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# Biometric Identification based Scores Discretization on Multi Forms of Image Structures

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Abstract

Multi-Biometric Identification is one of a well-known biometric in the area of pattern recognition and has always been under study through its important role in forensic science that could help government criminal justice community. In this paper, a novel multi-biometric identification of individuals by means of both physiological and behavioural biometrics is introduced. Different from the most conventional biometric identification, the extracted physiological and behavioural biometrics will go through a proposed Multi-Biometric Feature Discretization. The intention of Discretization in this study is to prevail over the deficiencies caused by poor features mining and to attain individual unique features that could reflect the individual varianceness in order to discriminate one person from another. The experiments was conducted on real-world multiple form of images which is adopted from INF/ENIT, HITMW and FVC databases. The experimental results supported our analysis by demonstrating a remarkable potential of the new structure of multi-biometric identification adaptability on multi forms of image structures and has excellent potency to conserve the distinctiveness of individual during identification.

**Keywords:** Pattern Recognition, Machine learning, Data Mining, Identification, Multi-Biometrics, Feature extraction, Discretization

## **1** Introduction

Data mining and machine learning both are an established field of extracting potentially useful knowledge from huge database including the process of identifying the comprehensible patterns in data [1,2] and has become a research

focus recently, especially in industrial applications [3-5]. In this paper, our main work is on biometric datasets, thus our discussion will be more on biometric datasets of data mining tasks which often involve continuous huge attributes. Thus, a discretization is introduced in this study to discretize those continuous input features. Discrete features compared to continuous features are easier to describe, understand because the datasets are well represented and simplified. This quality discretization of discrete attributes is an important criterion that ensures high performance on speed, accuracy and understandability of the classification models instead of continuous attributes [6]. In addition to this, discrete representation more to knowledge level concept which is practically useful for most of the pattern recognition applications. We believe that data mining algorithm using the discretized datasets can significantly impact the performance of individual identification used in the analysis of high dimensional based biometrical data.

A biometric identification is one of the established system that uses the concept of pattern recognition technology. Pattern recognition is a broad term which cover three common types of application fields; identification [7], verification [8] and recognition [9]. In order to serve for both academia and industry including potential applications, from security, forensics to archaeology activities, the scope of pattern recognition systems has been broaden up to a biometric based system. This includes identification and verification based on signature [10,11], iris [12,13], facial features [14,15], fingerprints [16] and handwriting [7,8].

## 2 Analysis of Behavioral and Physiological Biometrics

Biometric term can be categorized into two parts: behavioral biometrics that require individual characteristics of a person's behaviour for identification process (e.g. handwriting, signature, speech, gait) and physiological biometrics that gauge a physical features of the human body to perform identification (e.g. fingerprint, iris, face, DNA). In this paper, the proposed multi-biometric identification pertains to both categories of biometric properties; physiological and behavioural biometrics. The behavioral identification is based on Arabic and Chinese handwriting. They are discussed in sub section 4.1. As for physiological identification, our focus is on fingerprint; an overview of fingerprint structure can be read in sub section 4.2.

#### 2.1 Arabic and Chinese handwriting

Handwriting is an everyday form of communication for human being and commonly treated as one of the behavioral biometric which play a significant role in biometric system, a fundamentally pattern recognition system that acquire biometric data from an individual, extracting a feature set from the data or images, comparing the feature set against the template set in database and finally make decision on the pattern of feature set. In the present, most of research challenges and techniques in handwriting analysis are based on soft computing methods as described in [21] which idea on the fact that handwriting identification starts with discovery the basic features that are gathered together in order to identify the individuality of writer. Arabic, the second Semitic alphabetic writing system most widely used in the world besides Latin alphabet (the most widespread). Unlike Latin writing system, the writing direction of Arabic is performed from right to left. The most noticeable characteristics of the Arabic writing is that Arabic scripts are intrinsically cursive and Arabic alphabet composes of 28 basic letters which consists of strokes and dots (one, two, or three). Thirteen of Arabic characters don't have dots, ten of them have one dot, three have two dots, and two have three dots. Dots can be above ( $\dot{z}$ ,  $\dot{z}$ ), below ( $\dot{z}$ ,  $\dot{z}$ ), or in the middle ( $_{7}$ ) of the letter [31]. These dots and other diacritical marks intentionally for character vowelisation create ligatures that are difficult to segment during pre-processing [31].

