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Breast Cancer Classification Based on Improved Rough Set Theory Feature Selection

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Abstract. Breast cancer is one of the leading causes of death among the women. Mammogram analysis is the most effective method that helps in the early detection of breast cancer. In this paper we have made an attempt to classify the breast tissue based on Statistical features of a mammogram which extracted using simple image processing techniques with rough set theory. The proposed scheme uses texture models to capture the mammographic appearance within the breast. The statistical features extracted are the mean, standard deviation, smoothness, third moment, uniformity and entropy which signify the important texture features of breast tissue. Based on the values of these features of a digital mammogram, we have made an attempt to classify the breast tissue in to three basic categories normal, benign, and malignant given in the data base (mini-MIAS database). This categorization would help a radiologist to detect a normal breast from a cancer affected breast. Rough set theory can be regarded as a new mathematical tool for imperfect data analysis. Rough set based data analysis starts from a data table called a decision table. Each row of a decision table induces a decision rule, which specifies decision (action, results, outcome, etc.). We can know important data rules by using core and reduct which elimination of duplicate rows and elimination of superfluous values of attributes.

1. Introduction

Nowadays breast cancer detection and diagnosis is very important because the treatment is based on radio surgery. Breast cancer is the second leading cause of cancer death among women in the 40-55 ages[13]. Mammography is a key screening tool for breast abnormalities detection, because it allows identification of tumor before being palpable. Interpretations of mammographic images are difficult for the radiologists because of poor quality and noises in the images[1]. The proposed system is a novel scheme with rough set theory for abnormality detection and segmentation in Mammograms images, allows high degree of automation with less computation speed.

The breast is made up of many different types of cells. Breast cancers occur when one type of cell transforms from its normal characteristics and multiplies in an abnormal way. The extra cells form a mass of tissue called a tumor. Tumors are benign or malignant[12]. Many of them are benign and can be successfully removed. Malignant primary breast tumors cause problems by spreading into the normal tissue causing damage to the surrounding areas of the breast. According to the recent research results of

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University of Calgary, (Biomedical Engineering Research Group), these have been further classified based on some statistical features. This classification would help a radiologist to determine the breast anatomy (fibro glandular tissue) affected due to Estrogen secretion[2]. Rough set is founded on the assumption that with every object of the universe of discourse some information (data, knowledge) is associated. Objects characterized by the same information are indiscernible (similar) in view of the available information about them. Any set of all indiscernible (similar) objects is called an elementary set, and forms a basic atom of knowledge about the universe. Any union of some elementary sets is referred to as a crisp (precise) set otherwise the set is rough (imprecise, vague). Each rough set has boundary-line cases, i.e. objects which cannot be with certainty classified, by employing the available knowledge, as members of the set or its complement. Obviously rough sets, in contrast to precise sets, cannot be characterized in terms of information about their elements. With any rough set a pair of precise sets, called the lower and the upper approximation of the rough set, is associated. The lower approximation consists of all objects which surely belong to the set and the upper approximation contains all objects which possibly belong to the set. The difference between the upper and the lower approximation constitutes the boundary region of the rough set. Approximations are fundamental concepts of rough set theory. Rough set based data analysis starts from a data table called a decision table, columns of which are labeled by attributes, rows by objects of interest and entries of the table are attribute values. Attributes of the decision table are divided into two disjoint groups called condition and decision attributes, respectively. Each row of a decision table induces a decision rule, which specifies decision (action, results, outcome, etc.) if some conditions are satisfied. If a decision rule uniquely determines decision in terms of conditions the decision rule is certain. Otherwise the decision rule is uncertain. Decision rules are closely connected with approximations[3].

