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A Descriptive Tolerance Nearness Measure for Performing Graph Comparison

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Abstract. This article proposes the tolerance nearness measure (TNM) as a computationally reduced alternative to the graph edit distance (GED) for performing graph comparisons. The TNM is defined within the context of near set theory, where the central idea is that determining similarity between sets of disjoint objects is at once intuitive and practically applicable. The TNM between two graphs is produced using the Bron-Kerbosh maximal clique enumeration algorithm. The result is that the TNM approach is less computationally complex than the bipartite-based GED algorithm. The contribution of this paper is the application of TNM to the problem of quantifying the similarity of disjoint graphs and that the maximal clique enumeration-based TNM produces comparable results to the GED when applied to the problem of content-based image processing, which becomes important as the number of nodes in a graph increases.

Keywords: Graph matching, Tolerance nearness measure, Graph edit distance, Near sets, Maximal clique enumeration

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1. Introduction

This article proposes the tolerance nearness measure (TNM) as a computationally reduced alternative to the graph edit distance (GED) for performing graph comparisons. The graph comparison problem consists of quantifying the similarity or dissimilarity of graphs [1] and is required by many applications [2]. The TNM is example of a graph comparison measure, while the GED is an example of a graph matching algorithm. The chief difference between graph comparison and graph matching algorithms is that a graph matching algorithm maps similar substructures between the two graphs under comparison to produce a real value reflecting the degree these substructures align. Graph comparison methods are a generalization of graph matching approaches where the only requirement is assigning a real value indicating the degree of similarity between the two graphs, *i.e.* the likeness between the two graphs does not necessarily depend on matching substructures. In fact, the TNM approach proposed here is based on finding maximal cliques distributed across the two graphs in question. Further, it is important to note that the graph comparison problem is commonly solved by graph matching algorithms¹ [1]. This observation provides a basis for the comparison presented in this paper. In other words, both the GED and TNM make judgements regarding nearness and similarity of attributed graphs.

Many distance measures have been proposed to solve the graph comparison problem. The TNM is defined within the context of near set theory, where the central idea is that determining similarity between sets of disjoint objects is at once intuitive and practically applicable. More importantly, near set theory provides a mathematically-rigorous, theoretical framework for making decisions regarding nearness or apartness. The TNM grew out of work on practical applications of descriptive near set theory, and is calculated by using a maximal clique enumeration (MCE) algorithm. To date, the TNM has only been used to quantify the similarity of disjoint sets of objects with associated feature vectors. Here, the TNM is used to quantify the similarity of graphs as a solution to the graph comparison problem. In this regard, the TNM is applied to graphs in two ways: in its original form to assess the similarity between the vertices of two graphs (i.e. two disjoint sets of objects) based on their attributes; and to assess the similarity of two graphs using both vertex and edge information. The presented approach is contrasted with the well-known GED matching algorithm through application to the problem of content-based image retrieval (CBIR) on the Tarragona Rotation Zoom [3, 4], SIMPLIcity [5] and the Palmprint datasets [6]. The GED algorithm was selected due to its popularity and benchmark status within the field of graph comparisons [7]. The contribution of this paper is the application of TNM to the problem of quantifying the similarity of disjoint graphs – in the context of descriptive near set theory – and that the MCE-based TNM produces comparable results to the GED when applied to the problem of content-based image retrieval, which becomes important as the number of nodes in a graph increases. This article is organized as follows. Section 2 gives background on the GED and TNM, Section 3 describes the graph comparison measures and provides a detailed explanation of the GED and TNM. Section 4 presents the results along with discussions, and, finally, Section 5 is the conclusion of the article.

¹The graph comparison problem is also referred to as *graph similarity*, *graph distance*, or *graph measure* in the literature. Examples using this terminology are cited in the last paragraph of Section 2.