Besides Arabic, complication structure also applied in Chinese character. Chinese characters are ideographic in nature with more than 50,000 characters, of which 6000 characters are commonly used and have a wide range of complexity [32,33]. According to Fang, the complication structure in Chinese character mostly affected by multi stoke of each character. The characters may consist of one to thirty or more distinct strokes due to the variety of handwriting style. Fig. 1 shows the same Chinese character with different writing styles which has created a different number of strokes called multi strokes. Fang's proposed algorithms successfully unite some sub strokes into the proper and complete stroke to determine a Chinese character.



Fig.1 Same Chinese character with different number of strokes

#### 2.2 Fingerprint

A fingerprint pattern is formed of ridges and valleys over the surface of finger (Fig. 2). Different shape and structure of ridge and valley in each finger of an individual contributes to different global and local analysis. A global analysis is use to extract the common singular points namely loop, delta, whorl and etc. Fig. 3 shows examples of singular points such as core point, delta, loop and whorl. The core points or small circles shown in Fig. 3 is the center point of the highest loop in the singular region and generally use as a features to be pre aligned with other fingerprint pattern during matching process. These singular regions may be

classified into five classes: left loop, right loop, whorl, arch and tented arch as illustrated in Fig. 4.



Fig.2 Ridges and valleys on a fingerprint image



Fig.3 Singular regions and core points



**Fig.4 Fingerprint classes** 

In summary, global analysis provides an overall picture of fingerprint classification. Local analysis, on the other hand, gives more detail information of the image. These significant information including observe the regions of the ridge termination or ridge anomalies termed minutiae points. Some of typical ridge discontinuous structures are shown in Fig. 5. The ridges that suddenly break or

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discontinuous at the end are called termination. A terminate ridge with two split ridges at the end is called bifurcations. Fig. 5 shows a closer observation of real fingerprint image. We can see a ridge ending in Fig. 5 (a), a ridge bifurcation in Fig. 5(b). Fig. 5 (c) shows a lake. More details on different types of minutiae can be seen in Table 1.



Fig.5 Types of Minutiae (a) ridge termination or ridge ending (b) ridge bifurcation and (c) a lake

Minutiae	Types
	Termination
	Bifurcation
	Lake
	Independent Ridge
	Dot or Island
	Spur
	Crossover

**Table 1. Types of Minutiae** 

Overall, behavioural biometrics discussed here (Arabic and Chinese handwriting) can be expressed in variety of style, which is commonly based on the alignment and formation of multi directional, dots and stokes pattern. Naturally, the style of Arabic and Chinese character can be seen differently based on static, dynamic [34] and geometrical features. Static or stationary features are essential and it tells apart each character from other character. Dynamic features on the other hand represent generative aspect of the characters. Slant, ornamentals, aspect ratio, relative position and size of the strokes, corners and retraces included in geometrical structures. Because of these numerous variation of dots and stroke construction to write a certain character, the number of writing style of a character even by same person can be perpetuity. These features are those important points to be thought of when dealing with determining a writer of handwriting sample from a set of writers. The main challenge for behavioural identification is when dealing with different style of handwriting performed by the same author and how actually to acquire features that can reflect the authorship of the author. Same goes to the physiological biometrics identification as discussed in previous section. Table 2 summarizes the characteristics of behavioral and physiological biometric images.

	Table 2. Characteristics of Denvioral and Thysological biointerie images.							
<b>Biometric images</b>	Natural characteristic of Biometric images							
		1						
. ~	<ul> <li>Dots and other</li> </ul>	Cursive nature of						
(3 a) AW	diacritical marks	characters itself leads						
0 /- /	intentionally for	to variability between						
	character vowelisation	curve angles, shape						
<b>x</b> - <b>x</b>	create ligatures that is	and size						
السرايح	difficult to segment							
	during pre processing	(Those patterns vary greatly						
	[31]	among different writers and						
ou Di		even for same writer)						
• •								
DR• IFN/FNIT		Higher similarities						
	Multiple strokes	among features						
7 7	structures [35]							
1217	structures [55]							
	Difficult to determine							
112	various unique features							
4	of individuality							
I	or marriadancy							
<b>DB: HIT/MW</b>								

Table 2. Characteristics of Behavioral and Physiological biometric images.