In previous study Mohabey et al. (2000) proposed an interesting strategy for the color image segmentation using the rough set theory [21]. Pal et al. (2002) proposed an integration of the Expectation Maximization (EM) algorithm, rough-set theoretic knowledge Extraction and minimal spanning tree (MST) clustering [22]. Aboul Ella Hassanien (2006) proposed Feature Extraction and Rule Classification Algorithm of Digital Mammography based on Rough Set Theory [23]. Cai et al. (2007) A modified classic fuzzy C-means (FCM) algorithm based on Rough set for image segmentation [24]. Jirava pavel, krupka Jiri (2007) classification model based on rough set and fuzzy sets theory [25]. Jiang et al. (2008) proposed a new method for the image segmentation based on the rough set theory and neural networks [26]. Halder et al. (2012) Proposed a rough set approach for a gray scale image segmentation [27]. Anupama et al. (2013) Rough sets based clustering is analyzed with respect to K-means and Fuzzy C-means algorithms for MRI images of Brain Web database [28]. Reddy et al. (2013) proposed an algorithm which was based on the modified K-means clustering by using rough set theory [17].

2. Problem formulation

We have made an attempt to classify the breast tissue based on features of a mammogram using simple image processing techniques. We have used texture models to capture the mammographic appearance within the breast. Different features can be extracted for a digital mammogram as: Texture Features, Statistical Features and Structural Features. We have used MATLAB for extracting the tumors from input mammogram and for calculating various features. Based on the values of these features of a digital mammogram, breast tissue can be classified. This categorization would help a radiologist to detect a normal breast from a cancer affected breast so as to proceed with further investigation [4].

3. Our technique

The main goals are to perform the following operations:

- Input data
- Detection
- Feature Extraction

- Decision table
- Rough set rules
- Classification
- Segmentation

This forms a basic step in the detection of abnormal breast under computer aided detection system

3.1. Input image

The studying of medical imaging applications plays an important role. In this study, we use the Mini MIAS database that contains 320 images. This dataset consists of 12 class's defined by the Breast Imaging Reporting and Data System (BI-RADS). There are four breast tissue classes (fatty, fibroglandular, heterogeneously dense, and extremely dense) and three health status classes (normal, benign tumor, and malignant tumor) for each breast tissue type[5]. There are 206 normal images, 63 benign and 51 malign (which are considered abnormal)[14].

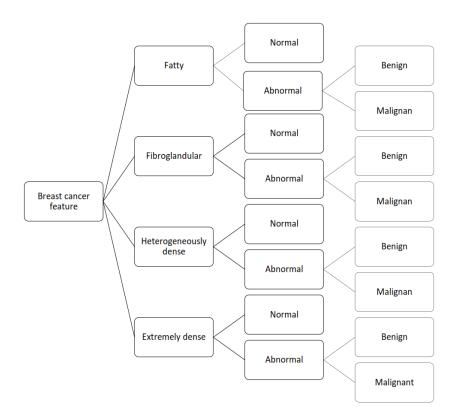


Figure 1: Flowchart designed for classify breast study.

3.2. Image processing

The preprocessing of digital mammograms refers to the enhancement of mammograms intensity and contrast manipulation, noise reduction, filtering, etc.

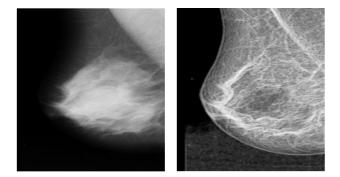


Figure 2: Image processing filter (a) image before filter. (b)image after filter.

3.3. Segmentation

The methods used to separate the region of interest from the background are usually referred as the segmentation process[6]. Segmentation can be carried out using any of the standard techniques like local thresholding, K-means clustering [15].

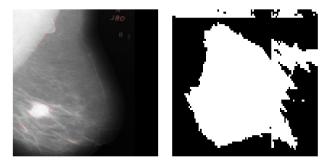


Figure 3: Segmentation image.

3.4. Feature Extraction

In our work, we have extracted Texture features, Statistical features and Structural Features for the segmented tumor from the given input mammogram image[7]. The features that we have extracted are:

no	moment	expression	Description
1	Contrast=inertia=sum of squares variance	$cont = \sum_{i,j} i - j ^2 p(i, j)$	A measure of the image contrast or the amount of local variations present in an image.
2	Inverse Difference Moment (IDM) =Homogeneity	$idm = \sum_{i,j} \frac{p(i,j)}{1+ i-j ^2}$	This descriptor has large values in cases where the largest elements in P are along the principal diago- nal.
3	Dissimilarity=Directional moment	$dissim = \sum_{i,j} i - j p(i, j)$	A measure of the image contrast that increase lin- early not exponentially.

