

K-Means Graph Database Clustering and Matching for Fingerprint Recognition

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Abstract

The graph can contain huge amount of data. It is heavily used for pattern recognition and matching tasks like symbol recognition, information retrieval, data mining etc. In all these applications, the objects or underlying data are represented in the form of graph and graph based matching is performed. The conventional algorithms of graph matching have higher complexity. This is because the most of the applications have large number of sub graphs and the matching of these sub graphs becomes computationally expensive. In this paper, we propose a graph based novel algorithm for fingerprint recognition. In our work we perform graph based clustering which reduces the computational complexity heavily. In our algorithm, we exploit structural features of the fingerprint for K-means clustering of the database. The proposed algorithm is evaluated using realtime fingerprint database and the simulation results show that our algorithm outperforms the existing algorithm for the same task.

Keywords

Pattern Recognition, Fingerprint Matching, Graph Matching, Clustering

1. Introduction

The graph is an important data structure for representing various multi-attributed data and their complex relations. Pattern recognition is the area of computer science which intends to find a particular pattern in the given input. A variety of graph based techniques had been proposed as a powerful tool for pattern representation and classification in the past years. For a longer duration, graphs were considered to be computationally expensive tool. But recently the graph based pattern recognition and image processing is gaining popularity. It is a fact that the computational complexity of the graph based methods is becoming feasible due to high end new generations of the computers and the research advancements. In this work, we exploit graph based fingerprint recognition. Fingerprint plays a distinguishing role in biometrics. It gives unique identification to the individual. It is permanent and non-changing character pattern over a long period of time. The fingerprint recognition consists of retrieving specific fingerprints from database. The characteristics or features of a fingerprint are extracted and used in identification. We have used attributed graph to represent the extracted features of the fingerprint. When the feature set size grows, graph matching becomes expensive. So we optimize the fingerprint database by K-Means clustering which drastically reduces the fingerprint matching time. In this section, we discuss some basic graph terminologies which are prerequisite for the further discussion. We also discuss the related literature in this section 2, we present the proposed algorithm. Section 3 discusses simulation results and section 4 concludes the paper. A graph G = (V, E) is a set of vertices V and edges E. V is the set of vertices and $E \subseteq (V \times V)$ is the set of edges of graph G. The order or size of a graph G is defined as the number of vertices of G and it is represented as |V|. If two vertices in G, say $u, v \in V$, are connected by an edge $e \in E$, this is denoted by e = (u, v) and the two vertices are said to be adjacent to each other. Edges can be directed or undirected. Graph matching can be applied to directed and undirected graphs both. A directed graph G = (V, E) is called complete when there is always an edge $(u_1, u_2) \in E = V \times V$ between any two vertices u_1, u_2 in the graph. When graphs are used to represent objects, images or fingerprints, vertices usually represent regions or features of the object or images, and edges between them represent the relations between the features.

Graph vertices and edges can contain large complex information. When this information is a label or attributes of vertices and edges, the graph is called an attributed graph. Formally, an attributed graph is a graph with six-tuple $G = (V, E, u, v, L_v, L_E)$, where

- *V* is a finite set of nodes,
- $E \in V \times V$ is a finite set of edges,
- *u* is a labeling function for vertices, $V \rightarrow L_V$
- v is a labeling function for edges, $E \rightarrow L_E$
- L_v is a set of node labels, and
- L_E is a set of edge labels

The two graphs *G* and *G'* can be compared by determining a mapping function *f* which associates nodes and edges of *G* with nodes and edges of *G'* and vice versa. A graph isomorphism between two graphs *G* and *G'* is given if there exists a bijective mapping *f* from the nodes of *G* to the nodes of *G'* such that the structure of the edges as well as all node and edge labels are preserved under *f*. More precisely: A bijective function $f: V \to V'$ is a graph isomorphism from a graph $G = (V, E, u, v, L_V, L_E)$ to a graph $G' = (V', E', u', v', L_V, L_E)$ if:

1) u(v) = u'(f(v)) for all $v \in V$

2) for any edge $e = (v_1, v_2) \in E$ there exists an edge $e' = (f(v_1), f(v_2)) \in E'$ such that v(e) = v(e'); for any $e' = (v'_1, v'_2) \in E'$ there exists an edge $e = (f^{-1}(v'_1), f^{-1}(v'_2)) \in E$ such that v(e') = v(e).

