these two strategies the *transitive closure regularization* (see Section 3.2). In addition, we find that with sufficient local capacity, the transitive closure regularization strategy works out better. Thus, we combine the above three techniques to build our algorithm. Note that these techniques are generally not admissible in the Euclidean space[1].

Finally, to prevent gradient vanishing at late stages, we provide an improved negative sampling strategy with a theoretical validation in Section A.4. For a node A, the strategy does not accept any node A' to be a negative sample of A if the distance between A and A' is larger than a certain threshold.

We conduct experiments on synthesize and real world datasets in Section A.5. Our algorithm significant reduces all types of illness and achieves better performances under various evaluation metrics.

2 RELATED WORK

Learning in hyperbolic space is initially proposed by Nickel & Kiela (2017). It is the most related work to our paper. Their method outperforms the Euclidean counterpart in low dimensions as to the task of learning embeddings for edge reconstruction. However, since their algorithm is directly adapted from the Euclidean space, it does not naturally leverage potentially useful geometrical

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[1]For example, the Euclidean space is shift/scale-invariant, which means translation and dilation will not change the relative relations of points. That makes dilation operation ineffective.

Algorithm 1 Geometry-Aware Algorithm

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for $i = 1$ **to** N_{epoch} **do**
    Compute local capacity according to equation 13
    if The local capacity is not sufficient **then**
        Perform the dilation operation according to equation 1
    end if
    if $i \leq N_{\text{tc}}$ **then**
        Loss $\leftarrow \mathcal{L}_{\text{tc}}(\Theta)$ according to equation 3
    else
        Loss $\leftarrow \mathcal{L}(\Theta)$ according to equation 5
    end if
    Apply the negative sampling strategy in equation 16
    Apply Riemannian gradient descent over Loss
end for

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properties of hyperbolic space (see Section A.1). As a consequence, there remain many bad cases even after convergence (see Figure 1a).

A series of work directly incorporate properties of hyperbolic space via optimization (Wilson & Leimeister, 2018; Bonnabel, 2013; Absil et al., 2009; Afsari et al., 2013). Specifically, Nickel & Kiela (2018) conduct training in the Lorentz space with a closed-form expression of the geodesics on the hyperbolic manifold. However, since the learning objective equation 5 is highly non-convex, obtaining more accurate gradients does not completely solve the problem.

Another group of work either implement hyperbolic versions of commonly used neural network modules Ganea et al. (2018c); Gulcehre et al. (2018); Chami et al. (2019b), or design models specifically tailored for hyperbolic space Vulić & Mrkšić (2017); Le et al. (2019); Cho et al. (2019); Leimeister & Wilson (2018); Weber et al. (2020); Chami et al. (2019a). These methods are task-specific and thus expensive to deploy in downstream applications.

Apart from the above learning approaches, Sala et al. (2018) presents a combinatorial algorithm that achieves better performance than Nickel & Kiela (2017; 2018) with even lower dimensions. The core idea is to extend the 2-dimensional results of Sarkar (2011) to arbitrary dimensions. However, this algorithm suffers from three vital weaknesses: (1) it requires complete information of the graph; (2) it is sensitive to addition/removal of data; and (3) most critically, it involves discrete operations and thus does not have gradients. Therefore, in scenarios where complete information is unavailable, the graph dynamically changes, or joint learning is needed, this approach does not suffice.

In this paper, we endorse the importance of leveraging geometrical properties in learning unsupervised hyperbolic embeddings. Based on this intention, we develop a geometry-aware algorithm that improves embedding performances, which, to the best of our knowledge, is original.

3 THE ALGORITHM

In this section, we develop a geometry-aware algorithm (Algorithm 1) targeting the three categories of illness. In detail, we propose three strategies: the dilation operation, the transitive closure regularization, and an improved negative sampling strategy.

3.1 DILATION

We introduce the dilation operation. Formally, under the same notations as in Section A.3, we define a mapping $g : \mathcal{B} \to \mathcal{B}$ as a k-dilation if for any $A \in \mathcal{B}$:

$$d(O, g(A)) = k \cdot d(O, A). \tag{1}$$

Notably, g can be computed explicitly. For instance, a 2-dilation can be formulated as

$$g(A) = \frac{2}{1 + \|A\|^2} A. \tag{2}$$

Figure 2: Visualization of the learning process of two-dimensional Poincaré embeddings with (1) the baseline algorithm in the upper row, and (2) the geometry-aware algorithm in the lower row. Dataset and plotting settings are identical as in Figure 1, except for that only intra-subtree illness and inter-subtree illness that exist consistently throughout the entire 3000 epochs are plotted.