2. Background

A comprehensive survey of most graph matching algorithms proposed since the late seventies, together with pattern recognition and machine vision applications, is presented in [8]. With respect to applications, there are numerous examples of solving practical problems using graph similarity measures. Examples include structural pattern recognition [9, 10], graph classification [11], character recognition [12], artificial intelligence [13, 14], chemoinformatics [15], document analysis [16], and bioinformatics [17]. Other interesting examples include shape analysis [18] and 3D segmentation [19].

In this work, the focus is on the popular GED [20, 21, 20, 22], which is considered one of the most well known graph matching approaches since it is generic and easily interpretable [22, 10, 7] (see, [23, 1] for a extensive discussions on the GED). The GED is an error-tolerant, inexact graph similarity measure. Accordingly, it can be applied to graphs with variable number of nodes, such as the graphs formed by the content-based image retrieval (CBIR) application presented in this paper.

The GED was first used for string matching, introduced as the *edit distance*, where it was used to find the minimum cost for all sequences of edit operations required to transform one string to other [24, 25]. This was followed by the adaptation of the edit distance for use in comparing trees [26]. In this case, it was modified to find the minimum amount of edit operations to convert one tree into the other, where each operation was associated with specific cost. Finally, the GED appears in its present form in 1983 [27, 21, 20]. Since then, it has been used in many applications: fingerprint classification [28], graph clustering problems [29], chemical graph recognition problem [30], molecular electrostatic potentials comparison [31] and diatom identification [32]. Some of the more recent examples include: database management [33], biological network alignment [34] and handwriting recognition [12, 35]. In this paper, the GED and the TNM approach are both applied to the problem of CBIR.

While the GED is well established, the TNM was recently introduced in 2010 [36] within the context of descriptive near sets theory². In general, the TNM follows from a rich and rigours mathematical history on the study and formalism of the notion of nearness [37]. J. B. Listing and F. Hausdorff first considered the ideas of near and far within a mathematical context. They were followed by J. H. Poincaré's work on resemblance and similarity, which was inspired by kinesthesia experiments performed by E. Weber in 1834, and G. T. Fechner's sensation sensitivity experiments [39, 40, 41, 42]. Poincaré's reflections led him to introduce resemblance in representative spaces (called physical continua) [43, 44, 45], where his work marked a shift from stimuli and sensations to an abstraction in terms of mathematical sets together with an implicit idea of tolerance. Only a few years later, F. Riesz published a paper in 1908 on the nearness of sets [46, 47, 48], initiating the mathematical study of proximity spaces and the eventual discovery of descriptively near sets. During the 1960s, E.C. Zeeman introduced tolerance spaces in modelling visual perception [49], and A.B. Sossinsky observed in 1986 [50] that the main idea underlying tolerance space theory comes from Poincaré, especially [44]. The idea of near set theory – now known as descriptively near set theory – was inspired by a collaboration in 2002 by Z. Pawlak and J. F. Peters on a philosophic poem entitled "How Near" [51]. In 2006, a formal approach to the descriptive nearness of objects was considered by J. Peters, A. Skowron and J. Stepaniuk [52, 53] in the context of proximity spaces [47, 54, 48, 55]. This work led to the

²A nice history of near set theory is available in [37, 38].

introduction of descriptively near sets [56, 57], which was soon followed by tolerance near sets in 2009 [58, 59]. Tolerance near set theory builds on near set theory through the use of tolerance spaces and relations (based on work by Zeeman [60] and Sossinsky $[50]^3$). Finally, the TNM was introduced in [36], which has been applied to the problems of image analysis and CBIR [62, 63, 64, 65, 66] and forms the main contribution of this article.