Formally, two graphs are isomorphic to each other if they are equal. Practically, for most applications this mapping is too much restrictive. In practice, graph isomorphism is of limited use and more relaxing concepts are necessary. One such approach is subgraph isomorphism. An injective function $f: V \to V'$ is a subgraph isomorphism from a graph G to a graph G' if there exists a subgraph $G_s \subseteq G'$ such that f is a graph isomorphism from G to G_{s} . The literature has focused on statistical and structural approaches for fingerprint classification. Statistical methods adopt the extraction of various mathematical characteristics to classify the patterns [1]. The structural approaches make use of syntactic [2] or structural constructs of the patterns [3] for the recognition methods. In the syntactic approach fingerprints are represented as grammar and the parsing techniques are used to classify the grammar. However some hybrid approaches have also been explored which combine the structural and statistical methods [4] [5]. The graph plays an important role in the structural and syntactic methods of fingerprint classification. In these methods the fingerprint patterns are represented by graphs or production rules in the grammar [6]. Graph isomorphism, sub graph isomorphism or grammar parsing is used to classify the fingerprints to their respective classes. Graph isomorphism and sub graph isomorphism approaches must be able to incorporate errors and distortions for the possibility of noisy fingerprint data. The syntactic approach [6] defines grammar in terms of terminal and non terminal symbols and production rules. The complex attributes considered in syntactic patterns make them difficult to build the parsing methods. The disadvantage is, it has high computational complexity [6]. The graph based methods can effectively use relational attributes of the fingerprint to build the features graph. The graph based fingerprint matching algorithm is highly dependant on the combination of fingerprint features and corresponding graph model used to represent the fingerprint. The effective graph prototypes representing "core" and "deltas" of the fingerprints were used in [7]

[8]. The graph derived from the fingerprint is classified to its corresponding owner class by using inexact graph matching [9]. The use of inexact graph matching makes the algorithm robust to absorb distortions in the fingerprints. The graph based fingerprint matching can be optimized by reorganization of the graph data in a preprocessing step. The graph database clustering effectively reduces the number of searches in the graph data. Various graph clustering policies like a hierarchical [10], indexed [11] [12], hashed [13] databases have been suggested for various pattern matching applications. Ultimate goal is to reduce the overall graph matching time by limiting the search area. A decomposition based hierarchical approach avoids matching each model graph individually onto the input graph [14]. The model graph is recursively decomposed offline into smaller subgraphs in [14]. At run-time, these subgraphs are matched onto the input graph and all detected subgraph isomorphisms are combined to form subgraph isomorphisms for complete model graphs. The main advantage of scheme is that subgraphs that appear multiple times in the same or in different model graphs must be matched only once onto the input. Consequently, the corresponding subgraph isomorphism detection process will be more efficient than the sequential matching of the input graph with each of the models. When all model graphs are highly similar, the method becomes independent of the size of the database. and algorithm's best case complexity becomes O(IM), while worst case complexity goes to $O(I^M M^2)$ where N is total number of model graphs in the database. I is the number of vertices in the input graph and M is number of vertices of the model graph [14]. For model graphs with some common substructures the algorithm is sublinearly dependent on the size of the database. Then the algorithm's best case takes O(NIM) amount of time, while worst case takes $O(NI^{M}M^{2})$ amount of time [14]. For large model graphs the algorithm is faster than traditional algorithms due to reappearing substructures that naturally evolve in large graphs. But for unlabeled, highly connected graphs, the algorithm performs poorly [14]. Median graph is an important concept used in graph matching. It acts as a representative for a set of graphs by extracting the essential information from them into a single prototype. Given a set of graphs S, the median is derived from S which has the smallest sum of distances (SOD) [15] to all the graphs in the set. Computing the median graph is an NP Complete problem [16]. The algorithm uses the concept of dissimilarity distance to compare the two graphs. If the the median itself belongs to the set of graphs, the set median is obtained, otherwise the generalized median is obtained. The algorithm guarantees that a good solution will be found [15]. The efficiency of the algorithm depends on reduction of the set of node labels and on a search strategy aiming at exploring the most likely median graph candidates. The graph can also be stored by means of hashing and indexing [13]. Each graph has a numerical key associated with it. A hash function maps this key onto the array of manageable size. The address thus calculated for an object is called as the hash code of the object. Database construction consists of computing such a hash-code for each object to be stored in the database. Several objects may have the same address. In such case a list of objects will be associated with each address. So the database becomes a table of lists, where each list stores a list of objects with same hash code. Indexing is the process of computing hash code for the unknown object for finding whether such object exists in the hash table or not. By using all the mathematical calculations based on Laplacian transform, [13] presented the graph hashing method. The method characterizes the graph by using its mathematical and computational properties [13]. The Laplacian matrix is a richer graph description than the well known adjacency matrix. However the indexing procedure has some limitations [13]. The decomposition of a graph into subgraphs is not optimal and it does not take into account the geometrical and structural constraints. Even if graphs are good for feature representation, they are harder to manipulate and operate. Some attempts have been made to combine the best of the graph and the vector domains [16] [17] in order to get the advantages of both. The median graph can be seen as the representative of a set of graphs. These methods generate approximate median graph using real databases containing large graphs. The approximate K-Means based algorithm computes the generalized median graph [18] for graph clustering in content-based image retrieval and analysis of DNA structures [18]. The median graph \overline{g} is the minimization of