Table 1: Contrast group mathematical representations.

no	moment	expression	Description
1	Angular second mo- ment(ASM)	$Asm = \sum_{i,j}^{n} p(i,j)^2$	A measure of the ho- mogeneity of an image. Hence it is a suitable mea- sure for detection of dis- orders in textures. For ho- mogeneous textures value of angular second mo- ment turns out to be small compared to non- homogeneous ones.
2	Energy=Uniformity	$Ene = \sqrt{Asm}$	Energy returns the sum of squared elements in the Grey Level Co-occurrence Matrix (GLCM). Energy is also known as uniformity. The range of energy is [0 1].
3	min probability	$Minimum{Xi i = 1, 2, 3,, N}$	This is simply the largest entry in the matrix.
4	max probability	$Maximum{Xi i = 1, 2, 3,, N}$	This is simply the smallest entry in the matrix.
5	Entropy	$Ent = -\sum_{i=1}^{n} p(i) \log_2 p(i)$	Entropy is a measure of in- formation content. It mea- sures the randomness of intensity distribution.

Table 2: Measures related to orderliness mathematical representations.

no	moment	expression	Description
1	Mean	$\mu = \sum_{i,j}^{n} p(i,j)$	The mean, m of the pixel values in the defined window, estimates the value in the image in which central clustering occurs.
2	Standard Deviation	$\sigma = \sqrt{\sum_{i=1}^{n} (p(i) - \mu)^2}$	The Standard Deviation, σ is the estimate of the mean square deviation of grey pixel value p (i, j) from its mean value Standard deviation describes the dispersion within a local region.
3	Variance	$var = \sqrt{\sigma}$	Variance is the square root of standard deviation.
4	Area descriptor	$ADes = \frac{\sigma}{\mu}$	Area descriptor is Outside the division between standard deviation and mean.
5	Skewness	$skew = \sum_{i=1}^{n} \left(\frac{p(i) - \mu}{\sigma}\right)^{3}$	Characterizes the degree of asymmetry of a pixel distribution in the specified window around its mean. Skewness is a pure number that characterizes only the shape of the distribution.
6	Kurtosis	$kurt = \sum_{i=1}^{n} \left(\frac{p(i) - \mu}{\sigma}\right)^{4}$	Measures the Peakness or flatness of a distribution relative to a normal distribution.
7	Fourth moment	$M4 = \sum_{i=1}^{n} (p(i) - \mu)^4$	Measures the Peakness of a distribution relative to a normal distribution expo- nentially (4).
8	Third moment	$M3 = \sum_{i=1}^{n} (p(i) - \mu)^{3}$	Measures the Peakness of a distribution relative to a normal distribution expo- nentially (3).
9	Mean energy	$\mu_{energy} = \frac{1}{n} \sum_{i=1}^{n} p(i)^2$	Measures the men energy intensity in the histogram.
10	Energy variance	$var_{energy} = \frac{1}{n-1} \sum_{i=1}^{n} (p(i)^2 - \mu_{energy})^2$	Measures the Energy variance intensity in a region.
11	Correlation	$\frac{\mu_{energy})^2}{\sum_{i,j}^{corr} \frac{(1-\mu_i)(j-\mu_j)p(i,j)}{\sigma_i\sigma_j}} =$	Correlation returns a measure of how correlated a pixel is to its neighbor over the whole image. The range of corre- lation is [-1 1].Correlation is 1 or -1 for a perfectly positively or negatively cor- related image. Correlation is Nan (Not a Number) for a constant image.
12	Smoothness	$smo = 1 - \frac{1}{1 + \sigma^2}$	A measure of grey level contrast that can be used to establish descriptors of relative smoothness.
13	Root Mean square(RMS)	$rms = \sqrt{\frac{\sum_{i=1}^{n} p(i) ^2}{n}}$	The RMS (Root Mean Square) com- putes the RMS value of each row or column of the input, along vectors of a specified dimension of the input, or of the entire input
14	Similarity	$sim = \sum_{i=1}^{n} \frac{p(i, j)}{1 + i - j }$	A first degree measure that increases with less contrast.