DB: FVC	• Formed of ridges and valleys over the surface of finger	• Two general approaches: minutiae based [36] and correlation based approach; encounter overall patterns of ridges and valleys including distort regions found in the
2000,2002,2004		ridges

# **3 Problem in existing methods**

### 3.1 Arabic and Chinese Identification

It is generally accepted that most of handwritten Arabic and Chinese identification is a difficult problem especially due inherently to the cursive nature of Arabic characters itself, which leads to a lot of variability between curve angles, shape and size though in this paper we are not focusing on this issue. However, those features still considered as one of the identification complications to identify the individuality for same writer. To the best of our knowledge and assumptions, we conclude a few common categories of difficulties in a writer identification for Arabic and Chinese handwriting.

First, the Arabic writing styles include the shapes of the descenders and even on Arabic calligraphic styles can be performed more than a dozen. Table 3 summarizes the Arabic characters in isolated, initial, medical and final form. Isolated Arabic characters could be performed into other words when those characters are joined from right with other characters [37]. Those patterns vary greatly among different writers and even for same writer. Second, Arabic characters composes of many dots and other diacritical marks intentionally for character vowelisation create ligatures that is difficult to segment during pre-processing [38-41]. Third, the Arabic can be splits into subwords, also known as Parts of Arabic Words (PAWs) and PAWs create many problems [42]. Finally the quality of the scanned handwritten images in particular for highly degraded historical images in some real world applications could also lead to specific problems.

No	Letter Name <sup>a</sup>	Isolated Form	Initial Form	Medial Form	Final Form	No	Letter Name	Isolated Form	Initial Form	Medial Form	Final Form
1	Alef <sup>b, c</sup>	ی ا	-	-	ال	16	Tah	ط	ط	ط	ط
2	Beh	ٻ	ł	÷	ب	17	Zah	ظ	ظ	ظ	ظ
3	Teh <sup>d</sup>	ة ت	ï	ï	ة ت	18	Ain	3	2	۶	ع
4	Theh	ث	ĵ	â	ث	19	Ghain	ġ	غ	ė	غ
5	Jeem	ج	Ş	Ę	ج	20	Feh	ف	ۈ	ف	ف
6	Hah	5	>	ג	5	21	Qaf	ق	ۊ	ä	ق
7	Khah	Ś	ż	ż	خ	22	Kaf	ك	٢	ک	ىك
8	Dal <sup>b</sup>	د	-	-	د	23	Lam	J	J	T	٦
9	Thal <sup>b</sup>	ذ	-	-	ذ	24	Meem	م	۵	۵	م
10	Reh <sup>b</sup>	ر	-	-	ر	25	Noon	ن	i	i	ن
11	Zain <sup>b</sup>	ز	-	-	ز	26	Heh	٥	ھ	<del>6</del>	٩
12	Seen	س	ш	ىبىر	س	27	Waw <sup>b</sup>	و	-	-	و
13	Sheen	ش	ش	ش	ݾ	28	Yeh	ي	ż	¥	ي
14	Sad	ص	ص	۵	ص	29	Hamza <sup>e</sup>	2	ţ	Å	Ĺ
15	Dad	ض	ض	ض	ض						

 Table 3. Four forms of Arabic characters [37]

Besides Arabic, complication of Chinese character is due to the alignment and formation of multi stokes pattern. Chinese characters are well known as large alphabet language; it has a variety of categories. Numerous variation of stroke construction to write a certain Chinese character, the number of writing style of a Chinese character can be perpetuity even from same writer.

Natural characteristic of Arabic and Chinese writing itself, which surrounded by multifaceted mechanism and various complications leads to many problems unsolve and most of the identification system tends to be not effective enough. Identification based on cursive handwriting is very difficult and the overall issues can be summarized as follows. First, difficult to obtain discriminant handwriting features as handwritings are usually not stable enough and have vast variability, so this creates a large amount of features data consequently slow down the computation progression and misclassification often occurs during learning and classification. Misclassification often happen causes by overlapping feature space. Second, too much of pre-processing at early stage such as normalisation to discard mixes noise of the character in order to refine classification would almost certainly trade off important individualistic of the writer [43]. This creates a poor classification which leads to a low identification rate either in terms of its capability and competency. Third, real world classifications often involve with continuous features. Unfortunately most of machine learning and data mining method such as Bayesian Network could not perform well with continuous features unless the continuous features are first pre-processed. This process is called Discretization, which will be discussed in more detail in next section. Moreover, in real world writer identification problems the number of writers to be identified is often unlimited and each writer has only few reference handwriting samples, extremely, only one reference handwriting sample served as training samples.