Finally, there is a large body of work on graph comparison methods other than those used in graph matching. This section concludes with recognition of these approaches. Tao et al. compare and contrast a graph measure defined with respect to topological indices with the GED [7]. Wallis et al. present an extension of the maximum common subgraph distance measure to propose a new graph distance metric. Similarly, [67] propose a new graph metric for measuring the similarity of graphs by identifying the relationship between the maximum common subgraph and the minimon common supergraph, and a new graph distance metric based on maximal common subgraph is presented by [68]. [2] present an approach for graph similarity scoring and matching based on structural similarity of local neighbourhoods. [69] demonstrate that graph similarity measures based on (sub)graph isomorphism and error-tolerant graph matching are a special case of a generic similarity measure proposed by [70]. A graph comparison methods for large graphs with hundreds of nodes is proposed in [71], and [72] present a graph similarity measure with low computational complexity based on the structural similarity of graphs. [73] propose a graph comparison measure based on the mutual information between graphs. Two further applications of graph comparison outside of graph matching are the work by [74] presenting an application of graph similarity measures for assessing attacks in dynamic networks, and an application of a pairwise graph comparison algorithm to the problem of automatic detection of conserved gene clusters reported by [75].

3. Graph Comparison Measures

The notion of quantifying the differences between two objects can be formalized through the definition of a similarity measure $d: X \times X \to \mathbb{R}$, where X is the representation space, containing a diversity of objects and their (multi-dimensional) descriptions, and \mathbb{R} is the set of real numbers. Further, d is considered a metric if conditions of non-negativity, uniqueness, symmetry and triangle inequality are satisfied. In this work, objects in X are attributed graphs and d provides a basis for decisions regarding the similarity of these graphs, which is known as graph comparison [1]. Here, the focus is on attributed graphs defined as a four tuple $G = (V, E, \alpha, \beta)$, where V is the set of vertices, E is a set of edges consisting of ordered pairs among elements $v \in V$, and α (resp. β) are functions that produce feature vectors containing information about node (edge) labels. With these definitions in mind, the remainder of this section describes the well-known GED metric and the recently reported tolerance nearness measure (TNM) for use in error-tolerant attributed graph matching.

³See, *e.g.*, [61]

3.1. Graph Edit Distance

The graph edit distance (GED)⁴ is the most common and well know method for performing errortolerant graph matching [10]. Briefly (and using the same notation as [76]), the GED is a measure of the amount of distortion required to transform one graph into another. Distortion actions, called edit operations, include insertion, deletion, and substitution of both nodes and edges. A series of distortions used to transition G^p into G^q is called an edit path, and is defined as $editPath(G^p, G^q) = (\epsilon_1, \dots, \epsilon_k)$, where ϵ_i represents edit operation i in this process of morphing one graph into another. The set of all edit paths for two graphs is labelled v, and for each $editPath \in v$ there is an associated bijection $f^{p,q} \in T$, where a bijection $f^{p,q} : V^p \to V^q$ assigns one vertex in G^p to only one vertex in G^q and T is the set of all possible bijections. Continuing, there is an associated cost function for each edit operation. Cost functions are combined to form an overall edit cost function, $EditCost(G^p, G^q, f^{p,q})$, used to find the minimum amount of operations required to transform G^p into G^q (defined in [77]) [76]. Finally, the GED is defined as $EditDistance(G^p, G^q) = \min_{f^p, q \in T} EditCost(G^p, G^q, f^{p,q})$.

The GED is generally determined by the A^* algorithm [78] when exactly evaluating the EditDist $ance(G^p,G^q)$ [76]. However, the complexity of this approach is exponential in the number of nodes. As a result, approximate methods are necessary when working on anything but small examples. Approximate methods are appropriately named sub-optimal because they strike a balance between computational cost and non-exact GED values. While there are many sub-optimal algorithms (see, e.g., [76]), this article focuses on the bipartite algorithm for generating GED values since it is considered one of the best sub-optimal algorithms [76], and the results presented here were generated using the implementation reported in [10], which also uses this algorithm. There are three variants of the bipartite algorithm, namely the original approach reported by [27], fast bipartite matching [76], and, recently, the square fast bipartite matching [13]. The fast bipartite matching algorithm was used for this article. Lastly, given graphs G^p , G^q of order n, m, respectively, the time complexity for the bipartite algorithm is $O(n+m)^3$ and the time complexity for both the fast and square fast bipartite matching algorithms is $O(max(n,m)^3)$ [13].