Table 3: Descriptive statistics group mathematical representations.

3.5. Decision table

A decision table specifies what decisions (actions) should be undertaken when some conditions are satisfied. Most decision problems can be formulated employing decision table formalism; therefore, this tool is particularly useful in decision making. A decision table has two subsets of attributes, called condition *C* and decision attributes *D* respectively. Decision table will be denoted T = (U, A, C, D). A decision table specifies what decisions (actions) should be undertaken when some conditions are satisfied. Most decision problems can be formulated employing decision table formalism; therefore, this tool is particularly useful in decision table has two subsets of attributes, called condition *D* respectively. Decision table formalism; therefore, this tool is particularly useful in decision making. A decision table has two subsets of attributes, called condition *C* and decision attributes *D* respectively. Decision table T = (U, A, C, D).

U			С		D			
	s_1	<i>s</i> ₂		Sm	d_1		d_k	
<i>x</i> ₁	f_{11}	f ₁₂		f_{1m}	g_{11}		g_{1k}	
<i>x</i> ₂	f ₂₁	f ₂₂		f_{2m}	g_{21}		g_{2k}	
•								
•								
x_n	f_{n1}	f_{n2}		f _{nm}	g_{n1}		g_{nk}	

Table 4: General fo	rm of decision table.
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3.6. Rough set

The idea of rough set was publicized by Professor Zdzislaw Pawlak, a Polish Mathematician, in 1982, for the development of automatic rule generation systems[8]. Using Rough set theory distorted regions of tumor can be detected. Rough set theory can be characterized as a new mathematical tool for imperfect data analysis. The theory has found applications in many domains, such as decision support, engineering, medicine and others[3].

3.6.1. Fundamental concepts in rough sets

Classical rough set theory often requires a large amount of labeled data, in which concepts information system, indiscernibility, approximation and attribute reduction are key issues.

1. Information system

The information system *T* can be expressed as a quadruple T = (U, A, C, D), where *U* is the set of entities (universe). *A* is the attribute set. If set *A* can be divided into conditional attribute set *C* and decisional attribute set *D*, i.e., $C \cup D = A$ and $C \cap D = \phi$.

2. Indiscernibility

If $P \subseteq R$ and $P \neq \phi$, then $\bigcap P$ (intersection of all equivalence relations belonging to *P*) is also an equivalence relation, and will be denoted by *IND*(*P*), and will be called an indiscernibility relation over *P*. Moreover

$[X]IND(P) = \bigcap_{(R \in P)} [x]R$

So U/IND(P) (i.e. the family of all equivalence classes of the equivalence relation IND(P)).

3. Approximation of set

Assume that $X, Y \in U$ and R is the equivalent relationship defined on the universe U. The lower-approximation set of the set X on R is defined as

$$\underline{\mathfrak{R}}(x) = \bigcup \{ Y \in U \nearrow R : Y \subseteq X \}$$
(1)

Where $\Re(x)$ is the maximum set of entities that for sure belong to the set *X*, also referred to as positive region, denoted by *POS*(*X*). Similarly, the upper-approximation set of the set *X* on *R* is defined as

$$\overline{\mathfrak{R}(x)} = \bigcup \{ Y \in U / R : Y \bigcap R \neq \phi \}$$
(2)

Where ϕ is an empty set and $\overline{\Re(x)}$ is the minimum set of entities which possibly belong to set *X*. Based on the above definitions, the boundary set can be defined as

$$BND(X) = \overline{\mathfrak{R}(x)} - \underline{\mathfrak{R}(x)}$$
(3)

If BND(X) is an empty set, the set X reduces to a crisp set on R. If on the other hand, BND(X) is non-empty; X is a rough set on R.

