Research Article

Gastrointestinal polyp classification through empirical mode decomposition and neural features



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Abstract

This study describes an automated detection of polyp type as it is very important to determine the existence of dysplasia—a stage leading to the development of gastrointestinal cancer. The polyp-type classification is performed by a multiclass support vector machine from feature-fusion of bi-dimensional empirical mode decomposition (BEMD) and convolutional neural network (CNN). An extensive experiment is performed using standard datasets by extracted features from the individual technique as well as a fusion of features from BEMD and CNN. The fusion technique confirms satisfactory performance compared to other techniques with an accuracy of 98.94%. Moreover, it shows potentiality in precisely classifying some challenging polyps even though these are somehow confusing for human experts.

Keywords Video endoscopy · Gastrointestinal polyp detection and classification · Convolutional neural network · Empirical mode decomposition

1 Introduction

According to the world health organization, cancer is becoming the most leading cause of death around the world. Gastrointestinal cancer is very dreadful among them. The American Cancer Society has published a statistic in 2020 that 53,200 cancer-related deaths are predicted from 147,950 cases of gastrointestinal cancers [1]. In the USA, the statistic showed that it stood third among the most pervasive diseases and the second-largest cause of death in cancer [1]. However, it can be curable at an early stage if the malignancy of polyps is detected and classified at the preliminary level.

The usual medical method for gastrointestinal polyp diagnosis is video endoscopy. Polyp's shape and type are very confusing to the operators as these depend on various parameters. Sometimes the misclassification causes severe malignancy in the later stage. But all the categories of polyp are not exhibiting the same risk. Therefore, accurate polyp classification is very crucial.

Clinical studies have found gastrointestinal polyp as the precursor of developing cancer cells [2–5]. Several diagnosis criteria including routine endoscopy have been introduced to help the medical doctors in polyp detection. Usually, a polyp can be classified as Lumen, Diverticula, Adenoma, Hyperplastic, Serrated. The operator-dependent procedure of endoscopic examination may lead to polyp misdetection as well as misclassification. To reduce this error, the computer-aided system is very much helpful.

Recently, computer-aided polyp detection systems have been developed to work on endoscopic video frames using color, shape, texture, and spatial features [6–8]. These help in the diagnosis process of polyp detection, and, also indicates the affected regions that might be examined with more attention [9, 10]. In [11], image feature extraction from the covariance matrix of textural

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behavior of wavelet decomposition and linear discriminant analysis (LDA) was proposed for colonic tumor detection. An intelligent system was proposed using the combination of a support vector machine (SVM) and color-texture. It achieved around 94% accuracy. Vassilis and Maria [12] proposed an adaptive neuro-fuzzy system that can extract texture spectra in video capsule endoscopy (VCE) for polyp detection. A comparison of texture-based and color-based methods of polyp detection is investigated by Alexandre et al. [13]. Li et al. [14] reported a classifier using a multilayer perceptron along with SVM for polyp detection using color and shape features and achieved about 94% accuracy. Tajbakhsh et al. [15] proposed an automated polyp detection technique using global geometric constraints and local intensity variations. Our group reported two endoscopic video-based polyp detection techniques: one using color wavelet and convolutional neural network (CNN) features [16] and the other using bi-dimensional empirical mode decomposition (BEMD) and CNN features [17]. Now, it is time to concentrate on the polyp classification [18, 19]. Hence, the main focus of this paper is endoscopic video-based gastrointestinal polyp classification by SVM using the fusion of two kinds of features: BEMD and CNN.

The major contributions of this paper are:

- i. We develop an efficient polyp classification through the fusion of BEMD and CNN features.
- ii. We perform a detailed experiment to confirm the effectiveness of the proposed technique and compared it with other existing techniques.
- iii. We overcome the difficulties of diagnostic experts to identify the different classes of polyps through an automated system with improved accuracy.

The remaining part of this paper is structured as follows: Section 2 describes the proposed methodology. Polyp region segmentation as well as identification is highlighted in Sect. 3. The experimental results are analyzed in Sect. 4. Finally, conclusions are drawn in Sect. 5.