3.2. Tolerance Nearness Measure

The focus of descriptive near set theory is to assess similarity in terms of the descriptions of objects within the sets. Until this article, the tolerance nearness measure (TNM) was used to quantify the similarity between two sets of objects, where these sets of objects were not graphs (see [65] for a detailed discussion). This process involves creating a single attributed graph, G^{XY} , based on the union of the two sets, $X, Y \subseteq O$, where the objects in O are used as graph vertices, i.e. $V^{XY} = X \cup Y$, and edges between vertices in V^{XY} are created based on the distance between feature vectors associated with each $v \in V^{XY}$. Specifically, a binary relation is defined between vertices based on their feature vectors, and edges between vertices are formed if the two objects satisfy the relation. Using G^{XY} , calculating the TNM requires finding all the maximal cliques, where, for each clique, the distribution of the objects between the original two sets is used to produce the final TNM value (see below).

⁴See [76] for an in-depth description of the GED.

⁵ [53] details the difference between traditional spatial nearness relations and relations defined in terms of descriptive nearness relations.

While useful, this process is not suited to assessing the similarity of graphs since new edges are produced during the calculation. For this reason, two approaches to using the TNM for quantifying the similarity of graphs are presented: one based solely on the vertices of the two graphs (using the original TNM approach), and the method presented below that incorporates the existing graph edge information in the calculation. Both of these approaches are graphically depicted in Figure 1.

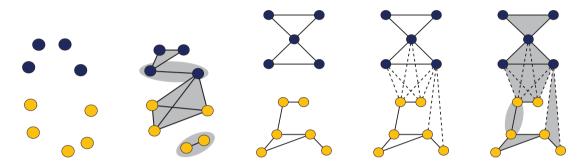


Figure 1. Visual example of two approaches for calculating the TNM. (a) and (b) represent the original approach, where (a) contains two sets of objects (represented by two colours) and (b) shows the edges created with a tolerance relation and the grey areas are maximal cliques. (c) - (e) depict the TNM in quantifying the similarity of graphs. (c) contains the original graphs (with edges), (d) shows new edges formed between the graphs using a tolerance relation, and the grey regions in (e) are maximal cliques.

Formally, the TNM is defined within the context of a perceptual system, which is denoted as $\langle O, \mathbb{F} \rangle$. A perceptual system consists of a non-empty set O of sample perceptual objects and a non-empty set \mathbb{F} of real-valued functions $\phi \in \mathbb{F}$, where $\phi : O \to \mathbb{R}$ [79, 36]. Recall, for this application, objects in O are graph vertices, and, for an attributed graph, each vertex has an associated feature vector generated by α . Within a perceptual system, α is defined as a feature vector given by $\alpha(x) = (\phi_1(x), \phi_2(x), \dots, \phi_i(x), \dots, \phi_l(x))$, where l is the length of the vector α , and each $\phi_i(x)$ in $\alpha(x)$ is a probe function value [80] forming part of the description of $x \in O$. In the descriptive near set literature, the feature vector α is called an object description.

The first step in finding the TNM between two sets of objects is the creation of G^{XY} using a binary tolerance relation [61, 36]. Briefly, a tolerance relation is one that is reflexive and symmetric, but not transitive. The specific relation used in this work is defined as follows. Given $\varepsilon \in \mathbb{R}_0^+$ and \mathbb{F} , the perceptual tolerance relation \cong_{ε} is defined as

$$\cong_{\varepsilon} = \{(x, y) \in O \times O : \| \alpha(x) - \alpha(y) \|_{2} \le \varepsilon\}, \tag{1}$$

where $\|\cdot\|_2$ is the L^2 norm [36]. Using \cong_{ε} , E^{XY} is formed by adding an edge between $x,y\in V^{XY}$ if, and only if, (iff) two objects satisfy this relation, *i.e.* the distance between the two feature vectors is less than ε . Fig. 1(a) & 1(b) give an example of forming edges in this manner. Now that G^{XY}

