

Jian Zhou<sup>1,2</sup>,  
Jun Wang<sup>1\*</sup>,  
Honggang Bu<sup>3</sup>

# Fabric Defect Detection Using a Hybrid and Complementary Fractal Feature Vector and FCM-based Novelty Detector

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<sup>1</sup>Ministry of Education,  
Key Laboratory of Textile Science & Technology  
Donghua University,  
Shanghai, 201620, China  
\*E-mail: junwang@dhu.edu.cn

<sup>2</sup>Ministry of Education,  
Jiangnan University,  
Key Laboratory of Eco-textiles,  
Wuxi, Jiangsu 214122, China

<sup>3</sup>North Valley Aircraft,  
11647 33rd St SE, Valley City, ND, 58072, USA

## Abstract

*Automated defect detection in woven fabrics for quality control is still a challenging novelty detection problem. This work presents five novel fractal features based on the box-counting dimension to address the novelty detection of fabric defect. Making use of the formation of woven fabric, the fractal features are extracted in a one-dimension series obtained by projecting a fabric image along the warp and weft directions, where their complementarity in discriminating defects is taken into account. Furthermore a new novelty detector based on fuzzy c-means (FCM) is devised to deal with one-class classification of the features extracted. Finally, by jointly applying the features proposed and the FCM based novelty detector, we evaluate the method proposed for eight datasets with different defects and textures, where satisfying results are achieved with a low overall missing detection rate.*

**Key words:** defect detection, box-counting dimension, fuzzy c-means, novelty detection.

## Introduction

In modern weaving mills, performing a visual examination of fabric products is a standard process before reaching customers, as defects can severely affect products' visual quality. Currently visual inspection is still manually performed by well-trained human inspectors, a job whose efficiency is greatly limited by the human inspector's experience. For this reason, the automation of fabric inspection based on computer vision and emerging artificial intelligence techniques have drawn considerable attention, and numerous approaches have been proposed to address this issue in recent decades [1].

Generally images of normal fabrics are dominated by the texture, which always exhibits high periodicity among sub-patterns. If there is a defect occurring in a fabric, the local regularity (periodicity) of the fabric will be disrupted, causing a local anomaly against its homogeneous texture background. Such anomalies always manifest in various ways, e.g. local structure or intensity change. Basically how to find versatile features that can robustly describe normal textures while sensitive to such anomalies (defects) is the basis for designing detection algorithms. Typical features used include the parameters of the textured model [2-3], features based on the grey-level co-occurrence matrix [4], Fourier features [5], multi-scale wavelet decomposition coefficients [6-7], and Gabor filtering fea-

tures [8]. With little taking into account of the complementarity of the features extracted, the aforementioned features or simple group of them generally result in performing well for certain types of fabric or defects and poorly for others.

Associated with texture roughness, the use of fractal based features such as the fractal dimension is another important approach for texture analysis. Fractal-based texture analysis was introduced by Pentland [9] in 1984, whose experimental results showed that there was a high positive correlation between texture roughness and fractal dimension, i.e. the rougher the texture, the larger the fractal dimension, with similar results found in [10-11]. Since the fractal dimension offers a decent measurement for the roughness of a texture surface, it is intuitive to use the fractal dimension as the feature for discriminating between a normal fabric texture and defective texture. Typical works include the following: Conci and Proença utilised the differential box-counting dimension for fabric defect detection [12], and Wen adopted another fractal feature, namely the Hurst coefficient, for identifying defects [13]. It is noted that the fractal features used in [12-13] are too few and simple, and only one fractal feature is used, which usually leads to a high false detection rate, e.g. up to 28% [12]. In fact, the single fractal feature, especially the fractal dimension, has a drawback in characterising local details of the texture, which might possess an identical or quite close fractal dimension for different textures. Some researchers were aware of the limitation of using a single feature, and different frac-

tal-based features were proposed in order to achieve better results. For example, Bu and Huang extracted four box-counting dimensions calculated from different measuring scale ranges [14]. In addition, the use of multi-fractals or feature combination can be seen in other applications [15-16].

In this paper, we introduce five new fractal features for fabric defect detection. The features proposed integrate local and globe texture information, which is able to describe the inherence of the fabric texture deeply. There are two distinctive differences between the fractal features proposed and previous features: (1) the features proposed make use of the woven fabric's horizontal and vertical details in projection operations; (2) the complementarity of the features extracted are taken into consideration in order to capture local and global details of the fabric texture. Besides this, all fractal features are extracted in a one-dimension series, rather than in two-dimension images. Considering fabric defect detection belongs to the category of one-classification or novelty detection problems, this paper also devises a new novelty detector based on fuzzy c-means clustering to handle the one-class or novelty classification of features extracted.

The rest of this paper is organised as follows: a brief introduction to the box-counting dimension and fuzzy c-means is firstly presented. The methodology appears in the next section, and the following is the experimental results and discussion. Conclusions are in the last section.