2 Methodology

This section outlines the proposed experiment to classify polyp classes. The system takes an endoscopic video as input and classifies the polyp in one of the five classes in the output. At first, frames are extracted from the video. Then after enhancement of each frame, a sliding window moves on the frame to capture image patches. After that, feature extractors run on the collected patches to grasp features using BEMD and CNN techniques. A final vector for a polyp classifier is prepared using the fusion of BEMD and CNN features. The SVM has performed to detect the polyp classes in the detection scheme. The self-explanatory major steps of the methodology are given in Fig. 1.

2.1 Dataset

In this paper, we have used 250 videos from two standard publicly available datasets: Endoscopic Vision Challenge dataset (https://polyp.grand-challenge.org/databases/) [20] and Department of Electronics, University of Alcala dataset (http://www.depeca.uah.es/colonoscopy_dataset/) [21]. From these 250 videos, we have extracted 14,900 frames with 5 different polyp types. Lumen and Diverticula typepolyp frames are extracted from Ref. [20] and Adenoma, Hyperplastic, and Serrated are from Ref. [21]. The detailed distributions of extracted video frames are given in Table 1.

2.2 Endoscopic video preprocessing

As mentioned earlier, the experiment has performed on 250 different endoscopic videos. In the training phase when any video feeds into the proposed system, then it transforms into the corresponding frames. A preprocessing technique is required to eliminate the unnecessary parts and irrelevant regions of the separated frames. Some of the extracted frames of different polyp classes are shown in Fig. 2.

Each frame is divided into sub-images of patch size 227×227 as shown in Fig. 3. The patch window goes from the top-left region to the bottom-right region of each frame with a stride of 15×15 . Through these patch collections, a sufficient amount of data has been produced from the captured frames. As supervised learning, this experiment demands huge quantities of training data to train the model. For the training scheme, it has required a sufficient number of the labeled dataset. Those image patches have alleviated the inadequacy of the labeled image. BEMD and CNN features are taken from these image patches.

2.3 Feature extraction

This section describes the feature extraction process using the proposed feature extractors: Bi-dimensional empirical mode decomposition (BEMD) and Convolutional Neural Network (CNN). The following Sects. 2.3.1 and 2.3.2 describe the BEMD and CNN feature extractors briefly.



Fig. 1 Major steps of polyp classification

 Table 1
 Polyp types with the extracted number of frames in endoscopic video

Polyp type	Number of extracted frames
Lumen [20]	2700
Diverticula [20]	2646
Adenoma [21]	3067
Hyperplastic [21]	3387
Serrated [21]	3100

2.3.1 Bi-dimensional empirical mode decomposition (BEMD)

Empirical mode decomposition (EMD) is a signal decomposition method, mostly used in signal processing to analyze the intuitive signal decomposition. In this research, we have used the self-adapting power of EMD in 2D image processing named bi-dimensional empirical mode decomposition (BEMD). BEMD, a 2D extension of EMD, has been developed for multiresolution image decomposition purposes [22–25]. It considers the 2D images as a series of bi-dimensional intrinsic mode functions (BIMFs) and a 2D residual with the same dimension of the original image. This process obeys two basic conditions:

- The number of extrema in the complete dataset must be equal to the number of zero crossings.
- The mean value of the local maxima and minima at any point of IMF should be zero.