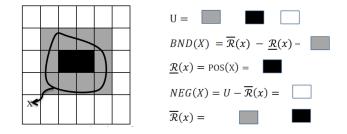


Figure 4: Approximations of set.

4. Attribute reduction

 $Q \subseteq P$ is a reduct of *P* if *Q* is independent and IND(Q) = IND(P). Obviously *P* may have many reducts[9]. The set of all indispensable relations in *P* will be called the core of *P*, and will be denoted CORE(P).

$$CORE(P) = \left(\begin{array}{c} RED(P) \end{array}\right) \tag{4}$$

Where RED(P) is the family of all reducts of *P*. Let *R* and *Q* be equivalence relations over *U*, *R*-positive region of *Q*, denoted $POS_R(Q)$ we understand the set

$$POS_R(Q) = \bigcup_{(x \in U \neq Q)} \underline{\mathfrak{R}(x)}$$
(5)

R- Positive region of *Q* is the set of all object of universe *U* which can be classified to class of U/Q.

3.6.2. Accuracy measure

Let $K = (U, \mathfrak{R})$ be the knowledge base and $R, Q \subset \mathfrak{R}$. We say Q depend in a degree k from knowledge R if and only if

$$K = \gamma_{R(Q)} = \frac{cardPOS_R(Q)}{cardU}$$
(6)

Where card denotes cardinality of the set statistics[18]. We have three phases

- 1. if k = 1, Q totally depends from R, and then all elements of the universe can be classified.
- 2. If 0 < k < 1, we say Q roughly depend from R, and then only elements that belong to positive region can be classified.
- 3. if k = 0, we say Q is totally independent from R, and then none of the elements of the universe can be classified.

3.7. Classification

Classification aim at simplifying the uncertain decision table and generating more significant decision rules to classify unseen objects[10]. K-nearest neighbor classification (KNN) is a classification model in which you can alter both the distance metric and the number of nearest neighbors. Because a Classification KNN classifier stores training data, you can use the model to compute resubstitution predictions. Classification KNN performance is directly related to the parameter k, there is no obvious information on the selection of k except that it should be positive and not a multiple of the total number of classes[16]. Alternatively, use the model to classify new observations using the predict method[11].

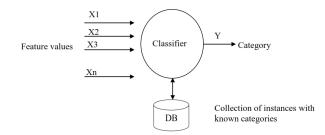


Figure 5: Classifier model.

The presented algorithm was implemented as a toolbox in MATLAB environment. This instrument is further applied

Algorithm 1 classification and segmentation data

	<i>Input</i> : breast cancer images from our data base
	Output: get new classification of breast cancer and segmentation
3:	procedure Get image from data set, then calculate feature extraction
4:	for each item Group no.1 feature extractions do
5:	Create matrix T called decision table
6:	From decision table apply reduct to remove repeat data and overflow values from Eq.4
7:	Compute lower and upper approximations $Eq.1, 2$
8:	Calculate value of k from <i>Eq.</i> 6
9:	if $0 < k \le 1$ then
10:	elements that belong to positive region can be classified
11:	else
12:	none of the elements of the universe can be classified
13:	calculate new feature extraction
14:	end if
15:	Apply KNN classifier
16:	if image normal then
17:	show it
18:	else
19:	compute canny segmentation method for abnormal images
20:	end if
21:	Show result data
22:	compute Accuracy from Eq.7
23:	end for
24:	return Accuracy
25:	end procedure

27

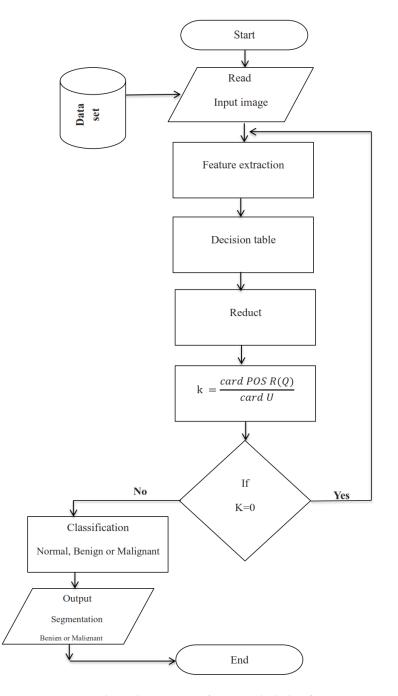


Figure 6: Shows the sequences of steps in which classification occur.