## Background

### Box-counting dimension

In fractal geometry, there are many methods that can be used to define the fractal dimension, such as the box-counting dimension (or box dimension), Hausdorff dimension and correlation dimension etc. In this work, the box-counting dimension, also known as the Minkowski dimension, is used to calculate the fractal dimension of a fabric image, as it can deal with non-self-similar fractals and be simply implemented with a computer. The key idea of the box-counting dimension is about covering with boxes, i.e. to count how many boxes are required to cover a set in different scales. Suppose that  $N(\delta)$  is the number of boxes of side length  $\delta$  required to cover a set  $S$ , the box-counting dimension is defined as:

$$D(S) = \lim_{\delta \rightarrow 0} \frac{\log N(\delta)}{\log(1/\delta)}. \quad (1)$$

Here the box dimensions are based on one-dimension sequences, and an example of a one-dimension curve is presented to demonstrate the procedures of calculating the box dimension. Let us denote the curve as  $S(x), x = 1, 2, \dots, M$  and  $N(\delta)$  is the minimal number of  $\delta \times \delta$  squares needed to cover  $S(x)$ . According to Equation (1), there is  $\log(N(\delta)) \propto D(S)\log(1/\delta)$ , and the  $D(S)$  can be estimated by the slope of the linear portion of  $\log(N(\delta))$  against  $\log(1/\delta)$ . Specifically let  $T$  be any sub-sequences of length  $\delta$  in  $S(x)$ , and then the minimal number of  $\delta \times \delta$  squares needed to cover  $S(x)$  is given by,

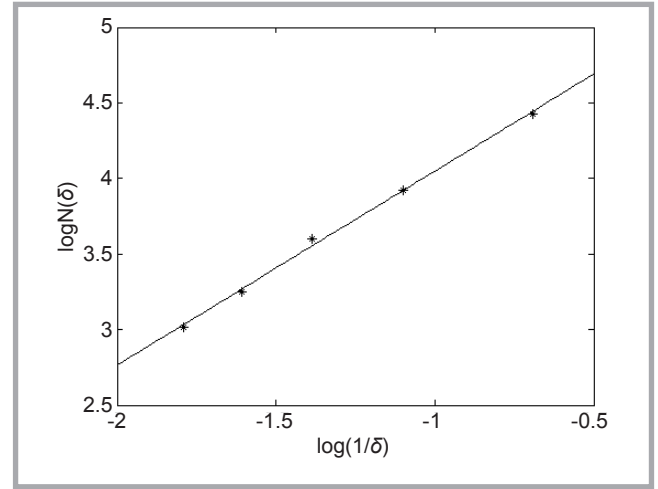
$$n(\delta) = \text{Int} \left[ \frac{\max(T) - \min(T)}{\delta} \right] + 1, \quad (2)$$

where,  $\text{Int}[\ ]$  is the integer rounding operator, and  $\max(T)$  and  $\min(T)$  are the largest and smallest values in the sequence  $T$ , respectively. Then  $N(\delta)$  can be obtained by calculating the mean of every sub-sequence in  $T$  with Equation (2), which is given by,

$$N(\delta) = \frac{E(n(\delta)) \times M}{\delta}, \quad (3)$$

where,  $E()$  is the expectation operator. Having obtained a different  $N(\delta)$  by changing the value of  $\delta$ , the slope of the linear portion of  $\log(N(\delta))$  against  $\log(1/\delta)$  can be estimated by means of fitting those points in the least-squares sense. Figure 1 depicts the fitting result of a data sequence of length 32 with  $\delta = 2, 3, 4, 5, 6$ , and the fitting equation is  $y = 1.28x + 5.33$ , i.e. the box dimension  $D(S)$  is 1.28.

Figure 1. Fitting result for calculating box dimension.



### Fuzzy c-means model

In machine learning, cluster analysis or clustering is one of the most important unsupervised learning methods of partitioning a set of unlabelled data into different groups according to a certain similarity. Different from supervised methods, the clustering techniques do not need any prior knowledge of the labels of data, potentially allowing to reveal the underlying structures within data. The clustering methods can be loosely classified into two categories: hard and soft clustering. In hard clustering, the data is grouped into hard-assigned clusters, where each data point completely belongs to one cluster, e.g. a k-means clustering algorithm. Rather than make a hard decision for each data point, soft clustering allows a data element to belong to more than one cluster, where the soft decision is made according to their degrees of belonging (quantified by the membership function). One of the most commonly used soft clustering is the fuzzy c-means (FCM) algorithm [17], which has been successfully applied to many applications, such as image segmentation [18] and fault diagnosis [19].