Image decomposition into IMFs follows a shifting algorithm [17]. If the input image window is denoted as f(x, y), the proposed algorithm can be described as,

Name of the polyp	Extracted polyp frames from videos				
(a) Lumen		The Copy			
(b) Diverticula					
(c) Adenoma		000 100 00 000 100 000 100			
(d) Hyperplastic	100	000			
(e) Serrated					

Fig. 2 Sample polyp classes in some extracted frames: a, b from Ref. [20] and c-e from Ref. [21]



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Sub-images

Standard deviation (
$$\sigma$$
): $\sigma = \sqrt{\frac{\sum_{i=1}^{N} (Gl_i - \mu)^2}{N}}$ (3)

images

Variance
$$(\sigma^2)$$
: $\sigma^2 = \frac{\sum_{i=1}^{N} (GI_i - \mu)^2}{N}$ (4)

Kurtosis
$$(\kappa)$$
: $\kappa = \frac{\frac{1}{N} \sum_{i=1}^{N} (Gl_i - \mu)^2}{\left(\frac{1}{N} \sum_{i=1}^{N} (Gl_i - \mu)^2\right)^4} - 3$ (5)

Skewness(
$$\gamma$$
): $\gamma = \frac{\mu - Mode}{\sigma}$ (6)

Root mean square (*RMS*):
$$RMS = \sqrt{\frac{\sum_{i=1}^{N} (GI_i)^2}{N}}$$
 (7)

Smoothness
$$(\eta)$$
: $\eta = 1 - \frac{1}{1 + \sum_{i=1}^{N} GI_i}$ (8)

Sum average (f) :
$$f = \sum_{i=1}^{N} GI_i(x, y)$$
 (9)

Along with the statistical features, the calculation of GLCM features has a great value on the textural feature extraction from images. The five GLCM features used in this work are represented using Eqs. (10)–(14).

Contrast:
$$C = \sum_{i=0}^{M-1} \sum_{j=0}^{N-1} (i-j)^2 P(i,j)$$
 (10)

Correlation:
$$\rho = \frac{\sum_{i=0}^{M-1} \sum_{j=0}^{N-1} i j \mathcal{P}(i,j) - \sigma_x \sigma_y}{\sigma_x \sigma_y}$$
(11)

Energy:
$$E = \sum_{i=0}^{M-1} \sum_{j=0}^{N-1} P(i,j)^2$$
 (12)

Entropy:
$$S = \sum_{i=0}^{M-1} \sum_{j=0}^{N-1} P(i,j) \log_2 P(i,j)$$
 (13)

Homogeneity:
$$H = \sum_{i=0}^{M-1} \sum_{j=0}^{N-1} \frac{P(i,j)}{1 + (i-j)^2}$$
 (14)

where the dimension of the image is $M \times N$ and P(i, j) is the absolute value of intensity change of the pixel compared to its previous pixel.

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Executing this algorithm, a set of BIMFs and a residual *r* are acquired. They can reflect the original image as

$$f(x,y) = \sum_{1}^{n} BIMF_{i} + r(x,y)$$
(1)

Here, full decomposition of 2D multispectral image has been performed using BEMD. The proposed method separates each RGB image into three color channels (Red, Green, Blue). From there, take the first five BIMFs for analysis as textures, shapes, edges among 9-level BEMD features. Hence, twenty multi-scale images for each channel in four different directions: 0°, 45°, 90°, and 135° are found. This work computes thirteen different features containing statistical and GLCM features for each image, hence, a total of 780 BEMD features for three color channels. If the total pixels in the non-polyp region are defined as *M* and total pixels in the polyp region is *N* while the gray-levelcolor-intensity in polyp region is represented as *GI* for each pixel, then the statistical features are calculated using Eqs. (2)–(9) as follows.

Mean
$$(\mu)$$
: $\mu = \frac{\sum_{i=1}^{N} GI_i}{N}$ (2)

2.3.2 Convolutional neural network (CNN)

CNN outperforms usual state-of-the-art feature extraction methodologies in recent years [26]. Experiment on different CNN models, such as AlexNet, VGG16, VGG19, ResNet50, confirms that the AlexNet model performs better in our dataset. The convolution parts of the AlexNet architecture have extracted 4096 CNN features for each sub-image of size 227 × 227 in 3 color channels. This model has used five convolutional layers followed by a ReLU, 3 max-pooling layers, and 3-fully-connected layers. As a fiveclass classification scheme, this model has used 5 neurons in the output layer. The dropout functionality has been applied in the fully-connected layers with a 70% probability implementation. Along with dropout and an L2 regularization is remarkably effective to reduce overfitting as well as welfare for generalization error. The 'softmax' activation has applied for output which was later removed by the classifier. The architecture of this model is given in Table 2.