$$\overline{g} = \arg\min_{g \in U} \sum_{i=1}^{n} d(g, g_i)$$
(1)

where U is the set of all graphs that can be constructed from a given set $S = \{g_1, g_2, \dots, g_n\} \subset U$ *i.e.* any graph whose number of nodes is between 1 and the sum of all the nodes of the g_i .

2. The Proposed Algorithm

In this work we present a new algorithm for automated graph based fingerprint representation and matching. We

achieve the promising results by clustering the graph feature set and then applying the graph isomorphism. The algorithm proves robust for noisy images as well. We extract various structural features like ridges, end points, bifurcations of the fingerprint. These end points and bifurcations are called as minutiae. Each fingerprint is represented as an attributed relational graph [19] where these features are represented as nodes of the graph. The edges of the graph represent relation attributes between the vertices of the graph. In this way the graph structure captures the structural relationships within the fingerprint graph. The algorithm is implemented and tested using a database of real fingerprint images. Most of the fingerprint recognition algorithms deal with large databases. So comparing the fingerprint with all images in the database takes long time and needs lots of computations. The algorithm performs two tier clustering of graphs. In the first level of clustering fingerprint features are clustered. In the second level the fingerprint graph database is clustered. The novelty of algorithm lies in the following considerations.

- 1) It uses graph as fingerprint data structure.
- 2) It optimizes fingerprint search by clustering fingerprint graph feature nodes (Tier 1 clustering).
- 3) It further optimizes the search by clustering fingerprint database (Tier 2 clustering).

2.1. Graph Based Fingerprint Recognition

This is a graph based approach to compare structural fingerprint features called as minutiae. We implement graph based algorithm given by [20]. The graph based representation of the features of the fingerprint make the procedure more flexible in nature. Each minutiae is represented as a vertex of the graph. We use attributed relational graph. Each minutiae is labeled with its own attributes and neighborhood features. An edge between two vertices indicates that they share neighborhood information between them. The graph isomorphism is applied to decide similarity or dissimilarity between the two graphs being compared.

The overall method involves following steps.

- 1) Fingerprint preprocessing
- 2) Features Extraction
- 3) Features representation as a graph

4) Graph isomorphism algorithm to classify the fingerprint. We are using ridge (R), ridge ending (E) and ridge bifurcation (B) minutiae for fingerprint recognition. **Figure 1** shows some typical examples of ridges and features extracted. If the feature extraction phase detects "n" minutiae for the fingerprint, an attributed relational graph with "n" nodes is formed. The graph isomorphism is applied between the input fingerprint graph G_I and database fingerprint graphs G_D . The algorithm returns true when the input graph G_I is isomorphic to the graph G_D in the fingerprint database. The graph isomorphism is performed by using linear comparison of each of the vertex from the sample fingerprint images.



Figure 1. Ridges, end points and bifurcations.