Given a set of interest  $\{x_i\}_{1 \leq i \leq N}$ , the FCM algorithm attempts to partition the data set into  $C$  different groups with cluster centres  $\{c_j\}_{1 \leq j \leq C}$ . Thus the objective function  $J_m$  for finding those centres is defined as:

$$J_m = \sum_{i=1}^N \sum_{j=1}^C u_{ij}^m \|x_i - c_j\|_2^2, \quad 1 \leq m < \infty \quad (4)$$

where,  $u_{ij}$  is the degree of belonging of  $x_i$  to the cluster  $j$ , and  $m \geq 1$  is the weight controlling the degree of fuzziness. Since the cluster centres  $c_j$  and degrees of belonging  $u_{ij}$  can be represented by each

other through the necessary conditions for minimising  $J_m$ , the alternating optimisation strategy can be used to minimise Equation (4). The implementation is composed of the following procedures: S1: Given  $C$  and  $m$ , initialise  $c_j^{(0)}$  with random numbers; S2: For  $t = 0, 1, \dots, n$ , update  $u_{ij}$  and  $c_j$  using the following formula:

$$\text{S3: Update } u_{ij}: u_{ij}^{(t)} = \frac{\left( \|x_i - c_j^{(t)}\|_2 \right)^{\frac{2}{m-1}}}{\sum_{k=1}^C \left( \|x_i - c_k^{(t)}\|_2 \right)^{\frac{2}{m-1}}}$$

$$\text{S4: Update } c_j: c_j^{(t+1)} = \frac{\sum_{i=1}^N (u_{ij}^{(t)})^m x_i}{\sum_{i=1}^N (u_{ij}^{(t)})^m}$$

S5: Repeat steps S3 and S4 until  $\|U^{(t+1)} - U^{(t)}\|_F < \epsilon$  is achieved, where  $U = [u_{ij}]$  is the membership matrix.

## Methodology

### Fractal feature extraction

Due to the fact that most real-world image textures do not exhibit stringent self-similarity, instead an approximate or statistical self-similarity, the fractal dimension calculated by the box-counting method does not possess the quality of scale invariance, which was also pointed out by Pruess [20], namely that limited observation resolution made the estimated box dimension vary with box scales. In our preliminary experiments, we noted that there was a positive association between the box dimension and box scales. Thus in order to alleviate the drawback of instability in estimating the box dimension with a fixed box scale, this work attempts to calculate the box dimension in a range of box scales, meanwhile addressing the

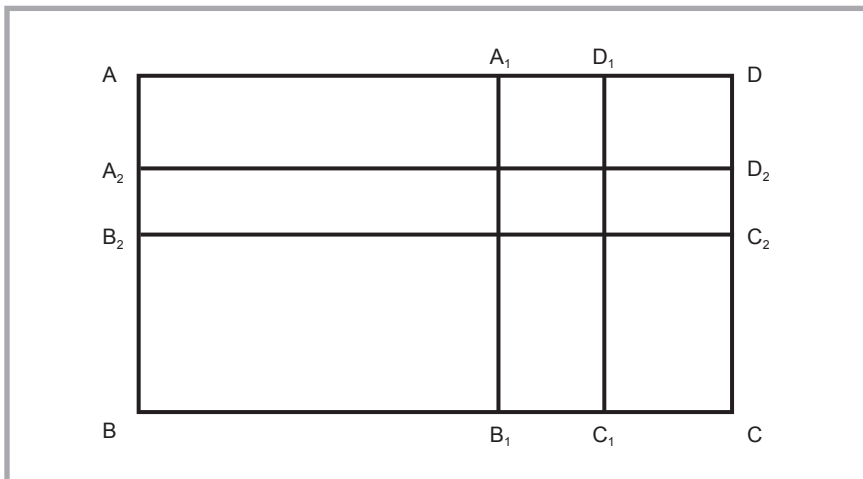


Figure 2. Illustration of feature extraction.

complementariness of the fractal features extracted. Moreover all the features proposed are extracted in one-dimension sequences by taking advantage of the formation of the fabric (interlaced with two mutually perpendicular yarn systems).

Without loss of generality, we illustrate the feature extraction scheme with a rectangle window ABCD, which is illustrated in Figure 2. Let us denote an image patch of size  $L_1 \times L_2$  ( $AB \times AD$ ) as  $W$ , a sub-window  $A_1B_1C_1D_1$  of size  $P_1 \times L_2$  as  $W_1$ , and a sub-window  $A_2B_2C_2D_2$  of size  $L_1 \times P_2$  as  $W_2$ . With the definition, the box dimension features are extracted based on  $W$ , given as follows:

- **Fractal feature 1:** Perform projection operation along AD and AB, i.e. compute the mean grey-level values along AD and AB, and denote the resultant two sequences as  $S_v$  and  $S_h$  with a length of  $L_1$  and  $L_2$ , respectively, combine  $S_v$  and  $S_h$  into a new sequence with a total length of  $L_1 + L_2$ , and calculate the box dimension of the new sequence using Equation (1), denoting fractal feature 1 as  $D_1$ .
- **Fractal features 2 and 3:** For  $W_1$ , perform a projection operation along  $A_1D_1$ , i.e. compute the mean grey-level values along  $A_1D_1$ , calculate the box dimension of the resultant sequence, transverse  $W_1$  through the whole window  $W$  along the horizontal direction, obtaining  $L_1 - P_1 + 1$  box dimensions, and extract the maximum and minimum box dimensions among them, denoting fractal features 2 and 3 as  $D_2$  and  $D_3$ .
- **Fractal features 4 and 5:** For  $W_2$ , perform a projection operation along the  $A_2B_2$  direction, i.e. compute the mean grey-level values along the  $A_2B_2$

direction, calculate the box dimension of the resultant sequence, transverse  $W_2$  through the whole window  $W$  along the vertical direction, obtaining  $L_2 - P_2 + 1$  box dimensions, and extract the maximum and minimum box dimension among them, denoting fractal features 4 and 5 as  $D_4$  and  $D_5$ .

Conveniently we combine the five fractal features extracted into a hybrid feature vector  $D = [D_1 D_2 D_3 D_4 D_5]$ . In particular,  $D_1$  mainly reflects the global information of fabric textures, which is more robust than the other four features, crucial for characterising the texture in most cases;  $D_2 \sim D_5$  are sensitive to local changes, capable of compensating for  $D_1$  in characterising details. Furthermore our preliminary experiments demonstrated that  $D_2$  and  $D_3$  are particularly sensitive to weft-wise defects, as the local abrupt of weft-wise defects can be well revealed; similarly,  $D_4$  and  $D_5$  are capable of finding the warp-wise defects. Note that all aforementioned features are calculated in a measuring scale range of 2~6 pixels.

#### Feasibility validation of features

In this section, we take two real-world fabric samples as examples to demonstrate the validation of the fractal features proposed. Here we hypothesise about the features from normal and defective samples obeying normal distribution, and the statistical hypothesis test, namely the t-test, is applied to objectively evaluate the effectiveness of the features proposed in differentiating between normal and defective fabric textures. It is notable that the t-test is not only able to judge the significance of the validation of a feature, but can tell the degree of effective-

ness of the various features, i.e. the larger a statistic, the more effective and discriminative the power. Suppose that there are two sets of feature samples,  $\{x_i\}$  and  $\{y_i\}$ , calculated from normal and defective samples, and the  $T$ -statistic to test whether the means of the two groups are statistically different from each other is given by,

$$T = \frac{\mu_1 - \mu_2}{\sqrt{\frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2}}} \quad (5)$$

where  $\mu_1$  and  $\mu_2$  are the means of samples  $\{x_i\}$  and  $\{y_i\}$ ,  $\sigma_1^2$  and  $\sigma_2^2$  the standard deviation samples  $\{x_i\}$  and  $\{y_i\}$ , and  $n_1$  &  $n_2$  are the total number of samples  $\{x_i\}$  and  $\{y_i\}$ . Here the  $T$ -statistic obeys the t-distribution with  $n_1 + n_2 - 2$  degrees of freedom and the significance level  $\alpha = 0.01$  is used. Thus if  $|T| > t_{0.99}$ , the feature is considered to be valid (significance).

**Example 1:** Figure 3 presents a typical coarse weft image, where Figure 3.a is a normal fabric image, Figure 3.b a defective image, and Figure 3.c has been divided into 64 patches of size  $32 \times 32$ . As shown in Figure 3.c, there are 8 patches containing defects in the last row. The total numbers of normal and defective samples are 64 and 8. Thus the associated statistics are computed as follows: The degree of freedom is equal to 70 ( $64 + 8 - 2$ ),  $t_{0.99} = 2.38$  (one-tailed), and the statistic  $|T|$  of the five fractal features  $D_1 \sim D_5$  are 6.99, 13.59, 10.77, 0.2125 and 8.03. Therefore, in this case, the four features (besides  $D_4$ ) are all valid with a high significance, especially for  $D_2$ .

**Example 2:** Figure 4 presents a typical double filling image, where Figure 4.a is a normal fabric, Figure 4.b a defective image, and Figure 4.c has been divided into 64 patches of size  $32 \times 32$ . As shown in Figure 4.c, there are 8 patches containing defects in the second last column. The total numbers of normal and defective samples are also 64 and 8. Thus the associated statistics are computed as follows: the degree of freedom is equal to 70 ( $64 + 8 - 2$ ),  $t_{0.99} = 2.38$  (one-tailed), and the statistic  $|T|$  of the five fractal features  $D_1 \sim D_5$  are 6.54, 1.84, 5.59, 12.93 and 9.03. Therefore, in this case, the four features (besides  $D_2$ ) are all valid with high significance, especially for feature  $D_4$ .

As demonstrated above, although the extracted features have different discriminative power for different defect types, the integration of them can ensure that at

least one feature is significant in defect detection practice.