 $(V^{XY}, E^{XY}, \boldsymbol{\alpha}, \boldsymbol{\beta})$ has been defined the TNM, between sets X and Y, is given as

$$TNM(G^{XY}, V^{X}, V^{Y}) = 1 - \left(\sum_{C \in H} |C|\right)^{-1} \cdot \sum_{C \in H} |C| \frac{\min(|C \cap V^{X}|, |C \cap V^{Y}|)}{\max(|C \cap V^{X}|, |C \cap V^{Y}|)}, \quad (2)$$

where H is the set of all maximum cliques obtained from G^{XY} , i.e. each $C \in H$ is a set of vertices representing a maximal clique in G^{XY} . Note, implicit in this definition is the requirement that H must be extracted from G^{XY} prior to performing the calculation given in Eq. 2.

As highlighted in the introduction, the focus of this article is the observation that Eq. 2 can also be used to assess the similarity between two graphs G^p and G^q . In this case, G^{pq} is formed in the following manner. The set of vertices is simply $V^{pq} = V^p \cup V^q$. E^{pq} , on the other hand, is obtained by the following two steps. First, set $E^{pq} = E^p \cup E^q$. Then, each $v_p \in V^p$ is compared to each $v_q \in V^q$, and an edge is only added to E^{pq} iff $v_p \cong_{\varepsilon} v_q$. Creating E^{pq} in this manner is illustrated in Fig. 1(c) - 1(e), and Algorithm 1 contains the pseudocode for calculating the TNM using this approach. Note, edges are only added between the graphs, not between vertices within the same graph. Also, $MCE_{BK}(V^{pq}, E^{pq})$ performs maximal clique enumeration using the Brong-Kerbosh algorithm [81], and it returns a collection (family) of sets, each consisting of the vertices contained in a maximal clique. In Line 8, the cliques are chosen from H in the order in which they are generated by the MCE algorithm $MCE_{BK}(V^{pq}, E^{pq})$.

Notice that the complexity for calculating TNM depends on the approach used to find all the maximal cliques in a graph. Specifically, the Bron-Kerbosh algorithm [81] was used in this work to produce H. While the results reported here use the original Bron-Kerbosh algorithm, the addition of adding pruning methods gives this algorithm a worst-case running time complexity of $O(3^{i/3})$ for a graph of n vertices. This is optimal since there exists up to $3^{i/3}$ maximal cliques in a graph with n vertices [82]. However, the TNM algorithm also consists of either: finding edges between all nodes when comparing two sets of objects (of sizes n and m), or only finding edges between the two sets of vertices when comparing graphs (also of sizes n and m). Thus, the final worst case runtime complexities are $O((n+m)^2+3^{(n+m)/3})=O((n+m)^2)$ and $O(\max(n,m)^2+3^{(n+m)/3})=O(\max(n,m)^2)$, respectively. Observe that both of these complexities are lower than those reported for the GED above.

4. Results and Discussion

This section compares the GED and TNM through their use in performing content-based image retrieval (CBIR) on two different datasets. CBIR is the task of retrieving all the images from a digital image dataset that match a given query image. The goal is to retrieve all the perceptually similar images, and in most cases rank the relevance of each image to the query image. Besides subjective evaluation of the results, CBIR methods are typically compared using datasets partitioned into cat-

 $^{{}^{6}\}beta$ is not used in this work, *i.e.* there is no extra information associated with edges.

Algorithm 1: TNM(edges) Algorithm

```
Input: G^p, G^q, \varepsilon

Output: TNM(G^{pq}, V^p, V^q)

V^{pq} \leftarrow V^p \cup V^q; E^{pq} \leftarrow E^p \cup E^q;

for v_p \in V^p do

| for v_q \in V^q do

| if v_p \cong_{\varepsilon} v_q then

| E^{pq} \leftarrow E^{pq} \cup (v_p, v_q);

| end

end

end

H = \text{MCE}_{\text{BK}}(V^{pq}, E^{pq});

w \leftarrow 0; TNM \leftarrow 0;

for C \in H do

| w \leftarrow w + |C|;

TNM = TNM + |C| \frac{\min(|C \cap V^p|, |C \cap V^q|)}{\max(|C \cap V^p|, |C \cap V^q|)};

end

TNM \leftarrow 1 - TNM/w;
```

egories of images. Taking this approach allows the use of metrics such as precision and recall to evaluate, compare, and contrast the reported methods.