#### **3.2 Fingerprint Identification**

From an extensive research of available literature, it was known that the popular fingerprint approaches are classified into two general classes: minutiae based approach and correlation based approach. Both approaches have some of its own shortcomings. The minutiae-based techniques first looks for the minutiae points and then match their chosen regions in a query finger with the stored patterns in database. Different fingerprint images have different number of minutiae points. The best total number of minutiae points for a good quality of fingerprint is around 60 to 80 minutiae points [44]. A disadvantage of minutiae-based technique is that it does not mend well to indexing mechanism.

Second approach called Correlation-based techniques [45,46], which consider the overall patterns of ridges and valleys found in a finger. These approaches encounter all information including distort regions found in the ridges. Thus, these ideas inferred to a noise problem as encountered in signal or image processing applications.

With regard to the fingerprint matching disadvantages (minutiae-based ones and correlation based approach), a great number of solutions have been proposed such as employing normalized cross correlation technique which can reduces the computational effort and decreases the error rate better than the minutiae based approach [47]. Meanwhile, in [48], they proposed a method that uses ridges joint with minutia. With this technique, the template cand query fingerprints can be easily aligned.

# 4 Related Work

Currently, many researchers have done great work enhancing feature extraction and classification methods as mentioned involves handwritten processing including normalization [59-60], feature extraction [61-62], and classification techniques based on soft computing methods including neural network [63] and rough set approaches. Only a few works focuses on data analysis and data representation. Besides understand and able to extract useful object's property at pre-processing phase, filter and intelligently discriminate unique features to represent data in a comprehensible way is important because this may intensify the individuality performance of identification process.

As such, we enlarge the scope of the possible use of the common biometric identification which has been, up to now, mainly evaluated on handwritings into biometrics identification coupled with the proposed Multi-Scores based Discretization which can work on any type of biometrics property in order to produce a wide variety of potential applications, from security, forensics and financial activities to archaeology. Framework of the Multi-Biometric Identification based Discretization as illustrated in Fig.5.

**Behavior and Physical Biometrics** 



4.1 Fig.5 Multi-Biometrics Identification based Discretization

It is generally accepted that most of handwritten Arabic and Chinese identification is a difficult problem especially due inherently to the cursive nature of Arabic characters itself, which leads to a lot of variability between curve angles, shape and size though in this paper we are not focusing

To achieve the objective of this study, previous feature extraction and discretization methods which is not restricted to any particular types of biometric behaviour and physiological property are enhanced. The concept of Discretization for data representation is adopted and the algorithm is based on Invariant Discretization originally proposed by Azah [64]. This technique has been applied to variety of handwriting types including our previous work on English handwriting

[64] and Chinese handwriting [21]. For this research, we improve Azah's Discretization to fit for multi-biometric images so-called Multi-Scores Discretization (MSD). The improved algorithm is able to deal with large and high dimensional feature vectors as well as to produce significant individual features which directly unique to those individual. Again, the improved Discretization here does not need the use of Normalization process.

In biometric identification, the proposed Multi-Biometric Feature Discretization (MBFD) transforms the multiple continuous feature set of individual closer to discrete feature sets (knowledge level representation). The description in two main steps below explains how the actual feature sets of an individual are discretized in multi-biometric for personal identification.

Step 1: Compute the size of interval,  $I_{width}$ . Given a set of features, the discretization algorithm first computes the size of interval,  $I_{width}$  as defined in Equation (4), i.e., it determines its upper and lower bounds. The range is divided by the number of features which then gives each interval upper approximation,  $AV_{upper}$  and lower approximation,  $AV_{lower}$  as demonstrated in Fig.6.

$$I_{width} = \frac{(f_{max} - f_{min})}{No_{feature}}$$
(1)



Fig.6 The improved Discretization; Multi-Biometric Feature Discretization.

The number of intervals generated is equal to the dimensionality of the feature vectors. This is to maintain the original number of extracted features from multiple extraction methods.