### Construction of novelty detector based on FCM

As mentioned in the preceding section, fabric defect detection is a kind of novelty detection problem, rendering the conventional FCM model useless for straightforward defect detection, as the FCM is normally used for the data, whose number of clusters is specified. If one tries to cluster normal samples into a centre and cluster the defective samples into several centres with respect to defect types, then one must know whether the input data contain defects or how many defect types there are in it in advance, which is unknown in practice. On the other hand, if the input data do not contain defects, a substantial false detection rate would be achieved, where clustering is conducted on it anyway; even though one tries to cluster the potential defective sample into one centre.

In this paper, we train the FCM with only normal samples, and more than one clustering centre is utilised to describe the stochastic variations of normal samples. For the decision rule, given an unknown sample  $y$ , if the minimal Euclid distances of  $y$  to the clustering centres (obtained after training FCM) exceeds the pre-defined threshold, then  $y$  is a defective sample, otherwise a normal one.

**Parameter optimisation:** There are two parameters: the number of clustering centres  $C$  and the fuzzy weight  $m$ . For  $C$ , a common way is to define a statistic to measure the validity of clustering in order to find the optimum number of clusters. However, a related work remarked that the selection of the number of clusters was a task-specific problem, and it was not justified to use one validity measurement for all the cluster analysis due to the random distribution of data. Compared with different validity measurements, we found that the validity measurement function proposed by Gao was suited to the clustering of normal fabric samples [21]. Given the number of cluster centres  $C$ , the fuzzy weight  $m$  and the total number of training samples  $N$ , the validity measurement is defined by,

$$FP(C) = \frac{1}{N} \sum_{j=1}^C \left( \sum_{i=1}^C \mu_{ij}^m / \sum_{i=1}^C \mu_{ij} \right) - \frac{1}{C} \sum_{i=1}^C \left( \sum_{j=1}^N \mu_{ij}^m / \sum_{j=1}^N \mu_{ij} \right) \quad (6)$$

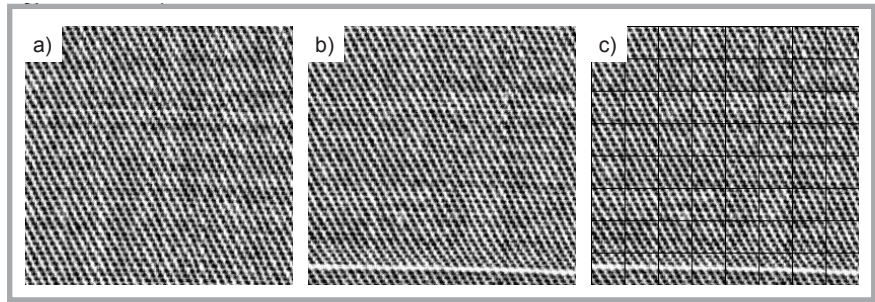


Figure 3. Fabric image samples: a) normal fabric image, b) defective image, c) illustration of patch division.

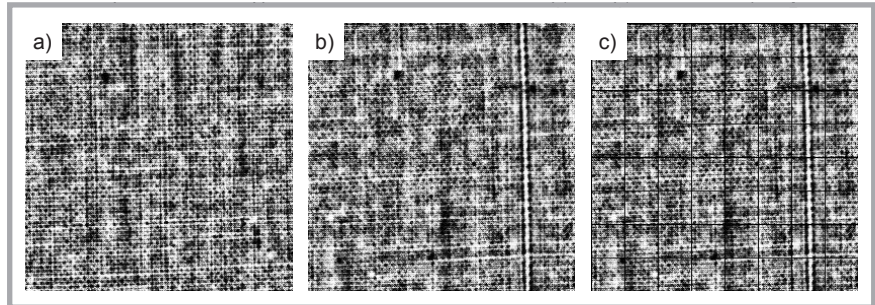


Figure 4. Fabric image samples: a) normal fabric image, b) defective image, c) illustration of patch division.

Thus, fixing  $m$  and calculating  $FP(C)$  with different  $C$ , the optimum  $C$  is the minimum one of set  $FP(C)$ , where the searching range of  $C$  is from 2 to  $2\ln(N)$  (a commonly used upper boundary).

In terms of  $m$ , there is no theoretical way of reference, and most researchers choose it empirically or experimentally. For example, Bezdek pointed out that the choice of  $m$  was an application specific problem, and suggested a range (1.1 5) for reference [17]; Pal and Bezdek found the range (1.5 2.5) is empirically best for most cases [22]. In this work, our primary experimental results showed that the suitable range of  $m$  for normal sample clustering is from 2 to 5. Considering the computing efficiency (a larger  $m$  needs less iteration times for the convergence of Equation (4)), here we select  $m$  empirically according to the total number of training samples as well, i.e. when the number of training is larger than or equal to 3000, setting  $m=4$ , otherwise setting  $m=2.5$ .