2.2. Tier 1 Clustering: Graph Based Fingerprint Recognition by K-NN Clustering of the Minutiae

When the fingerprint minutiae are more in numbers, comparing each one of them becomes very expensive for the machine. So in this algorithm we cluster the graph data by using K-NN clustering based on Euclidean distance between the vertices of the graph. A features template for each fingerprint is build after clustering the neighborhood relative vertices in similar classes. The advantage of this type of clustering is that the graph comparison is first done with the cluster features template of the fingerprint instead of direct comparison of the vertices of the graph. If the cluster features match with the input fingerprint graph then further graph isomorphism is performed. Otherwise the next cluster is visited in a hope to get matching candidate in it. The algorithm involves following steps. The steps 1 to 3 are same as discussed in previous section.

- 1) Fingerprint preprocessing
- 2) Features extraction
- 3) Features representation as a graph
- 4) KNN based graph node clustering
- 5) Graph isomorphism algorithm to classify the fingerprint.

2.3. The K-Nearest Neighbor Clustering

The K-nearst-neighbor (KNN) clustering algorithm measures the distance between a query scenario and a set of scenarios in the data set. It cluster the K nearest neighbor minutiae in the fingerprint space. The distance between two scenarios is computed by using distance function d(x, y), where x, y are scenarios composed of n features such as $x = \{x_1, x_2, \dots, x_n\}$ and $y = \{y_1, y_2, \dots, y_n\}$.

The Euclidean distance metric is chosen to determine the closeness between the minutiae (vertices). The Euclidean distance is defined as the K-NN distance because of the structural neighborhood properties of the minutiae (or vertices). The Euclidean distance is given by:

$$d(x, y) = \sqrt{(x_1 - y_1)^2 + (x_2 - y_2)^2 + \dots + (x_n - y_n)^2} = \sqrt{\sum_{i=1}^n (x_i - y_i)^2}$$
(2)

The K-NN clustering proves to be the best due to its lesser execution time and good clustering accuracy. Generally a node is classified by a majority votes of its neighbors. The nodes are assigned to the class that is closest amongst its K nearest neighbors, where K is a small positive integer. If K = 1, then the node is assigned to the class of its nearest neighbor.

1) Each node within the feature set has a class label in the set, $Class = c_1, \dots, c_n$.

2) The K-nearest neighbors are found by calculating the distance matrix where K is the number of neighbors.

3) The K-closest nodes are analyzed to determine which class label is the most common among the set.

4) The most common class label is then assigned to the node being analyzed.

After graph node clustering the graph based fingerprint matching is performed. The identification of the fingerprint is performed in this module. It reads each fingerprint clustered graph template from the database linearly and compares it with the clustered input fingerprint graph template. If the initial graph template matches then only the total graph isomorphism [20] is applied. The graph matching is continued till the input fingerprint is identified otherwise no match is found.

2.4. Tier 2 Clustering: Graph Based Fingerprint Recognition by K Means Graph Database Clustering

In the earlier section we cluster the features of each fingerprint prior to fingerprint matching. In K-means graph database clustering approach we cluster the fingerprint database based on K means clustering algorithm. It clusters the similar class of fingerprints in the database in similar clusters. It saves further matching time drastically as the fingerprint search needs to search a specific sub part of the database. We compare the performance of our algorithm with the graph based fingerprint recognition [20] which do not apply any database clustering.

3. Simulation Results

For the fingerprint detection we have tested the results on real life data set with 150 images. In fingerprint

recognition we implemented following algorithms

Method 1: Fingerprint recognition with graph matching [20]

Method 2: Fingerprint recognition with K-NN clustering of the features and graph matching

Method 3: Fingerprint recognition with K means graph database clustering

The results are depicted in this section. We notice that the graph clustering is reducing the graph matching time drastically than the time taken by graph matching without clustering. Figure 2 shows sample database used. The features extraction phase extracts the minutiae in the fingerprint. It identifies ridges, endpoints and bifurcations present in the image. An attributed relational graph is build for each fingerprint. These features are stored as a characteristic template for each image. Figure 3 shows the features extracted from the image. Each minutia is identified by its (x, y) position, angle, and its type as end point, ridge, bifurcation etc. Each minutia is represented as a vertex and the features are attributes of the vertex.

In the experiments conducted, we recorded fingerprint processing time and fingerprint matching time as comparison criterions. Figure 4 shows the time difference for fingerprint graph processing/clustering for all the three methods for the same sample set. The graph processing clustering time taken for K-NN feature clustering is much higher than processing time taken by method 1 and method 3.