In comparing the two measures in this article, test datasets used in previously published GED and TNM papers were employed. Consequently, the datasets used in this work are the Tarragona Rotation Zoom [3, 4], SIMPLIcity [5] and the Palmprint datasets [6]. The Tarragona Rotation Zoom dataset was selected as this article uses the same GED implementation from [10] – as well as the learnt edit-costs identified in the paper. SIMPLIcity was similarly chosen for its use in earlier TNM papers [36, 65]. The Palmprint dataset was picked to demonstrate the increase in computational complexity as the size of the graph grows.

The Tarragona dataset consists of four categories: boat, east park, east south, and residence. Each category consists of 10 images, where each image produces a single graph. The features associated with each vertex in a graph are extracted using the SpeedUp Robust Features (SURF) algorithm [83] and the edges between the vertices are created using Delaunay triangulation [84]. Each graph consists of 50 vertices and 64 features. There are no attributes associated with graph edges.

The SIMPLIcity dataset contains 10 categories, where each category consists of 100 images. Again, a single graph is extracted from each image, consisting of 228 vertices and 18 features per vertex. Features are extracted using texture based techniques, moments in CIELUV colour space, edge based features and Zernike moments [36]. Edges between vertices are created using Eq. 1, where an edge is formed if two vertices satisfy the perceptual tolerance relation. The graphs used to generate the results were created using $\varepsilon = 0.2^7$ as this value produced the best results in [65]. The

⁷See [36] for a discussion on how the TNM varies with ε .

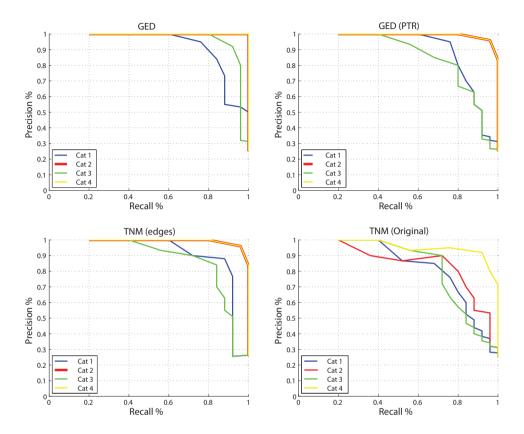


Figure 2. Average precision vs. recall plots for the GED and TNM on the Tarragona dataset. (a) GED, (b) GED (PTR), (c) TNM (edges), and (d) TNM (Original).

value of ε is typically selected based on trial and error and falls into the range $[0,\sqrt{l}]$, where l is the feature vector length, $\varepsilon=0$ produces no edges and $\varepsilon=\sqrt{l}$ produces a fully connected graph.

A subset of the Palmprint dataset is extracted because of very high computational time which contains 4 categories and each category consits of 8 images (graphs). The dataset contains variable number of nodes. Maximum nodes in a graph are 1385 and minimum is 574, where the average number of nodes is \sim 841. There are total 5 features associated with each nodes *i.e.* position, type, quality and type of minutiae. The features associated with each node in a graph are extracted using the algorithm described in [85] and the original edges between the nodes were created using delaunay triangulation with no attributes [6]. The edges used in this experiment are created using the perceptual tolerance relation with $\varepsilon=0.4$.