Step 2: Compute a single discriminatory value, *disFeature* of each interval. Instead of taking the range between the interval, the *disFeature* here is considered by taking the midpoint of the  $AV_{upper}$  and  $AV_{lower}$  interval as defined in Equation (2) (improvised from Azah's Invariant Discretization).

$$disFeature = \left(\frac{AV_{lower} + AV_{upper}}{2}\right) \tag{2}$$

With this improved procedure in computing intervals, the estimated representation feature values are more close to the actual multi-biometric features distribution (true feature values). This preserves the discriminative power of the original multi-biometric features and enhances the statistical distinctiveness between individuals. Therefore, here, the discretization is said to be robust and efficient enough to handle the issues in feature distribution of Multi-biometric identification. The graphical representation of the improved Discretization compare to normalization and previous Discretization is shown in Fig.7.



**Fig.7** Comparison of the actual, discretized multi-biometric features (proposed), previous discretized and normalized feature distributions on a fingerprint dataset.

Unlike other discretization approaches, the main motivation behind this algorithm is to maximize inter-class distances for all biometric samples that do not belong to the same individual class. By representing the features into a set of intervals, the issues of dimensionality caused by overlapping features can be avoided. This makes the identification process easier and faster due to the easier clarification by the classification and decision tasks. Fig.8 and Fig.9 each presented the samples of pre-discretized datasets for Arabic and Fingerprint, which is the actual extracted datasets before executed with the proposed Multi-Scores based Discretization algorithm. There are nine extracted vectors in each row and the last column represents the participant's class.

```
\begin{array}{c} 0.0027\ 0.1198\ 0.0306\ 0.4324\ 0.6136\ 0.9662\ 0.8006\ 0.0796\ 0.1145\ 1\\ 0.0148\ 0.0099\ 0.0142\ 0.8289\ 0.0562\ 0.0314\ 0.2361\ 0.0233\ 0.1537\ 3\\ 0.0075\ 0.0837\ 0.0062\ 0.5652\ 0.7277\ 0.5844\ 0.0149\ 0.0142\ 0.8720\ 2\\ 0.0029\ 0.0367\ 0.0082\ 0.5854\ 0.0838\ 0.2044\ 0.0371\ 0.2278\ 0.0659\ 1\\ 0.0106\ 0.0530\ 0.0075\ 0.8783\ 0.1322\ 0.0292\ 1.7893\ 0.1524\ 0.0629\ 3\\ 0.0030\ 0.1742\ 0.5573\ 0.0438\ 0.0373\ 0.5001\ 0.1037\ 0.2453\ 0.1610\ 1\\ 0.0135\ 0.0693\ 0.0538\ 0.1839\ 0.0404\ 1.4319\ 0.0190\ 0.1452\ 0.0153\ 3\\ 0.0172\ 0.0670\ -1.0000\ 0.0353\ 0.1841\ 0.0205\ 0.1077\ 0.0416\ 0.3497\ 2\\ 0.0017\ 0.0766\ 0.0562\ 0.2755\ 0.0328\ 0.0850\ 0.1933\ 0.2309\ 0.0028\ 1\\ 0.0020\ 0.0120\ -1.0000\ 0.0080\ 0.0305\ 0.0099\ 0.0187\ 2.0052\ 0.0939\ 2\\ 0.0317\ 0.2651\ 0.0122\ 0.7570\ 0.6109\ 0.7127\ 2.3325\ 0.1340\ 0.0580\ 3\\ 0.0045\ 0.0111\ 0.0063\ 0.0402\ 0.0142\ 0.3099\ 0.4702\ 0.8670\ 0.0121\ 2\\ \end{array}
```

Fig.8 Samples of Pre-discretized Arabic handwriting written by 3 writers

3.9500	7.0000	6.0000	9.1500	12.7500	16.9500	1.8500	5.8500	10.6000	1
1.9500	7.2000	5.6500	8.0000	14.4000	8.6500	3.4500	7.0000	3.5500	2
11.7000	15.8500	7.5500	14.9000	21.7500	13.8000	8.4500	15.0500	14.7000	3
1.6500	2.5500	2.2500	6.3500	9.0000	5.8000	3.5000	3.5500	9.1000	1
4.9500	7.6000	5.4500	12.1000	17.6000	8.6500	3.8000	7.1000	4.3500	2
3.5500	2.5500	1.2500	8.2000	11.5500	6.7500	4.3000	5.6000	7.3500	1
5.8000	6.7500	6.2000	16.2000	22.3000	15.4500	6.5000	9.7000	3.5000	2
9.8500	16.0500	6.7000	8.5500	7.9000	8.5500	3.9500	5.7000	5.2500	3
4.9500	8.8500	5.5500	13.0000	17.8500	11.5500	4.3500	6.2500	6.7000	2
13.5000	34.9500	22.3500	19.2500	31.7000	27.9000	9.7500	21.1500	17.3000	3
6.1000	8.0000	9.5000	14.0000	18.8000	17.0000	4.5500	7.5000	10.4500	1
4.1000	10.6500	8.1500	19.4500	31.9000	21.8000	10.9000	19.5500	23.5500	3