4. Numerical results

We have used MATLAB for extracting the tumors from input mammogram and for calculating various features.

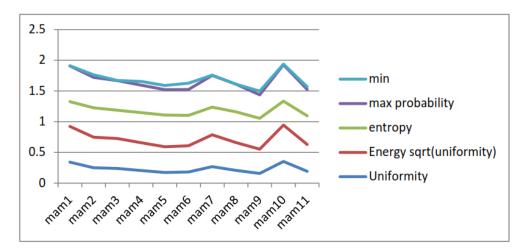
First: Numerical results for measures related to orderliness data set

Image Sam- ples	Uniformity	Energy sqrt(uniformity)	Entropy	max probability	min probability
mam1	0.3401	0.5832	0.40196	0.5814	0.0038147
mam2	0.2487	0.4987	0.47658	0.4963	0.042915
mam3	0.2379	0.4877	0.45951	0.4819	0.0019073
mam4	0.2044	0.4521	0.49022	0.4457	0.062943
mam5	0.1734	0.4164	0.51761	0.412	0.068665
mam6	0.181	0.4254	0.49559	0.4193	0.10395
mam7	0.2684	0.5181	0.44958	0.5158	0.005722
mam8	0.2058	0.4537	0.49878	0.4505	0.00095367
mam9	0.1562	0.3952	0.50232	0.3816	0.060081
mam10	0.3507	0.5922	0.39049	0.5906	0.01049
mam11	0.191	0.4370	0.46683	0.4268	0.044823
mam12	0.1424	0.3774	0.53311	0.3682	0.00095367
mam13	0.3358	0.5795	0.40655	0.5781	0.0047684
mam14	0.2718	0.5213	0.45606	0.5195	0.005722
mam15	0.3619	0.6016	0.36236	0.5985	0.005722
mam16	0.3873	0.6223	0.37094	0.6214	0.014305
mam17	0.4467	0.6684	0.3063	0.6648	0.00095367
mam18	0.382	0.6181	0.32479	0.6097	0.00095367
mam19	0.2073	0.4553	0.50349	0.4528	0.024796
mam20	0.199	0.4461	0.49606	0.4423	0.02861
mam21	0.1434	0.3787	0.52569	0.3653	0.019073
mam22	0.2634	0.5132	0.4379	0.5085	0.002861
mam23	0.145	0.3808	0.51044	0.3654	0.0047684
mam24	0.1458	0.3818	0.50057	0.3652	0.00095367
mam25	0.1701	0.4124	0.54099	0.4096	0.00095367
mam26	0.1908	0.4368	0.52046	0.4342	0.0019073
mam27	0.1388	0.3726	0.53774	0.3652	0.00095367
mam28	0.137	0.3701	0.55805	0.3652	0.0047684
mam29	0.1478	0.3844	0.4855	0.3652	0.00095367
mam30	0.2564	0.5064	0.426977	0.5	0.00095367

Table 5: Measures related to orderliness data set.