The dilation operation rescales the embedded structure so that each point is pushed to a location with sufficient local capacity. Given $A \in \mathcal{B}$ with degree k, let r_A be the distance between A and its k-nearest neighbor in $\mathcal{V}$ (in Poincaré distance). The operation helps increase r_A and thus increasing the local capacity of (A, r_A).

3.2   Transitive Closure Regularization

In this section, we present the transitive closure regularization strategy. It contains the following two operations.

Adding transitive closure edges.   The transitive closure edges $\mathcal{E}_{tc}$ are edges between nodes and their non-parent ancestors. These edges are also considered as positive samples in addition to $\mathcal{E}$ in the objective equation 5. The purpose of adding these auxiliary edges is to push the subtrees apart so they are less likely to overlap in the Poincaré ball.

Re-weighting.   We modify the weights of transitive closure edges to prevent overfitting in early epochs. Let η_{tc} be a real number between 0 and 1. Then, the objective becomes

$$\mathcal{L}_{tc}(\Theta) = \mathcal{L}(\Theta) + \eta_{tc} \sum_{(i,j) \in \mathcal{E}_{tc}} \mathcal{L}_{i,j}(\Theta). \tag{3}$$

It is noteworthy that these operations are not admissible in general spaces like the Euclidean space. Since the local capacity of any $(A, r) \in \mathbb{R}^d \times \mathbb{R}$ is a constant with respect to d, we cannot alleviate the tremendous capacity illnesses generated by transitive closure edges.

4   Experiments

Figure 2 shows how our algorithm outperforms the baseline (Nickel & Kiela, 2017) on the synthetic dataset. Our method not only achieves perfect MAP (**0.998**) but also yields better reconstructed geometry. We do extensive experiments on multiple real-world datasets of various scales and characteristics. Results show our algorithm consistently outperform the baseline algorithms (Nickel & Kiela, 2017; 2018), especially on extremely bushy datasets. See Section A.5 for detailed statistics.

5   Conclusion

In this paper, we analyze three categories of illness and develop a geometry-aware algorithm that targets at reducing these illnesses and improving performance. Our algorithm shows superior performance over baseline models on both synthetic and real world datasets.

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A APPENDIX

A.1 PRELIMINARIES

A hyperbolic space H^d is a d-dimensional Riemannian manifold with a constant negative sectional curvature. Models of hyperbolic space include the hyperboloid model and the Poincaré ball model, etc. In this paper, we focus on the Poincaré ball model. However, our methods can also be adapted to other models via isometries.

Let $\mathcal{B} = \mathcal{B}^d$ denote the d-dimensional Poincare ball. The distance between any two points $B_1, B_2 \in \mathcal{B}^d$ is given by

$$d(B_1, B_2) = \text{arcosh}\left(1 + 2\frac{\|u - v\|^2}{(1 - \|u\|^2)(1 - \|v\|^2)}\right), \tag{4}$$

where u and v are the Euclidean vectors of B_1 and B_2. In the rest of the paper, we denote the Poincaré distance by $d(\cdot, \cdot)$ and the Euclidean distance by $\|\cdot\|$. Given a set of points $\mathcal{V} = \{x_i\}_{i=1}^n$ and the relation set $\mathcal{E} \subset [n]^2$, the goal is to learn an embedding $f : \mathcal{V} \to \mathcal{B}$ that preserves the inherent structure. To achieve this goal, the following loss function $\mathcal{L}$ is minimized.

For $(i, j) \in \mathcal{E}$, define $\mathcal{N}(x_i, x_j)$ as the set of negative samples of (i, j). Let $\Theta = \{\theta_i\}_{i=1}^n$, where each $\theta_i \in \mathcal{B}^d$ is the embedding of x_i. Define $d(x_i, x_j) = d(\theta_i, \theta_j)$. The loss function is then defined as

$$\mathcal{L}(\Theta) = -\sum_{(i,j)\in\mathcal{E}} \log \frac{e^{-d(x_i, x_j)}}{\sum_{x'\in\mathcal{N}(x_i, x_j)\cup\{x_j\}} e^{-d(x_i, x')}}. \tag{5}$$

For conciseness, we denote each term in the sum in equation 5 as $\mathcal{L}_{i,j}(\Theta)$. This objective can be optimized via Riemannian gradient descent Nickel & Kiela (2017).