Fig.9 Samples of Pre-discretized Fingerprint from 3 participants

Fig.10 and Fig.21 show the discretized datasets for Arabic handwriting and fingerprint after process. Discretized datasets are also known as Post-Discretized scores. The circled vectors illustrate the *unique values* obtained from each individual. From Fig. 20, the different colours of circled vectors indicate that each writer has it own style of handwriting and this is demonstrated by its *unique values* here used as a discriminative role to portray the uniqueness features of individuality from others.

Biometric Identification based Scores

0.0535833 0.16245 0.0535833 0.48395 0.591117 0.80545 0.0535833 0.16245 1  $0.0535833 \ 0.0535833 \ 0.0535833 \ 0.591117 \ 0.0535833 \ 0.16245 \ 0.0535833 \ 0.269617 \ 0.0535833 \ 1$  $0.0535833 \ 0.16245 \ 0.591117 \ 0.0535833 \ 0.0535833 \ 0.48395 \ 0.0535833 \ 0.269617 \ 0.16245 \ 1$ 0.0535833 0.0535833 0.0535833 0.269617 0.0535833 0.0535833 0.16245 0.269617 0.0535833 1 0.129167 0.129167 0.129167 0.911667 0.129167 0.129167 0.129167 0.129167 0.129167 3 0.129167 0.129167 0.129167 0.911667 0.129167 0.129167 1.68667 0.129167 0.129167 3 0.129167 0.129167 0.129167 0.129167 0.129167 1.42833 0.129167 0.129167 0.129167 3 0.129167 0.129167 0.129167 0.653333 0.653333 0.653333 2.20333 0.129167 0.129167 3 0.168689 0.168689 0.168689 0.5026 0.836511 0.5026 0.168689 0.168689 0.836511 2 0.168689 0.168689 0.166956 0.168689 0.168689 0.168689 0.168689 0.168689 0.5026 2 0.168689 0.168689 0.166956 0.168689 0.168689 0.168689 0.168689 1.83824 0.168689 2 0.168689 0.168689 0.168689 0.168689 0.168689 0.168689 0.5026 0.836511 0.168689 2 ..... . . . . . . . . . . . . .

Fig.10 Samples of Post-discretized Arabic handwriting scores by 3 writers

Meanwhile, for fingerprint, different shape and structure of ridge and valley in each finger are the useful information to determine an individual from others. After going through the proposed Multi-Biometric Feature Discretization, the unique values of an individual are obtained as presented in Fig.11 below.

0.975 0.975 0.975 6.125	8.075 6.125	4.175 4.175 10.025	1	
4.175 6.125 6.125 10.023	5 11.975 17.825 (	0.975 6.125 10.025	1	
4.175 0.975 0.975 8.075	11.975 6.125	4.175 6.125 8.075	1	
6.125 8.075 10.025 13.92	5 17.825 4.175 8	8.075 10.025 8.075	1	
1.13056 7.60278 5.34167	7.60278 14.3861	7.60278 1.13056	7.60278 1.13056	2
5.34167 7.60278 5.34167	12.125 16.6472	7.60278 1.13056	7.60278 5.34167	2
5.34167 7.60278 5.34167	16.6472 21.1694	4 14.3861 7.60278	9.86389 1.13056	2
5.34167 9.86389 5.34167	12.125 18.9083	3 12.125 5.34167	5.34167 7.60278	2
12.5611 16.0056 16.0056	22.8944 12.5611	9.11667 9.11667	16.0056 16.0056	3
9.11667 16.0056 1.72222	9.11667 9.1166	7 9.11667 1.72222	1.72222 1.72222	3
12.5611 22.8944 19.45	33.2278 26.338	9 9.11667 19.45	16.0056 16.0056	3
1.72222 9.11667 19.45	33.2278 22.894	4 (9.11667) 12.5611	19.45 22.8944	3
•••••				