U	С							
	Uniformity	Energy sqrt(uniformit	Entropy y)	max prob- ability	min prob- ability	decision		
mam1	0.3401	0.5832	0.40196	0.5814	0.0038147	1		
mam2	0.2487	0.4987	0.47658	0.4963	0.042915	1		
mam3	0.2379	0.4877	0.45951	0.4819	0.0019073	0		
mam4	0.2044	0.4521	0.49022	0.4457	0.062943	0		
mam5	0.1734	0.4164	0.51761	0.412	0.068665	1		
mam6	0.181	0.4254	0.49559	0.4193	0.10395	0		
mam7	0.2684	0.5181	0.44958	0.5158	0.005722	0		
mam8	0.2058	0.4537	0.49878	0.4505	0.00095367	0		
mam9	0.1562	0.3952	0.50232	0.3816	0.060081	0		
mam10	0.3507	0.5922	0.39049	0.5906	0.01049	1		
mam11	0.191	0.4370	0.46683	0.4268	0.044823	0		
mam12	0.1424	0.3774	0.53311	0.3682	0.00095367	1		
mam13	0.3358	0.5795	0.40655	0.5781	0.0047684	1		
mam14	0.2718	0.5213	0.45606	0.5195	0.005722	0		
mam15	0.3619	0.6016	0.36236	0.5985	0.005722	1		
mam16	0.3873	0.6223	0.37094	0.6214	0.014305	0		
mam17	0.4467	0.6684	0.3063	0.6648	0.00095367	1		
mam18	0.382	0.6181	0.32479	0.6097	0.00095367	0		
mam19	0.2073	0.4553	0.50349	0.4528	0.024796	1		
mam20	0.199	0.4461	0.49606	0.4423	0.02861	0		
mam21	0.1434	0.3787	0.52569	0.3653	0.019073	1		
mam22	0.2634	0.5132	0.4379	0.5085	0.002861	0		
mam23	0.145	0.3808	0.51044	0.3654	0.0047684	2		
mam24	0.1458	0.3818	0.50057	0.3652	0.00095367	0		
mam25	0.1701	0.4124	0.54099	0.4096	0.00095367	1		
mam26	0.1908	0.4368	0.52046	0.4342	0.0019073	0		
mam27	0.1388	0.3726	0.53774	0.3652	0.00095367	0		
mam28	0.137	0.3701	0.55805	0.3652	0.0047684	2		
mam29	0.1478	0.3844	0.4855	0.3652	0.00095367	0		
mam30	0.2564	0.5064	0.426977	0.5	0.00095367	1		

Table 6: Measures related to orderliness decision table.



In decision table for Measures related to orderliness group 0=normal, 1=Benign and 2=Malignant.

Figure 7: Data representation for Measures related to orderliness group.

Second: Descriptive statistics group data set

mam9

53.5637

mam10 43.1755

73.2029

69.3156

5.36E+03

4.80E+03

0.0761

0.0688

	Table 7: Some descriptive statistics and their decision table.										
U		С									
	mean	Standard	deviation	smoothn	esstrea De-	skewne	sskurtosis	Third	Fourth	contrast	de
			vari-		scriptor			mo-	mo-		
			ance					ment	ment		
mam1	36.5348	63.4199	4.02E+03	0.0583	1.7359E+00	1.5346	3.8219	0.0236	0.0146	9.85E+04	1
mam2	49.3319	73.0232	5.33E+03	0.0758	1.4802E+00	1.1902	2.8879	0.0279	0.0194	1.02E+05	1
mam3	52.0066	80.0201	6.40E+03	0.0896	1.5387E+00	1.1485	2.6306	0.0355	0.0255	1.07E+05	0
mam4	59.4984	82.6967	6.84E+03	0.0952	1.3899E+00	0.9093	2.0935	0.031	0.0232	1.10E+05	0
mam5	64.4811	69.7978	4.87E+03	0.0697	1.0825E+00	0.4249	1.5451	0.0087	0.0087	1.24E+05	1
mam6	70.0907	72.7462	5.29E+03	0.0753	1.0379E+00	0.2896	1.4227	0.0067	0.0094	1.07E+05	0
mam7	49.86	74.2953	4.86E+03	0.0695	1.4901E+00	0.913	2.1428	0.0186	0.012	1.10E+05	0
mam8	61.9692	74.2953	5.52E+03	0.0782	1.1989E+00	0.5931	1.6595	0.0147	0.012	9.70E+04	0

1.3667E+00

1.6054E+00

1.1 . 1.1

D decision

1.32E+05 0

5.91E+04 1

0.0219

0.0242

0.0155

0.0148

2.2758

2.7173

0.9236

1.2032

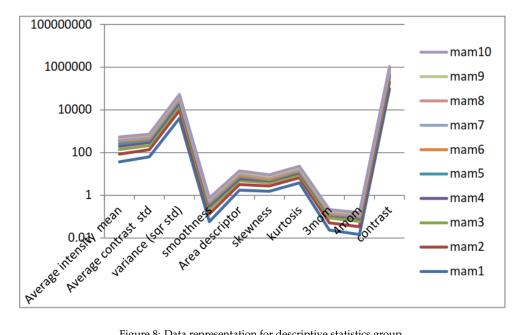


Figure 8: Data representation for descriptive statistics group.