A.2 ILLNESS

In this section, we formally define illness that harms the performance of hyperbolic embeddings and is hard to optimize. Let $\overrightarrow{AB}$ be a ground-truth edge in G, and $\overrightarrow{AB'}$ be the inferred edge from the hyperbolic embeddings, where $B' \neq B$. We call this situation *illness with respect to* A. Let C be the nearest common ancestor of B and B'. We categorize three kinds of illness according to the pairwise relationships among B, B', and C. Formally, we define capacity illness, intra-subtree illness, and inter-subtree illness in **Definition** 1.

Definition 1 (Categories of Illness). *We define the illness to be capacity illness if B is the parent of B'. We define the illness to be an intra-subtree illness if B is the ancestor but not the parent of B'. We define the illness to be an inter-subtree illness if $C \neq B$.*

It is straightforward to see that the union of capacity illness and intra-subtree illness are exactly situations where $C = B$. Therefore, the above three kinds of illness are a partition of all illness.

We visualize these three kinds of illness in Figure 3. Intuitively, capacity illness is confusion between sibling nodes, intra-subtree illness is confusion between nodes from the same subtree, and inter-subtree illness is confusion between nodes from different subtrees. Notably, the negative impact of having different kinds of illness may vary according to specific tasks[2].

A.3 LOCAL CAPACITY

In this section, we define local capacity and theoretically relate it to capacity illness. We begin with **Definition** 2.

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[2]For example, when embedding the *is-a* relationship in WordNet, the harm caused by intra-subtree illness may not be as serious as the other two kinds. This is because, in any subtree, its nodes automatically have the *is-a* relationship with the root node of the subtree, even when they are not connected by a ground-truth edge.

Figure 3: Illustration of the three categories of illness. A is the source node and B is the ground-truth target node. It is called (1) *capacity illness* if A connects to B_1, (2) *intra-subtree illness* if A connects to B_2, and (3) *inter-subtree illness* if A connects to B_3.

Definition 2 (Local Capacity). *Given a geodesic space $(\mathcal{X}, d)$ and a geodesic sphere[3] $\mathcal{S}_r$ centered at $A \in \mathcal{X}$ with radius r. The local capacity of (A, r) is defined as*

$$\max\left\{|\mathcal{C}| : \mathcal{C} \in \mathcal{S}_r; \forall C_1, C_2 \in \mathcal{C}, C_1 \neq C_2, \atop d(C_1, C_2) > d(C_1, A) \vee d(C_2, A)\right\}. \tag{6}$$

According to **Definition** 2, for $A \in \mathcal{V}$ and a radius r, if $|\{C : C \text{ is a child of } A, d(A, C) \leq r\}|$ exceeds the local capacity of (A, r), then capacity illness must exist. To obtain bounds on local capacity, we first bound the r-packing number of a geodesic sphere centered at point $A \in \mathcal{B}$ with radius r.

First, the packing number of the geodesic sphere centered at the origin O is the same as that of A. To see why we use a result from Paupert (2016):

$$\mathbf{Isom}(\mathcal{B}) = \mathbf{M\ddot{o}b}(\mathcal{B}). \tag{7}$$

That is, the group of isometries from $\mathcal{B}$ to itself coincide with the group of all Möbius transformations preserving $\mathcal{B}$. In the Poincaré ball model, $\mathbf{M\ddot{o}b}(\mathcal{B})$ is generated by inversions in generalized spheres $\mathcal{S}'$ such that $\mathcal{S}' \perp \partial\mathcal{B}$, therefore once we extend $\overrightarrow{OA}$ to C with $\|OC\|^2 - 1 = \|OC\| \cdot \|CA\|$, then by taking the restriction in $\mathcal{B}$ of the inversion in generalized sphere $\mathcal{S}'$ centered at C we get an isometry from $\mathcal{B}$ to itself which maps A to O. Specifically, it is an isometry between any geodesic sphere centered at A and the geodesic sphere centered at the origin O (with the same radius r).