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Fig.11 Samples of Post-discretized fingerprint scores of 3 participants
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# 5 Experimental Results and Discussion

Several experiments were carried out to seek for the efficiency of the generic identification on both types of biometric images using the proposed MBFD in Biometric Identification. The identification performance is compared with the traditional one. To illustrate this, the experiments are done with different classification methods which are based on Rough set theory [65]. Four reduction algorithms based on rough set were chosen from ROSETTA Toolkit namely Johnson's algorithm, Holte IR algorithm, Genetic algorithm and Exhaustive algorithm. The experiments were conducted on different biometric properties of

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datasets: physiological and behavioural biometrics as summarized in Table 4 and Table 5 accordingly.

B	Images	ROSETTA Built-in	Data	70% Training Data	60% Training Data	50% Training Data	Average
i		Methods on	Types	30% Testing Data	40% Testing Data	50% Testing Data	Acc
0		Reductions					(%)
m							
t							
r							
i							
с							
		Johnson's Algorithm	Pre_Dis	11.11	8.33	6.89	8.78
	l		Post_Dis	94.44	95.83	93.10	94.46
	Chinese	Holte 1R Algorithm	Pre_Dis	11.11	12.50	10.34	11.32
	datasets		Post_Dis	100.00	100.00	100.00	100.00
		Genetic Algorithm	Pre_Dis	11.11	12.50	10.34	11.32
		T 1 2 41 21	Post_Dis	100.00	100.00	100.00	100.00
тт		Exhaustive Algorithm	Pre_Dis	11.11	12.50	10.34	11.32
н	HII/MW		Post_Dis	100.00	100.00	100.00	100.00
a n							
d							
w							
r							
i							
t							
i							
n a							
g		Johnson's Algorithm	Dra Dia	66 67	75.00	60.00	67.22
	Arabic	Johnson's Argorium	Post Dis	100.00	100.00	100.00	100.00
	datasets	Holte 1R Algorithm	Pre Dis	66.67	75.00	60.00	67.22
	uatasets	fione fit ingointin	Post_Dis	100.00	100.00	100.00	100.00
		Genetic Algorithm	Pre_Dis	66.67	75.00	60.00	67.22
		, and the second s	Post_Dis	100.00	100.00	100.00	100.00
		Exhaustive Algorithm	Pre_Dis	66.67	75.00	60.00	67.22
	IFN/ENIT		Post_Dis	100.00	100.00	100.00	100.00
		Johnson's Algorithm	Pre_Dis	67.22	26.11	32.78	42.04
	Fingerprint		Post_Dis	100.00	100.00	58.89	86.30
	datasets	Holte 1R Algorithm	Pre_Dis	73.89	26.11	32.78	44.26
F			Post_Dis	100.00	100.00	100.00	100.00
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		Genetic Algorithm	Pre_Dis	73.89	26.11	32.78	44.26
			Post_Dis	100.00	100.00	100.00	100.00
	FUG	Exhaustive Algorithm	Pre_Dis	73.89	26.11	32.78	44.26
	FVC		Post_Dis	100.00	100.00	100.00	100.00

Table 4. Identification accuracy after reduction algorithm

From the above-obtained result in Table 4, it clearly shows that the first approach based on Rough set theory, four reduction methods namely Johnson's algorithm, Holte 1R, Genetic algorithm and Exhaustive algorithm perform well with post-discretized datasets. Each method successfully achieved the overall average accuracy of more than 90.0%. From the same result table, it can be seen that the same four algorithms presented the worst performance with pre-discretized datasets, which provide low identification rates; below 75.0%.

Clearer illustration and discussion are summarized in Table 5 below. The comparison with the proposed Multi-Biometric Identification based Discretization is presented. The proposed identification successfully achieves more than 95.0% on an average when compared to the existing system which obtain an average accuracy lower than 50.0% on every biometric datasets. This illustrates that the proposed architecture is much competent than existing system; can extract useful features in much better way during pre-processing phase and able to bear a higher discriminative power of individuality during post-processing phase. This can be seen through the significant improvement in the identification rates.