2.3.3 Support vector machine (SVM)

Detection and classification are usually done by using SVM [27–30], neural networks [16, 17, 31, 32], linear discriminant analysis (LDA) [11], adaptive neuro-fuzzy inference system [12], etc. Here, we used the SVM classifier, as it is robust for noisy and sparse data [29]. 780 color EMD features and 4096 CNN features are combined to construct the SVM input vector of 4876 features.

In this research, the classification layer of CNN is replaced by a multiclass SVM, as it uses less epoch and less processing time than the CNN classifier. SVM mimics the principle of one-against-all classification. A

Layer type	Size	Filter	Stride	Activation
conv1	55×55	96	4	ReLU
pool1	27×27	96	2	-
conv2	27×27	256	1	ReLU
pool2	13×13	256	1	-
conv3	13×13	384	1	ReLU
conv4	13×13	384	1	ReLU
conv5	13×13	256	1	ReLU
pool5	6×6	256	2	-
fc6	1×1	4096	-	ReLU + drop- out + L2-reg- ularization
fc7	1×1	1024	-	ReLU + drop- out + L2-reg- ularization
fc8 (output + clas- sification layer)	1×1	5	-	Softmax

hyperplane is considered that will divide each class with all other classes. This hyperplane tries to fit as the maximum distance between the data points of two halves. In this scheme, the SVM is interested to solve L_n binary classification where each current class L_i is classified versus other L_{n-1} class data. For the total of y data points, SVM will get L_y weight vectors as $W_1, W_2, ..., W_{Ly}$. When trying to classify any new point (x^*) , it will impose in a position further from the decision boundary by using $h(x^*)$. The class can be set by the following optimization equation.

$$y^* = \arg\max h(x^*).W_j \tag{15}$$

Figure 4 depicts the principle of SVM for five classes in this research.

3 Polyp region identification

After classifying polyp, this work has developed a procedure to identify and mark the polyp regions. Whenever any patch is detected as a polyp, the polyp region is marked by a square. A new circle is formed covering the most overlapped squares.

There are many possible polyp regions of various sizes. Figure 5a shows such possible polyp regions (see the marked regions). The classifier evaluates each marked region by a score. The final polyp region is extracted by averaging the higher score regions (see Fig. 5b). The detected polyp is classified as Hyperplastic.



Fig. 4 Multiclass SVM implementation for five classes

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Fig. 5 Polyp detection and classification: **a** several regions are marked as the candidate for a possible polyp, **b** a recognized polyp (marked region) and it is classified as Hyperplastic in a video frame taken from Ref. [20]

4 Experimental results and discussions

As mentioned in Sect. 2.1, we have done experimentation using 14,900 frames with different polyp types taken from two publicly available standard datasets. This experiment splits the whole dataset into two groups: 70% data (9900 polyp frames) are for training and 30% data (5000 polyp frames) are for testing. The training features are extracted from the training data and used to train the classifier. After that, testing is performed using testing data and classification performance is evaluated through the well-known metrics derived from the confusion matrix: accuracy, sensitivity, specificity, precision, F-value. These can be presented by the following equations.

$$Accuracy = \frac{TP + TN}{TP + TN + FP + FN} \times 100\%$$
(16)

Sensitivity =
$$\frac{TP}{TP + FN} \times 100\%$$
 (17)

Specificity =
$$\frac{TN}{FP + TN} \times 100\%$$
 (18)

$$Precision = \frac{TP}{TP + FP} \times 100\%$$
(19)

$$F - value = 2 \times \frac{Sensitivity \times Precision}{Sensitivity + Precision}$$
(20)

where TP is the true positive, FP is the false positive, TN is the true negative, and FN is the false negative. These metrics are performed on different image transform techniques based on GLCM and statistical features.