Then, we compute the r-packing number of the geodesic sphere S_r centered at O with Poincaré radius r. For $B_1, B_2 \in S_r$, let $u = \overrightarrow{OB_1}, v = \overrightarrow{OB_2}$ and r be the Poincaré norm of u. Then, as long as the angle θ between u and v satisfies

$$\theta \geq \theta_r = 2\arcsin\left(\frac{1}{2\cosh(r/2)}\right), \tag{8}$$

we have $d(B_1, B_2) \geq d(B_i, O)$, $i = 1, 2$. For not very small r, $\theta_r \approx 2e^{-r/2}$. Then, the r-packing problem is equivalent to evaluating the size of the largest spherical code of angle θ_r in dimension d, defined as $\mathcal{A}(d, \theta_r)$. According to Jenssen et al. (2018), we have

$$\mathcal{A}(d, \theta) \geq (1 + o(1))\frac{c_\theta \cdot d}{s_d(\theta)}, \tag{9}$$

where $c(\theta) = \log \frac{\sin^2(\theta)}{\sqrt{(1-\cos\theta)^2(1+2\cos\theta)}} \approx \log\frac{2}{\sqrt{3}}$ for small θ, $s_d(\theta) = (1+o(1))\frac{\sin^{d-1}\theta}{\sqrt{2\pi d}\cdot\cos\theta}$. Specifically, when $d \leq 16$ with small θ, Shannon (1959) provides a better bound:

$$\mathcal{A}(d, \theta) \geq \frac{1}{s_d(\theta)} = (1+o(1))\frac{\sqrt{2\pi d}\cdot\cos\theta}{\sin^{d-1}\theta}. \tag{10}$$

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[3]By considering the extension of radius, one can define local capacity on geodesic balls instead of spheres, which leads to the same conclusion.

To sum up, for not very small r we have the following lower bounds under different dimensions:

$$\mathcal{A}(d, \theta_r) \gtrsim \begin{cases} \pi e^{\frac{r}{2}} & d = 2 \\ \sqrt{2\pi d} \cdot 2^{1-d} e^{\frac{d-1}{2}r} & 3 \le d \le 16 \\ \sqrt{2\pi} \log \frac{2}{\sqrt{3}} \cdot d^{\frac{3}{2}} \cdot 2^{1-d} e^{\frac{d-1}{2}r} & d \ge 17 \end{cases} . \tag{11}$$

As for the upper bound, according to Kabatiansky & Levenshtein (1978),

$$\mathcal{A}(d, \theta) \le e^{\phi(\theta)d(1+o(1))}, \tag{12}$$

where $\phi(\theta) > -\log \sin \theta$ is a certain function. Therefore,

$$\mathcal{A}(d, \theta_r) \lesssim \begin{cases} \pi e^{\frac{r}{2}} & d = 2 \\ 2^d e^{\frac{dr}{2}} & d \ge 3 \end{cases} . \tag{13}$$

A.4 Improved Negative Sampling

We theoretically observe that gradient vanishing can happen during the late stages of the training process. We then show that reducing distances between negative pairs helps prevent gradient vanishing. Based on this observation, we present an improved negative sampling strategy.

First, we make the theoretical analysis. For a fixed positive pair $(i, j) \in \mathcal{E}$, we analyze the norm of the Riemannian gradient of $\mathcal{L}_{i,j}$ with respect to θ_k for every k. For conciseness, let $|\mathcal{N}(x_i, x_j)| = m$, and denote $\sum_{x' \in \mathcal{N}(x_i, x_j)}$ as $\sum_{x'}$. The loss with respect to positive pair (i, j) is given by $\mathcal{L}_{i,j}(\Theta)$ from equation 5. Suppose the naive sampling strategy provides negative pairs $\{(x_i, x')\}_{x' \in \mathcal{N}(x_i, x_j)}$. For simplicity's sake, since we have restricted our negative samples in a local ball instead of the whole Poincaré ball, we assume for any $x' \in \mathcal{N}(x_i, x_j)$, the corresponding new negative sample $\widetilde{x}' \in \widetilde{\mathcal{N}}(x_i, x_j)$ satisfies $d(x_i, x_j) \le d(x_i, \tilde{x}') = \tau d(x_i, x')$ with some constant $\tau = \tau(\mathcal{N}(x_i, x_j)) \le 1$. We then consider the following modified loss:

$$\tilde{\mathcal{L}}_{i,j}(\Theta) = -\log \left(1 + \sum_{\tilde{x}'} e^{d(x_i, x_j) - d(x_i, \tilde{x}')} \right) . \tag{14}$$