The graph processing/clustering time difference between method 1 and method 3 is not clearly distinguishable in Figure 4. Figure 5 elaborates the graph processing/clustering time difference between graph processing without clustering (method 1) and graph processing with K-means graph database clustering (method 3).

Figure 6 shows the significant time difference for fingerprint graph matching for all the three methods. Once the fingerprint database is clustered, fingerprint matching takes minimum time in method 3. The method 2 takes intermediate matching time in between method 1 and method 3. The method 1 where no clustering is applied, graph matching time the maximum.

4. Conclusions

In this work we have proposed two tier graph clustering and matching algorithms for fingerprint recognition. The fingerprints are preprocessed and structural features, the minutiae are extracted. The fingerprint is represented as an attributed graph by relating all the features with each other. When a query fingerprint image is given as an input for fingerprint matching, graph isomorphism is applied. For the graph of size "n" vertices the algorithm performs " n^{2} " comparisons. When size of "n" becomes larger, graph matching proves expensive. So we apply tier-clustering; Euclidean distance based K-NN clustering of the feature set. It clusters the feature vertices in similar groups



UgaleHN Right.BMF Figure 2. Few examples of fingerprints used for testing.

UgaleHN Left.BMP

```
<FingerprintTemplate Version="2" OriginalDpi="96" OriginalWidth="260" OriginalHeight="300">
<Minutia X="125" Y="969" Direction="64" Type="Bifurcation" />
<Minutia X="1089" Y="1401" Direction="50" Type="Ending" />
<Minutia X="208" Y="1453" Direction="192" Type="Bifurcation" />
<Minutia X="505" Y="833" Direction="104" Type="Bifurcation" />
<Minutia X="1266" Y="1125" Direction="178" Type="Ending" />
<Minutia X="422" Y="693" Direction="13" Type="Bifurcation" />
<Minutia X="302" Y="375" Direction="178" Type="Ending" />
<Minutia X="135" Y="453" Direction="215" Type="Ending" />
<Minutia X="1260" Y="807" Direction="32" Type="Ending" />
<Minutia X="120" Y="234" Direction="232" Type="Ending" />
<Minutia X="109" Y="854" Direction="64" Type="Bifurcation" />
<Minutia X="802" Y="276" Direction="64" Type="Ending" />
<Minutia X="859" Y="328" Direction="141" Type="Ending" />
<Minutia X="693" Y="688" Direction="178" Type="Ending" />
<Minutia X="1073" Y="469" Direction="242" Type="Bifurcation" />
<Minutia X="1161" Y="943" Direction="64" Type="Ending" />
<Minutia X="557" Y="635" Direction="192" Type="Ending" />
<Minutia X="146" Y="1479" Direction="87" Type="Ending" />
<Minutia X="458" Y="958" Direction="77" Type="Ending" />
<Minutia X="1172" Y="661" Direction="178" Type="Ending" />
```

Figure 3. Feature extraction like ridges, end points, bifurcations.



Figure 4. The difference in fingerprint processing time needed for all three methods. Method 1: Graph based fingerprint recognition (without clustering), Method 2: Graph based fingerprint recognition by K-NN clustering of the minutiae, Method 3: Graph based fingerprint recognition by K means graph database clustering.



Figure 5. The difference in fingerprint processing time needed for Method 1 and Method 3. Method 1: Graph based fingerprint recognition (without clustering) Method 3: Graph based fingerprint recognition by K means graph database clustering.



Figure 6. The difference in fingerprint matching time needed for Method 1, Method 2 and Method 3.

based on their neighborhood relationships. Because of the clustering, the algorithm compares cluster properties before comparing contents of the cluster individually. The approach drastically reduces the time taken by comparison without clustering. Further in tier-2, the fingerprint database is clustered with K-means algorithm.

We have tested the results on real time data set available with 150 fingerprints. The observation is that graph clustering definitely reduces time needed for the graph matching purpose. The results of graph matching remain the same. The size of graph, we are considering is restricted due to memory limits of the processing machine. These techniques increase the processing time though. In that case, it is expected that the clustering of the databases is to be done off line prior to graph matching process for best results as a combination.

In the future work, we intend to analyze and test the results on larger data sets along with detail performance evaluation of the algorithms. Definitely graphs based algorithms have their own pros and cons. The selection of appropriate graphs for specific applications with appropriate graph matching algorithm is the key. It is always a challenge to achieve a general graph based platform for various applications and their needs.

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