The first experimentation is done using only BEMD features. The classification results are shown in Table 3. In this table, we have also shown the obtained results from the direct original image features and the wavelet decomposition method for comparison. Performance figures are calculated using the average of all the polyp classes. The results confirm that the BEMD feature-based method is more effective.

The second experimentation is done with various architectural models of CNN. From Table 4, it is confirmed that AlexNet gives better accuracy compared to VGG16, VGG19, and ResNet50 models. For this, in the feature fusion, we used AlexNet model. Moreover, instead of SVM, we have also tried to classify the polyps using CNN. We have found that the SVM classifier is more efficient than the CNN classifier in terms of computation time and accuracy.

The third experimentation is done with feature fusion of BEMD and CNN to classify five types of polyps: Lumen, Diverticula, Adenoma, Hyperplastic, Serrated. Table 5 shows the confusion matrix for each type of polyp classification. The system faces no difficulty in classifying hyperplastic polyp as it is clear enough and quickly detect using the learning features. In a few instances, Lumen and

 Table 3
 The polyp classification performance measure using only BEMD features. Performances of other techniques are shown for comparison

Methods	Accuracy (%)	Sensitivity	(%) Specif	icity (%) F	Precision (%)	F-value (%)
Using the whole original image	68.74	67.98	69.52	e	8.59	68.70
Wavelet decomposition	74.73	74.12	75.11	7	'3.94	73.99
BEMD	80.81	79.38 81.57 80.07		30.07	80.46	
Table 4 The polyp classification performance	CNN architecture	Accuracy (%)	Sensitivity (%)	Specificity (%)	Precision (%)	F-value (%)
measure using various CNN	VGG16	91.94	89.91	92.42	90.19	90.60
models	ResNet50	91.97	91.03	92.09	91.83	91.91
	VGG19	92.71	91.18	92.87	91.27	92.66
	AlexNet	94.04	93.30	93.12	91.81	94.02

 Table 5
 Confusion matrix of polyp classification using featurefusion of BEMD and CNN

True class	Predicted class						
	Lumen	Diverticula	Adenoma	Hyperplas- tic	Serrated		
Lumen	883	17	0	0	0		
Diverticula	13	832	0	0	0		
Adenoma	0	0	1027	0	24		
Hyperplas- tic	0	0	0	1148	0		
Serrated	0	0	26	0	1030		

Diverticula look similar and thus making the classification confusing. Similar phenomena are occurring for Serrated and Adenoma. Table 6 shows the performance measure (accuracy, sensitivity, specificity, precision, and F-measure) for the classification of different polyp types. In this table, for comparison purposes, we have also shown the obtained results using CNN only features in classifying different polyp types. The results confirm that our proposed fusion method performs nicely compared to the CNN only feature-based method.

Finally, we have observed the performance of k-fold (k=5) cross-validation. The whole dataset has randomly partitioned into k equal subsamples. Among the k subsamples, all the k – 1 subsamples data are considered for the training set and only the remaining subsample has been used for testing the model. This process has repeated for k times by considering each subsample of k once for validating while all other k – 1 is for training. Table 7 shows the 5-fold cross-validation results for BEMD, CNN, and the fusion of BEMD and CNN. The results clearly show that the proposed fusion method outperforms both BEMD only and CNN only feature-based methods.

A few works have been performed on endoscopic polyp classification due to data scarcity. Zou et al. [33] used

 Table 7
 Accuracy of 5-fold cross-validation

Methods	5-fold cross- validation Accuracy
BEMD only	80.81
CNN (Alexnet) only	94.04
The fusion of BEMD and CNN (AlexNet)	98.94

DCNN in colonoscopy polyp classification. Experimenting with our data set we obtain 94% accuracy for DCNN. Pablo Mesejo et al. [21] used the computer-aided classification for Hyperplastic, Serrated, and Adenoma polyps. They analyzed texture and color features and used a multi-SVM as a classifier and obtained 90% accuracy. The study of Ribeiro et al. [18] classified only two polyp classes—Hyperplastic and Adenoma using the CNN model. They obtained 90.96% accuracy. The average accuracy in polyp classification using the proposed technique is compared with other polyp detection techniques and is shown in Table 8. The table confirms that our proposed fusion-based method shows improved accuracy than the existing polyp detection techniques.