With some arithmetic, we have

$$\begin{aligned} \frac{\partial \tilde{L}_{i,j}(\Theta)}{\partial \theta_i} &= \frac{\sum_{\tilde{x}'} e^{-\tau d(x_i, x')} \left(\frac{\partial d(x_i, \tilde{x}')}{\partial \theta_i} - \frac{\partial d(x_i, x_j)}{\partial \theta_i} \right)}{e^{-d(x_i, x_j)} + \sum_{x'} e^{-\tau d(x_i, x')}}, \\ \frac{\partial \tilde{L}_{i,j}(\Theta)}{\partial \theta_j} &= -\frac{\sum_{\tilde{x}'} e^{-\tau d(x_i, \tilde{x}')} \frac{\partial d(x_i, x_j)}{\partial \theta_j}}{e^{-d(x_i, x_j)} + \sum_{\tilde{x}'} e^{-\tau d(x_i, x')}}, \\ \frac{\partial \tilde{L}_{i,j}(\Theta)}{\partial \theta_k} &= \frac{e^{-\tau d(x_i, x_k)} \frac{\partial d(x_i, \tilde{x}_k)}{\partial \theta_k}}{e^{-d(x_i, x_j)} + \sum_{\tilde{x}'} e^{-\tau d(x_i, x')}}, \end{aligned} \tag{15}$$

where $k \ne i, j$ and $\tilde{x}_k \in \tilde{\mathcal{N}}(x_i, x_j)$. When $\tau = 1$ and $\tilde{\mathcal{N}} = \mathcal{N}$, the gradients would vanish if $d_m = \min_{x'} d(x_i, x')$ is large (which usually occurs when $\|x_i\| \approx 1$). With the new negative sampling strategy whose τ is small enough, the norm of the gradients could be much larger.

Based on the above analysis, we define our negative sampling strategy $\mathcal{N}_\gamma(x_i, x_j)$ as follows: it randomly selects m samples from the set

$$\{x_k : (i, k) \notin \mathcal{E} \cup \mathcal{E}_{\text{tc}}, d(x_i, x_k) < \gamma\}. \tag{16}$$

A.5 Experiments

We apply our algorithm to both a synthetic dataset and real-world datasets on the graph reconstruction task. We evaluate the performance by mean average precision (MAP) and Mean Rank (MR) defined below. For $A \in \mathcal{V}$ with degree $\deg(A)$ and neighborhood $\mathcal{N}_A = \{B_1, ... B_{\deg(A)}\}$, let R_{A,B_i} be the smallest subset of $\mathcal{V}$ containing B_i and all points closer to A than B_i. Then, the MAP is defined as

$$\text{MAP}(f) = \frac{1}{|\mathcal{V}|} \sum_{A \in \mathcal{V}} \frac{1}{\deg(A)} \sum_{i=1}^{|\mathcal{N}_A|} \frac{|\mathcal{N}_A \cap R_{A,B_i}|}{|R_{A,B_i}|}. \tag{17}$$

(a) Capacity illness   (b) Intra-subtree illness   (c) Inter-subtree illness

Figure 4: Number of different kinds of illness under different η_{tc}.

and the MR is defined as

$$\mathrm{MR}(f) = \frac{1}{|\mathcal{V}|} \sum_{A \in \mathcal{V}} \sum_{i=1}^{|\mathcal{N}_A|} \left(|R_{A,B_i}| - i \right). \tag{18}$$

In addition, we report the number of three kinds of illness defined in **Definition** 1 after the algorithm's convergence.

A.5.1 SYNTHETIC DATASET

We synthesize a balanced tree T of 5 layers including the root node. Each non-leaf node in T has 5 children. The edges are directed, pointing from children to parents.

We compare our algorithm (Algorithm 1) to the baseline Poincaré embedding algorithm (Nickel & Kiela, 2017). In both algorithms, we set the dimension to be 2, learning rate to be 0.5, batch size to be 50, and the number of negative samples m to be 50.