The results are compared in this article using precision and recall [86] and are given in Tables 1 & 2 & 3 and Figures 2 & 3 & 4 Here it is important to note a few points. First, the TNM results presented here are broken into two categories, namely the original approach (depicted in Figs. 1(a) & 1(b)) and the method that incorporates the existing graph edge information in the calculation (depicted in Figs. 1(c)-1(e)). These two approaches are labelled *original* and *edges*. Next, with respect to the SIMPLIcity & Palmprint dataset, the TNM *original* and *edges* results are the same

since the SIMPLIcity & Palmprint graphs are created with the perceptual tolerance relation. Thus, only one set of figures is presented. Lastly, the Tarragona dataset consists of graphs, however, for comparison with the TNM, new edges are also created between vertices using the perceptual tolerance relation. These results are labelled GED (perceptual tolerance relation (PTR)) and were generated using $\varepsilon=1.5$. This value of epsilon was selected as it produces graphs which generate the same final learnt costs when using the approach reported in [10].

 $Table \ 1. \quad Comparison \ of \ GED \ and \ TNM \ on \ Tarragona \ dataset \ using \ Precision(P) \ and \ Recall(R)$

Category	GED		GED (PTR)		TNM (edges)		TNM (Original)	
	P	R	P	R	P	R	P	R
Boat	0.9500	0.7600	0.9500	0.7600	0.9000	0.7200	0.8500	0.6800
East Park	1.000	0.8000	1.000	0.8000	1.000	0.8000	0.9000	0.7200
East South	1.000	0.8000	0.8500	0.6800	0.9000	0.7200	0.9000	0.7200
Residence	1.000	0.8000	1.000	0.8000	1.000	0.8000	0.9500	0.7600
Average	0.9875	0.7900	0.9500	0.7600	0.9500	0.7600	0.9000	0.7200
Timing (ms)	43.33		45.71		40.29		16.29	

Table 2. Comparison of GED and TNM on SIMPLIcity dataset using Precision(P) and Recall(R)

Category	G	ED	TNM		
	P	R	P	R	
1. Africans	0.6635	0.1327	0.7230	0.1446	
2. Beaches	0.5245	0.1049	0.4755	0.0951	
3. Buildings	0.4385	0.0877	0.4745	0.0949	
4. Buses	0.6985	0.1397	0.5145	0.1029	
5. Di- nosaurs	1.000	0.2000	1.000	0.2000	
6. Elephants	0.7680	0.1536	0.7210	0.1442	
7. Flowers	0.9470	0.1894	0.8615	0.1723	
8. Horses	0.8375	0.1675	0.7360	0.1472	
9. Mountains	0.6425	0.1285	0.6325	0.1265	
10. Food	0.7130	0.1426	0.7425	0.1485	
Average	0.7233	0.14466	0.6881	0.13762	
Timing (ms)	11.36		0.61		

The results were generated as follows. First, a single image was selected as the query image, which

Category	G]	ED	TNM		
	P	R	P	R	
1. Cat 1	0.6875	0.3438	0.6563	0.3281	
2. Cat 2	0.4063	0.2031	0.5625	0.2813	
3. Cat 3	0.5938	0.2969	0.6875	0.3438	
4. Cat 4	1	0.5000	1	0.5000	
Average	0.6719	0.3359	0.7265	0.3633	
Timing (s)	1127.348		283.144		

Table 3. Comparison of GED and TNM on Palmprint dataset using Precision(P) and Recall(R)

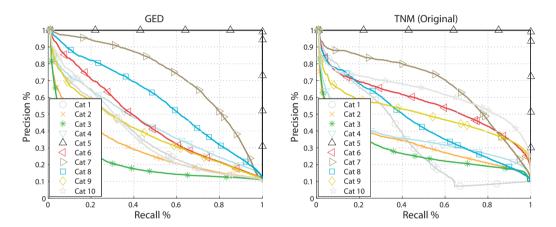


Figure 3. Average precision vs. recall plots for the GED and TNM on the SIMPLIcity dataset. (a) GED, and (b) TNM.