Figure 6 shows some instances where a human expert faces difficulty, but these are correctly classified by our proposed method. Hence our automated system helps human radiological experts.

5 Conclusions

Machine-based polyp classification helps the physicians precisely diagnosis of gastrointestinal cancer at the early stage of polyps. With this motive, here we investigated an automated system for gastrointestinal polyp classification by SVM using the feature-fusion of BEMD and CNN techniques. We did three vital experiments using

Table 6	The performance measure o	f polyp	classification	using feature	e-fusion c	of BEMD	and CNN
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Classes	Method	Accuracy (%)	Sensitivity (%)	Specificity (%)	Precision (%)	F-value (%)
Lumen	CNN only	92.71	94.11	89.54	92.52	90.20
	Proposed method (BEMD + CNN fusion)	98.98	97.22	98.40	97.33	97.28
Diverticula	CNN only	92.54	92.47	90.42	91.98	90.07
	Proposed method (BEMD + CNN fusion)	98.91	97.16	98.39	97.04	97.11
Adenoma	CNN only	92.38	92.04	90.16	92.11	90.70
	Proposed method (BEMD + CNN fusion)	98.27	97.52	98.35	97.61	97.58
Hyperplastic	CNN only	96.25	92.97	95.63	94.23	93.42
	Proposed method (BEMD + CNN fusion)	99.60	99.82	99.84	99.83	99.86
Serrated	CNN only	93.71	91.98	91.90	92.17	90.87
	Proposed method (BEMD+CNN fusion)	98.96	97.63	98.32	97.54	97.58

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Table 8 Comparison of the proposed fusion-based classification with different polyp detection methods

Paper	Methodology	Dataset	Accuracy (%)
lakovidis et al. [3]	KL+Wavelet+SVM	86 videos	95
Alexandre et al. [13]	RGB + XY + SVM	4620 images	94.87
Li et al. [14]	RGB + MLP + SVM	450 images	94.20
Billah et al. [<mark>16</mark>]	CW+CNN+SVM	100 videos	98.60
Ribeiro et al. [18]	CNN	100 images	90.96
Zhu et al. [31]	CNN + SVM	180 images	85
Zou et al. [<mark>33</mark>]	DCNN	25 videos	96
Mesejo et al. [21]	Color + Texure + mSVM	76 videos	90
Proposed system	Color EMD + CNN + mSVM	250 videos	98.94



- (a) Adenoma (Serrated)
- **(b)** Lumen (Diverticula)



(d) Serrated (Adenoma)

Fig. 6 Some examples of challenging instances (b from Ref. [20] and a, c and d from Ref. [21]) where a human expert faces difficulty but these are correctly classified by our proposed method

standard datasets to classify the polyps into five classes. In the first experiment, we have found that BEMD feature-based method is effective than the wavelet decomposition technique. In the second experiment, we have found that AlexNet gives better accuracy compared to VGG16, VGG19, and ResNet50 CNN models for our dataset. In the third experiment, we have performed an indepth performance analysis of polyp classification using our BEMD and AlexNet CNN feature-fusion method. We have found that our fusion-based polyp classification method shows promising performance (accuracy over 98%) than the individual feature-based method as well as other existing polyp detection techniques. In the future, we will do more detailed experiments and try to fusion other important features for further improvement of accuracy to avoid misclassification for real-life implementation.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

Ethical approval (involvement of animals) This article does not contain any studies with animals performed by any of the authors.

Ethical approval (involvement of human subjects) This article does not contain any studies with human participants performed by any of the authors. The datasets used in the paper are publicly available. Proper acknowledgments with citation guidelines are maintained for the use of these datasets.

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