The comparison between learning procedures is presented in Figure 2. As demonstrated, our algorithm learns visually more balanced embeddings with less illness. Quantitative results of the comparison, including the number of illness, MAP and MR, are presented in Figure 5 and Figure 6. Specifically, we compare different N_{tc} in Figure 5 and different η_{tc} in Figure 6. Our algorithm produces significantly less illness than the baseline algorithm and achieves the highest MAP and MR.

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A.5.2 REAL-WORLD DATASET STATISTICS

We display statistics of several datasets (including the Yelp Challenge Dataset [4], the Commodity Catalog Dataset, and the WordNet dataset Miller (1998)) used in our experiments in Table 1. We report the number of nodes $|\mathcal{V}|$, the number of edges $|\mathcal{E}|$, maximum degree, variance of all degrees, and tree depths. We make remarks to these datasets on the size and characteristics.

A.5.3 REAL-WORLD DATASET EXPERIMENT RESULTS

We run baseline algorithms (Nickel & Kiela, 2017; 2018) and our algorithm on Yelp Challenge (Tree) (see Table 2), WordNet Verbs (Tree) (see Table 3), WordNet Nouns (Tree) (see Table 4), Commodity Catalog (Tree) (see Table 5) and WordNet Nouns (Closure) (see Table 6). We report the reconstruction Mean Rank, MAP, number of capacity errors, number of intre-subtree errors and number of inter-subtree errors respectively.

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[4] https://www.yelp.com/dataset/documentation/main

(a) Capacity illness (b) Intra-subtree illness (c) Inter-subtree illness

(d) log MR (e) MAP

Figure 5: Performances of our algorithm under different hyperparameter settings on the synthetic tree. Base denotes the baseline algorithm, GA is our algorithm where DL is the dilation operation, RW is the re-weighting strategy and number is N_{tc}.

(a) Capacity illness (b) Intra-subtree illness (c) Inter-subtree illness

Figure 6: Number of different kinds of illness under different η_{tc}.

The results for Yelp Challenge (Tree) are illustrated in Table 2. In all algorithms, we set the learning rate to be 1.0, batch size to be 10, and the number of negative samples m to be 50.

The results for WordNet Verbs (Tree) are illustrated in Table 3. In all algorithms, we set the learning rate to be 1.0, batch size to be 10, and the number of negative samples m to be 50.

The results for WordNet Nouns (Tree) are illustrated in Table 4. In all algorithms, we set the learning rate to be 1.0, batch size to be 50, and the number of negative samples m to be 50.

The results for Commodity Catalog (Tree) are illustrated in Table 5. In all algorithms, we set the learning rate to be 1.0, batch size to be 10, and the number of negative samples m to be 50.

Dataset	$|\mathcal{V}|$	$|\mathcal{E}|$	Max deg	Std of deg	Depth	Size	Characteristics
Yelp Challenge (T)	1587	1586	194	7.882	4	small	shallow
Commodity Catalog (T)	134812	134811	726	26.383	6	large	shallow; bushy
WordNet Verbs (T)	13542	13541	361	9.122	12	medium	deep
WordNet Nouns (T)	82115	82114	666	6.574	15	large	deep
WordNet Nouns (C)	82115	769130	82114	913.591	15	large	deep; dense

Table 1: Dataset statistics (T: Tree; C: Closure)

		Dimensionality	MR	MAP	Capacity	Intra	Inter
YELP CHALLENGE Reconstruction	**Euclidean**		27.612	0.107	1440	82	**28**
	Nickel & Kiela (2017)		1.681	0.855	292	**45**	64
	Nickel & Kiela (2018)	2	1.520	0.894	163	47	74
	GA-DL (Ours)		1.351	**0.926**	**86**	65	47
	GA-DL-RW (Ours)		**1.202**	0.914	162	54	37
	Euclidean		11.320	0.326	1236	9	44
	Nickel & Kiela (2017)		1.063	0.988	**1**	**0**	28
	Nickel & Kiela (2018)	5	1.101	0.984	**1**	11	26
	GA-DL (Ours)		1.062	0.987	3	9	20
	GA-DL-RW (Ours)		**1.030**	**0.989**	3	9	**18**
	Euclidean		1.528	0.910	183	1	14
	Nickel & Kiela (2017)		1.042	0.990	**0**	**0**	25
	Nickel & Kiela (2018)	10	1.064	0.987	**0**	9	23
	GA-DL (Ours)		1.051	0.989	1	5	21
	GA-DL-RW (Ours)		**1.015**	**0.993**	5	6	**9**