Third: Contrast group data set

U		С					
	contrast	Inverse	Dissimilarity	decision			
		Difference					
		Moment					
mam1	9.85	4.9	2.434129	1			
mam2	10.19	3.9	2.604502	1			
mam3	10.73	4.3	2.549482	0			
mam4	10.98	4.1	2.648189	0			
mam5	12.36	3	2.940145	1			
mam6	10.71	3.7	2.659529	0			
mam7	11.05	3.8	2.633394	0			
mam8	9.70	4.1	2.473971	0			
mam9	13.16	3.3	2.937944	0			
mam10	5.91	4.5	2.002846	1			
mam11	12.83	3.4	2.891276	0			
mam12	10.75	4	2.630655	1			
mam13	10.50	4.5	2.529737	1			
mam14	8.96	4.1	2.375325	0			
mam15	10.64	4.2	2.623059	1			
mam16	6.61	4.4	2.09246	0			
mam17	9.13	4.3	2.412082	1			
mam18	6.92	4	2.194056	0			
mam19	13.12	3	3.020435	1			
mam20	10.01	3.9	2.527979	0			

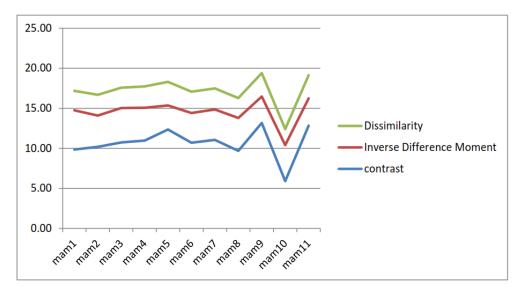


Figure 9: Data representation for contrast group.

- 1. with the reduction result Y = [1, 2], which means that the feature Energy and uniformity in our system irreducible. The features Entropy, max probability and min probability can be removed without affecting the recognition of system.
- 2. degree of dependence $k = \frac{2}{5} = 0.4$ this means elements that belong to positive region can be classified.
- 3. Accuracy of the classification is obtained by using the given equation:

$$Accuracy = \frac{(TP + TN)}{(TP + TN + FN + FP)} \times 100$$
(7)

Where:

True positive=TP: Correctly classified as having breast cancer.

True negative=TN: Correctly classified as not having breast cancer.

False positive=FP: Classified as having breast cancer but actually they don't have cancer.

False negative=FN: Classified as not having breast cancer but actually they have cancer[19].

no	sample	Knn classi- fier	TP	TN	FP	FN	accuracy	average
1	5		1	4	0	0	100	86.8
2	50		30	12	6	2	84	
3	100		56	22	12	10	78	
4	100		64	20	10	6	84	
5	200		121	55	9	15	88	

Table 9: Classification result accuracy.

5. Conclusion

In this study, the tumor identification and the investigation are carried out with the help of rough theory for the potential use of mini-breast data for improving the tumor classification. Using rough set theory feature extraction is performed efficiently and this made segmentation easier varying tissue characteristics[20]. Although there is no known way to prevent breast cancer, mortality can be reduced only with early diagnosis. Therefore, the computer aided diagnosis (CAD) systems are very important as they allow radiologists to reconsider mammogram images with increased sensitivity of detection and diagnosis. The health status classification is performed with two consequent stages, where the normal and abnormal mammograms are determined first, and the abnormal defined mammograms are then classified as benign and malignant. These results helpful for radiologists to make more accurate breast cancer diagnoses.

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