## Experiments

To evaluate the efficiency of the fractal features extracted and the FCM novelty detector described in the preceding sections, eight datasets are chosen for testing, including three twill fabrics and five plain fabrics. A detailed configuration

of the eight datasets is listed in Table 1, where the samples refer to sub-windows of size  $32 \times 32$  pixels. Besides this, necessary pre-processing operations are conducted to obtain comparable features for classification, and histogram equalisation is firstly adopted to help enhance the contrast of defects in the raw fabric images acquired before partitioning into sub-windows. Since the values of raw features vary widely, each fractal feature extracted is normalised to have zero mean unit variance, whose normalised version is given by,

$$x' = \frac{x - \bar{x}}{\sigma}, \quad (7)$$

Where,  $\bar{x}$  and  $\sigma$  are the mean and standard derivation of  $x$ .

Having calculated the five normalised fractal features, the FCM classifier proposed needs to be trained with a normal sample in order to find the optimal parameter for practical testing of unknown samples. Here, dataset 1 (see Table 1) is taken as an example to demonstrate the training process:

- S1: Choose  $m=4$  as the total number of training samples larger than 3000;
- S2: Calculate the search range of clusters  $C$ :  $2 \leq C \leq 16$  ( $2\ln(4472) = 16.8$ );
- S3: Calculate  $FP(C)$  in the range from 2 to 16, which is listed in Table 2.

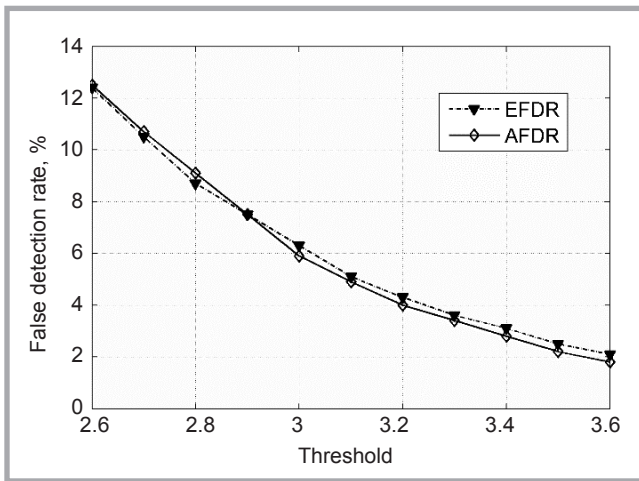


Figure 5. Trend curves of EFDR and AFDR under varying thresholds.

we adopt a simple and effective way to determine this parameter, which is to collect an additional sample set from normal samples as the validation set, and conduct a test on it by increasing the threshold with a small step to obtain its false detection rate (FDR) with respect to the threshold used. Then the FDR obtained can be used as an estimated false detection rate (EFDR) in favour of choosing a threshold in practical detection. Figure 5 presents the trend curves of EFDR and the actual false detection rate (AFDR), where EFDR and AFDR are obtained from training samples and testing samples of dataset 1.

From Table 2, it can be seen that  $FP(C)$  has the minimum when  $C=3$ . Thus the optimal clustering numbers for datasets 1 is 3 are  $(-0.0057, -0.0047, -0.0060, 0.0011, -0.0031)$ ,  $(0.4532, 0.4246, 0.4392, 0.1631, 0.1474)$  and  $(-0.4511, -0.4369, -0.4273, -0.1790, -0.1215)$ . Since the features extracted have been normalised with a zero mean, the features from normal samples should fluctuate around zero. That is to say, the first cluster centre is almost located at the

origin, and the other two clusterings are symmetrical with respect to the origin. Therefore with the three optimal cluster centres, unknown features will be classified into a defect if any of their distance to the three clusters exceeds the predefined threshold.

Generally the choice of threshold is a non-trivial task due to the absence of negative samples for the estimation of the missing detection rate (MDR). Here

As shown in Figure 5, the values of EFDR and AFDR are quite close to each other at a specific threshold, whose trend curves exhibit good agreement under different thresholds. It indicates that EFDR can provide a good prediction for the AFDR, i.e. it is reasonable to use it for the selection of the threshold in practical detection. On the other hand, since the actual MDR is unknown, the threshold selected referring to the EFDR can only make sense in controlling the AFDR in some manner, leaving no clues for actual MDR. Fortunately the FDR and MDR have a negative tendency towards each other, i.e. the larger the FDR, the smaller the MDR. Thus, from the application point of view, if a low MDR is important, it is recommended to choose a threshold with respect to a higher EFDR. Furthermore Table 3 summaries the detection results for all eight datasets.