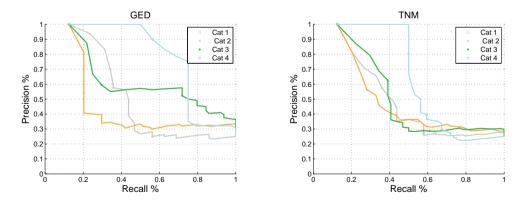


Figure 4. Average precision vs. recall plots for the GED and TNM on the Palmprint dataset. (a) GED, and (b) TNM.

was compared to each other image in the dataset using either the GED or TNM approaches mentioned above. The measure values were then sorted and the image category information was used to calculate precision and recall. This process was then repeated for each image in the dataset. Finally, precision and recall values were averaged across each category and the results are giving in Figures 2 & 3 & 4. Similarly, the results reported in Tables 1 & 2 & 3 represent the values obtained for the top 4, top 20 and top 4 images, respectively, in each category. Note, this process consisted of 210 comparisons for the Tarragon dataset, 500,500 comparisons for SIMPLIcity and 528 comparisons for the Palmprint.

Given this information, some interesting observations about the results can be made. Begining with Tables 1 & 2 notice that the TNM results are all quite similar to the GED. With respect to the original TNM (using only vertex information) and the Tarragona dataset, TNM precision values are lower than both GED and GED (PTR)⁸. Recall values are closer than precision, but still all lower than GED. The TNM using the graph edge information performs better, where precision and recall values tie GED in East Park and Residence categories. The better performance makes sense since more of the original, domain-specific graph information is used. With respect to the SIMPLIcity dataset, TNM out performs GED in the Africans, Buildings, and Food categories. As before, GED performs better in the majority of cases, but the results are close. With respect to Figures 2 & 3, it is harder to make a direct comparison. In general, Figure 2 shows that TNM produces comparable results to the GED, where GED performs slightly better. On the other hand, from Figure 3, TNM performs better in the Beaches, Buildings, Elephants, and Mountains categories, while GED performs better in Buses, Horses, and Food categories. Both measures perform comparably in categories Africans, Dinosaurs, and Flowers. Table 3 shows the results for the Palmprint dataset for both TNM and GED; where GED performs slightly better than TNM in first category. On the contrary, the TNM performs better than GED in second and third category and the GED & the TNM tie results for fourth category. The overall average precision and recall represents that the TNM performs better than the GED with much less computational time for single comparison. Figure 4 shows the results for both TNM and GED are very close, however, the TNM performs better in most of the categories.

The average time per graph comparison for each approach are given in the bottom of Tables 1 & 2 & 3. Notice, as expected, the runtime for the TNM approach is less than the GED approach in all cases. Further, the TNM(Original) approach is much faster than the TNM(edges) approach since the graphs produced by the perceptual tolerance relation with $\varepsilon=1.5$ contain less edges than the original Tarragona graphs. Likewise, the timing results presented in Table 3 shows that computational time increases with the increase in size of graph which in this case is much less for the TNM than the GED. Finally, these timing results were generated using Matlab version R2012a on a system running Windows 10 with an Intel i5-4200U CPU @ 1.6GHZ and 4GB or memory.

5. Conclusion

This article presented a new application of the TNM for use as a graph comparison measure, and demonstrated it is a competitive alternative to the popular and well-known GED. Graph-based results were presented with respect to the original TNM definition (using only vertices in the calculation) as

⁸Recall, GED (PTR) uses the same edges as used in the TNM calculation (rather than the edges that are present in the dataset).

well as a new approach that also uses graph edge information. The results demonstrate that both measures produced comparable precision and recall values. Further, the TNM approach is advantageous in that it is not dependent on costs, *i.e.* cost functions must be defined for each edit operation in the GED, whereas the TNM approach is only dependent on ε from Eq. 1. Additionally, the runtime complexity of the TNM was shown to be less than that of GED, which is important as graph sizes grow ever larger. Future work will consist of exploration into learning optimal ε values for a given problem domain, and applications of these methods to other datasets since results suggest TNM may perform as good (or better) than GED in many real world applications.

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