Classification	Biometric Datasets	Conventional Biometric Identification (%)	Novel Multi-Biometric Identification (%) (Proposed Multi-Biometric Feature Discretization)
Reduction algorithms based	Chinese Handwriting	10.68	98.61
on Rough set Theory:	Arabic Handwriting	67.22	100.00
Johnson's Algorithm	Fingerprint	43.70	96.57
Holte 1R Algorithm Genetic Algorithm Exhaustive Algorithm			
Overall Acc (%)		40.53	98.39

Table 5. Comparisson of Identification Accuracy on each FVC databases

On the whole, to quantify the performance of the proposed algorithm on Multi-Biometric identification in brief, a *t-test* is conducted to illustrate the significance difference between the proposed and the existing one. For hypothesis testing, the scalar mean of an average accuracy for pre and post discretized scores after several training and testing procedure is used as a statistic and basis for one-to-one individual identification performance comparisons of the three biometric datasets.

Given two paired sets; pre-discretize datasets,  $\mathcal{X}_i$  and post discretized of each selected biometric datasets,  $\mathcal{Y}_i$  of *n* total datasets for each biometric databases. Examples, if the selected biometric datasets is fingerprint. Then, the given two paired sets would be pre-discretize of fingerprint datasets,  $\mathcal{X}_i$  and post

discretized of fingerprint datasets,  $\mathcal{Y}_i$  of *n* total fingerprint for each FVC database, the paired *t*-*test* determines if they differ from each other in a significant way. The t value is computed using the mathematical equation (1), (2) and (3) respectively.

$$\hat{x}_{i} = (x_i - \overline{x}) \tag{1}$$

$$\hat{y}_{i} = (y_i - \overline{y})$$
(2)

$$_{t=} (\bar{x} - \bar{y}) \sqrt{\frac{n(n-1)}{\sum_{i=1}^{n} (\hat{x}_{i} - \hat{y}_{i})^{2}}}$$
(3)

 Table 6. Comparison of Identification Accuracy on each biometric image

Biometric Databases			Mean	Std. Dev.	Τ	Р	Significance
Handwriting	IFN/ENIT (Arabic)	Pre Post	67.22 100.00	6.41 0.00	9.77234E-10	0.05	Highly Significance
	HIT/MW (Chinese)	Pre Post	10.68 98.61	1.68 2.57	2.22631E-20	0.05	Highly Significance
	FVC2000	Pre Post	72.22 100.00	6.64 0.00	8.19455E-09	0.05	Highly Significance
Fingerprint	FVC2002	Pre Post	26.11 93.06	5.74 16.60	3.71138E-08	0.05	Highly Significance
	FVC2004	Pre Post	32.78 89.72	6.41 18.93	3.49287E-07	0.05	Highly Significance

As per Table 6, when *t* test analysis was done on post-discretized biometric scores and those of actual biometric datasets (pre- discretized datasets), the results were highly significant with *t* 8.19455E-09 and P<0.05 for fingerprint datasets. When *t* test analysis was done on IFN/ENIT Arabic datasets, the results were also highly significant with *t* 9.77234E-10 and P<0.05. Same analysis achievement on pre and post-discretized HIT/MW Chinese datasets which shown a highly significant with *t* 2.22631E-20 and P<0.05, stating that individual identification shows a better performance with discretized datasets as compared to the actual datasets. Thus, the analysis of the data reveals that there is a definite improvement in the individual identification performance after enhancing the previous biometric identification to overcome the discrepancy caused by poor generalization and able to extract useful information from any form structure of biometric property correctly. Moreover, the concept of discretization in biometric is able to achieve lower identification error than those one without the use of discretization.

# 6 Conclusion

In this paper we presented a study of a complex characteristics of biometric images adopted from three standard databases INF/ENIT, HITMW and FVC databases. The process of feature extraction is presented along with the results. Analysis of the usefulness features was conducted via searching and extraction of the best feature sets using the morphological operations and 3x3 zoned windows method. Beside this, in the most of the individuality analysis of handwriting studied, Azah's discretization method does possess good discriminative power for handwriting identification. This successfully justified their use in identification analysis. Thus, in this study, we enhance and propose a novel discretization to fit for Multi-Biometric identification so called Multi-Biometric Feature Discretization (MBFD). The algorithm handle multiple behaviour and physiological images by search, induce, organize and represent a list of intervals into a better individual informative representation. This algorithm was tested on a few classification methods to seek for its effectiveness on multiple biometric images. The biometric identification based on proposed MBFD extensively boosted up the individual identification accuracy of 99.0% on any form of biometric images and effectively prevails over the existing identification that had positional and oriental weaknesses.

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