Table 2: Yelp Challenge Dataset

		Dimensionality	MR	MAP	Capacity	Intra	Inter
WORDNET VERBS Reconstruction	**Euclidean**		66.302	0.124	7366	3778	1879
	Nickel & Kiela (2017)		8.088	0.521	4826	2582	1683
	Nickel & Kiela (2018)	2	6.875	0.448	5459	**1769**	2812
	GA-DL (Ours)		8.319	0.510	5023	2361	1908
	GA-DL-RW (Ours)		**3.559**	**0.563**	4543	2741	**1399**
	Euclidean		21.245	0.295	10383	**1273**	**277**
	Nickel & Kiela (2017)		2.329	0.854	235	2321	454
	Nickel & Kiela (2018)	5	2.389	0.853	285	2277	468
	GA-DL (Ours)		2.195	**0.878**	**211**	1443	920
	GA-DL-RW (Ours)		**1.722**	0.841	378	1306	1893
	Euclidean		6.240	0.677	4205	1424	300
	Nickel & Kiela (2017)		1.950	0.856	232	2322	457
	Nickel & Kiela (2018)	10	1.945	0.855	257	2338	**441**
	GA-DL (Ours)		**1.654**	**0.884**	**184**	1456	828
	GA-DL-RW (Ours)		**1.654**	0.842	241	**1255**	2110

Table 3: WordNet Verbs

The results for WordNet Nouns (Closure) are illustrated in Table 6. In all algorithms, we set the learning rate to be 1.0, batch size to be 50, and the number of negative samples m to be 50. [5]

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[5]Results of Nickel & Kiela (2017) are based on their official implementation and hyperparameters in `https://github.com/facebookresearch/poincare-embeddings`.

		Dimensionality	MR	MAP	Capacity	Intra	Inter
WORDNET NOUNS **Reconstruction**	Euclidean		151.321	0.235	46563	21942	2993
	Nickel & Kiela (2017)		71.271	0.322	48844	**17293**	438
	Nickel & Kiela (2018)	5	74.625	0.230	50950	18777	311
	GA-DL (Ours)		19.313	0.481	35119	21158	444
	GA-DL-RW (Ours)		**2.869**	**0.697**	**13394**	23907	**27**
	Euclidean		23.113	0.278	23978	31018	15745
	Nickel & Kiela (2017)		41.014	0.324	49152	**17276**	154
	Nickel & Kiela (2018)	10	38.395	0.241	50986	18690	100
	GA-DL (Ours)		7.900	0.754	10647	13547	5551
	GA-DL-RW (Ours)		**2.738**	**0.722**	**9483**	24844	**32**

Table 4: WordNet Nouns

		Dimensionality	MR	MAP	Capacity	Intra	Inter
COMMODITY CATALOG **Reconstruction**	Euclidean		163.044	0.024	132276	1480	700
	Nickel & Kiela (2017)		86.759	0.063	130479	**2001**	174
	Nickel & Kiela (2018)	5	50.393	0.082	127437	1968	238
	GA-DL (Ours)		5.405	**0.745**	**49837**	1971	301
	GA-DL-RW (Ours)		**2.951**	0.683	56922	7711	**26**
	Euclidean		59.133	0.052	131806	1676	289
	Nickel & Kiela (2017)		67.283	0.071	130212	**1998**	147
	Nickel & Kiela (2018)	10	36.836	0.112	124366	1964	196
	GA-DL (Ours)		2.167	**0.978**	**1871**	**1861**	390
	GA-DL-RW (Ours)		**2.015**	0.881	19902	5525	**26**

Table 5: Commodity Catalog

		Dimensionality	MR	MAP	Capacity	Intra	Inter
WORDNET NOUNS CLOSURE **Reconstruction**	Nickel & Kiela (2017)	10	4.736	0.772	46192	151324	**9942**
	GA-DL (Ours)	10	4.788	0.781	41756	144358	11987
	GA-DL-RW (Ours)	10	**4.270**	**0.797**	**38277**	**134541**	12575

Table 6: WordNet Nouns Closure

A.5.4 EFFECTIVENESS OF OUR PROPOSED NEGATIVE SAMPLING

We display a zoom-in of Figure 7b in Figure 8.

(a)   (b)

Figure 7: Effectiveness of negative sampling.

Figure 8: Effectiveness of negative sampling.