By examining Table 3, it can be seen that: The number of clusters vary with the datasets, i.e. its optimal are specific to fabric textures. Explicitly the larger the variation of input features, the more clusters are required to describe their loose distribution, i.e. a texture with lower regularity would need more clusters than that of higher regularity. It can be observed from dataset 7 that there are 12 cluster centres for the fabric of low regularity (see Table 3). Moreover twill fabric always needs less centre than plain fabric, as it generally has higher regularity than plain ones.

The average EFDR and AFDR for all datasets are quite close to each other, indicating the robustness of the EFDR in forecasting the AFDR. The reasons for the poor agreement of datasets 4 and 7 are mainly due to the low texture regularity and too few training samples being used.

Table 1. Sample configuration of the eight datasets.

Dataset	Fabric type	Training samples	Testing	
			Normal samples	Defective samples
1	Twill	4472	4472	302
2	Twill	2960	2960	84
3	Twill	1792	1792	256
4	Plain	1820	1820	170
5	Plain	5010	5010	320
6	Plain	1608	1608	216
7	Plain	1168	1168	101
8	Plain	1474	1474	76

Table 2.  $FP(C)$  for different  $C$ .

C	2	3	4	5	6	7	8	9
FP	0.433	<b>-0.934</b>	0.078	-0.229	-0.786	-0.134	-0.387	-0.190
C	10	11	12	13	14	15	16	/
FP	-0.348	-0.288	-0.173	-0.258	-0.402	-0.320	-0.222	/

Table 3. Summary of detection results.

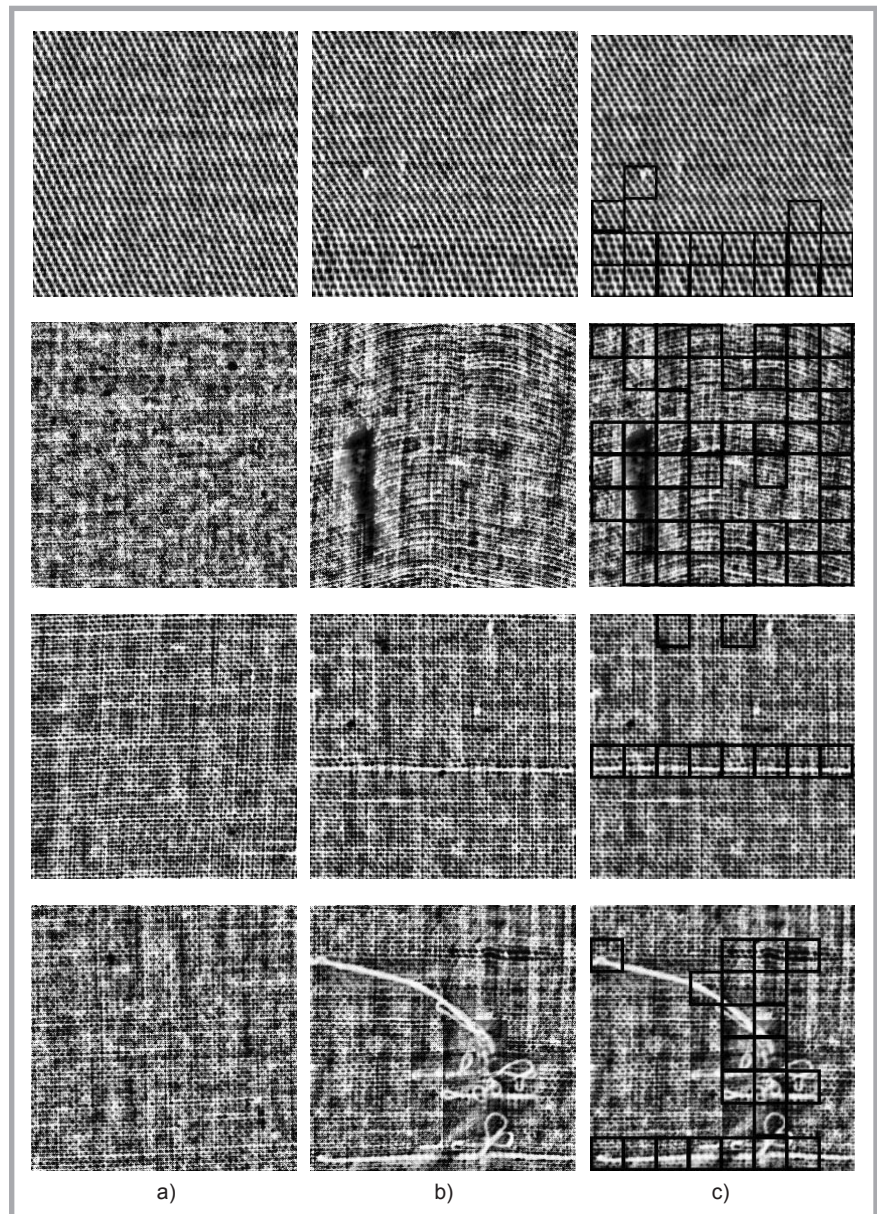
Dataset	C	EFDR,%	AFDR,%	MDR,%
1	3	8.12	8.23	5.63
2	4	2.97	3.17	1.19
3	3	5.64	5.64	1.56
4	7	7.78	9.04	10.0
5	3	4.73	4.39	5.94
6	5	4.98	5.29	3.24
7	12	7.71	6.33	7.92
8	3	6.11	5.77	2.63
Average		6.01	5.98	4.76

Overall the fractal features proposed and the FCM based novelty detector can achieve a 4.76% MDR with a low 5.98% FDR on average. And we also note that those missing detection regions are usually located in the margins or weakness regions of the defects. Undoubtedly the method proposed can successfully label the major regions of defects.

Some figurative examples of detection results are given in *Figure 6*, where *Figure 6.a* and *6.b* are the normal and defective images, and *Figure 6.c* presents the detection results of *Figure 6.b*. Note that the red squares represent  $32 \times 32$  pixel sub-windows, which are identified as abnormal regions (defects). By viewing *Figure 6*, it can be found that the features proposed exhibit a good discriminating power on defects of linear shape or structural changes (see the first three columns in *Figure 6.c*). For the defects in the last column of *Figure 6.c*, the method proposed is able to detect the majority of defective regions, and the reasons for the missing regions are because they are located at the margins between detecting patches, providing not enough abnormal regions to identify them. Hence, in practical applications, we recommend using smaller patch sizes or overlapping patch division depending on specific process requirements.

## Conclusions

In this work, five fractal features have been extracted in a one-dimension projection series of fabric images by making use of the formation of fabric for defect detection. To better discriminate defects from the normal texture, the complementarity of fractal features extracted are fully taken into account, where the preliminary feasibility of those features in finding defects is verified via the t-test. In feature classification, a new FCM based novelty detector has been constructed to conduct an efficient novelty classification of features extracted, which has shown its effectiveness in handling the random distribution of normal samples. The experimental results for eight fabric datasets show that the method proposed can achieve a 4.76% MDR with a low 5.98% FDR on average, which further confirms the usefulness of the fractal features proposed for the FCM based novelty detector.



*Figure 6.* Figurative examples of detection results: a) normal fabric images, b) defective fabric images, c) detection results.

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## INSTITUTE OF BIOPOLYMERS AND CHEMICAL FIBRES

### LABORATORY OF BIODEGRADATION

The Laboratory of Biodegradation operates within the structure of the Institute of Biopolymers and Chemical Fibres. It is a modern laboratory with a certificate of accreditation according to Standard PN-EN/ISO/IEC-17025: 2005 (a quality system) bestowed by the Polish Accreditation Centre (PCA). The laboratory works at a global level and can cooperate with many institutions that produce, process and investigate polymeric materials. Thanks to its modern equipment, the Laboratory of Biodegradation can maintain cooperation with Polish and foreign research centers as well as manufacturers and be helpful in assessing the biodegradability of polymeric materials and textiles.

The Laboratory of Biodegradation assesses the susceptibility of polymeric and textile materials to biological degradation caused by microorganisms occurring in the natural environment (soil, compost and water medium). The testing of biodegradation is carried out in oxygen using innovative methods like respirometric testing with the continuous reading of the CO<sub>2</sub> delivered. The laboratory's modern MICRO-OXYMAX RESPIROMETER is used for carrying out tests in accordance with International Standards.



The methodology of biodegradability testing has been prepared on the basis of the following standards:

- **testing in aqueous medium:** 'Determination of the ultimate aerobic biodegradability of plastic materials and textiles in an aqueous medium. A method of analysing the carbon dioxide evolved' (PN-EN ISO 14 852: 2007, and PN-EN ISO 8192: 2007)
- **testing in compost medium:** 'Determination of the degree of disintegration of plastic materials and textiles under simulated composting conditions in a laboratory-scale test. A method of determining the weight loss' (PN-EN ISO 20 200: 2007, PN-EN ISO 14 045: 2005, and PN-EN ISO 14 806: 2010)
- **testing in soil medium:** 'Determination of the degree of disintegration of plastic materials and textiles under simulated soil conditions in a laboratory-scale test. A method of determining the weight loss' (PN-EN ISO 11 266: 1997, PN-EN ISO 11 721-1: 2002, and PN-EN ISO 11 721-2: 2002).



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The following methods are applied in the assessment of biodegradation: gel chromatography (GPC), infrared spectroscopy (IR), thermogravimetric analysis (TGA) and scanning electron microscopy (SEM).

#### Contact:

INSTITUTE OF BIOPOLYMERS AND CHEMICAL FIBRES  
ul. M. Skłodowskiej-Curie 19/27, 90-570 Łódź, Poland  
Katarzyna Dziedziczak Ph. D.,  
tel. (+48 42) 638 03 31, e-mail: lab@ibwch.lodz.pl